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Relationship between renin-angiotensin-aldosterone system and renal Kir5.1 channels

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Abstract

K_{ir}5.1 (encoded by *Kcnj16* gene) is an inwardly rectifying $K^+(K_{ir})$ channel highly expressed in the aldosterone-sensitive distal nephron of the kidney where it forms a functional channel with $K_{ir}4.1$. $K_{ir}4.1/K_{ir}5.1$ channels are responsible for setting the transepithelial voltage in the distal nephron and collecting ducts and are thereby major determinants of fluid and electrolyte distribution. These channels contribute renal blood pressure control and have been implicated in salt-sensitive hypertension. However, mechanisms pertaining to the impact of $K_{ir}4.1/K_{ir}5.1$ mediated K^+ transport on the renin-angiotensin-aldosterone system (RAAS) remain unclear. Herein, we utilized a knockout of Kcnj16 in the Dahl salt-sensitive rat (SS^{Kcnj16−/-}) to investigate the relationship between $K_{ir}5.1$ and RAAS balance and function in the sensitivity of blood pressure to the dietary Na^+/K^+ ratio. The knockout of *Kcnj16* caused substantial elevations in plasma RAAS hormones (aldosterone and angiotensin peptides) and altered the RAAS response to changing the dietary Na^{+}/K^{+} ratio. Blocking aldosterone with spironolactone caused rapid mortality in SS^{Kcnj16-/-} rats. Supplementation of the diet with high K⁺ was protective against mortality resulting from aldosterone-mediated mechanisms. Captopril and losartan treatment had no effect on the survival of SS^{Kcnj16-/−} rats. However, neither of these drugs prevented mortality of SSKcnj16−/− rats when switched to high Na+ diet. These studies revealed that the knockout of Kcnj16 markedly altered RAAS regulation and function, suggesting K_i 5.1 as a key regulator of the RAAS, particularly when exposed to changes in dietary sodium and potassium content.

Declarations of interest

The authors declare that they have no conflicts of interest with the contents of this article.

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Keywords

RAAS; salt-sensitive hypertension; Kcnj16; Kcnj10; hypokalemia; ENaC; HSD11β2

Introduction

In the kidney, Na^+ and K^+ transport in the late distal convoluted tubule (DCT2), connecting tubule (CNT), and cortical collecting duct (CCD), which collectively comprise the aldosterone-sensitive distal nephron (ASDN), is a critical determinant of the pressurenatriuresis relationship responsible for the long-term control of arterial pressure [1, 2]. Basolaterally expressed inwardly rectifying $K^+(K_{ir})$ channels are critical for K^+ transport in these segments [3–5]. The K_{i} 4.1/ K_{i} 5.1 heterotetramer is the predominant basolateral K+ channel in the ASDN and sets the basolateral membrane potential, determines the driving force for ion transport, and senses and controls plasma K^+ levels [6–8]. We previously reported that the K_i -4.1/ K_i -5.1 heteromeric channel is required for the kidney to maintain numerous homeostatic mechanisms, including acid/base balance, whole body K+ homeostasis, and blood pressure control. In particular, we found that a high salt (HS) diet caused elevated blood pressure accompanied by increased expression of $K_{ir}5.1$ in the ASDN of the Dahl salt-sensitive rat, an established model of salt-sensitive (SS) hypertension. Our previous studies revealed that a knockout of Kcnj16 in the Dahl saltsensitive rat (SS^{Kcnj16−/−}) eliminated renal K_{ir}4.1/K_{ir}5.1 channel expression and function in the kidney and resulted in hypotension, salt wasting tubulopathy, and hypokalemia, which was exacerbated and led to mortality within a few days when SS^{Kcnj16-/-} rats were fed a high salt diet [9]. K_{ir} 5.1 has been proposed to be involved in the responsiveness of blood pressure to diet, and dysfunction of this channel may contribute to the pathophysiology of salt-sensitive hypertension. Although it is clear that $K_{ir}4.1/K_{ir}5.1$ channels regulate ion transport and influence blood pressure in the ASDN [5, 10], little progress has been made in defining the mechanism and extent of these channels' effect on the renin-angiotensinaldosterone system (RAAS).

The RAAS is a hormonal homeostatic system responsible for sustaining blood pressure in spite of inevitable fluctuations in plasma electrolytes resulting from variable dietary intake. The main effector peptides of the RAAS, angiotensin II (Ang II) and aldosterone, respond to deviations in plasma Na^+ and K^+ , respectively, and act on the kidney to maintain blood pressure homeostasis. RAAS dysfunction or dysregulation is associated with numerous cardiorenal pathologies, and overactivity of the RAAS is associated with the development of hypertension and progression of chronic kidney disease (CKD). The capacity for RAAS hormones to alter renal ion transport is key to its effects on blood pressure. In the ASDN, the Na+-Cl− co-transporter (NCC) and the epithelial Na+ channel (ENaC) are both responsive to aldosterone signaling and have been established as significant contributors to the development of SS hypertension [11, 12]. We and others previously reported that NCC and ENaC were inappropriately activated in Dahl SS rats fed a HS diet, and inhibition of these Na+ transporters was sufficient to prevent the development of SS hypertension [11, 13–17]. However, modulation of Na⁺ transport by basolateral $K_iA_1/K_i.5.1$ channels in the pathophysiology of SS hypertension has not been fully elucidated. The mechanism of

 $K_{ir}4.1/K_{ir}5.1$ channels' involvement in salt-sensitive hypertension and the capacity of these channels to influence the RAAS requires investigation and is the main goal of this study. Despite their overlapping functionality and a shared site of action in the ASDN, there is thus far little evidence directly implicating renal $K_{ir}4.1/K_{ir}5.1$ channels in RAAS homeostasis or function. We hypothesize that $K_{ir}5.1$ subunit exerts its control over blood pressure both through its established role in determining fluid and electrolyte homeostasis and via an additional regulatory role in RAAS modulation. We utilized a rat model with a knockout of Kcnj16 on the Dahl SS rat background to investigate the capacity of $K_{ir}5.1$ to influence RAAS signaling in the sensitivity of blood pressure to the dietary Na^+/K^+ ratio.

Methods

Animals.

The genetic background used for the studies is the re-derived Rapp Dahl SS rat (SS/ JrHsdMcwi; referred to as SS^{WT}), which has been inbred for over 50 generations at the Medical College of Wisconsin (MCW). This strain is a proven model for the study of saltsensitive hypertension [11, 18–20]. A Kcnj16 knockout rat model ($SS^{Kcnj16-/-}$) on the Dahl SS rat background was created at the MCW Gene Editing Rat Resource Center using zinc finger nuclease technology to induce an 18 base-pair deletion in the second transmembrane domain (TM2) of the $K_{ir}5.1$ protein, resulting in the knockout of $K_{ir}5.1$ subunit as previously described [9, 21]. Animals were housed in a temperature and humidity-controlled facility and maintained on a standard 12/12 dark/light cycle with water and food provided ad libitum. SS^{Kcnj16-/-} and SS^{WT} rats were fed a purified AIN-76A diet (#113755, Dyets, Inc., Bethlehem, PA) containing 0.4% NaCl and 0.36% K⁺ from wean unless otherwise noted (LS LK; 0.4% NaCl represents normal salt content but referenced here as a LS for simplicity in comparison to high salt (HS) diet). For experiments investigating the impact of altering dietary sodium and potassium intake the following diet compositions are used and specified in the text: HS LK (# 113756), HS HK (# 113522) and LS KH (# 113521) all form Dyets, Inc., Bethlehem, PA. All rats except SS^{Kcnj16-/−} rats on a HS LK diet (this diet results in rapid mortality of $SS^{Kcnj16-/-}$ rats [9] requiring samples to be taken for measurement after 24 hours) were fed the specified diet for a 5 week duration before measurements were taken at approximately 12 weeks of age. Numbers of male and female rats included in all experimental groups are indicated in all figure legends.

Quantification of RAAS hormones.

The characterization of RAAS components was performed by quantification of the steadystate angiotensin peptide and aldosterone levels in equilibrated heparinized plasma samples by Attoquant Diagnostics (Vienna, Austria). Stable isotope-labeled internal standards for both classical and alternative renin-angiotensin system (RAS) branch constituents (Ang I (1-10), Ang II (1-8), Ang III (2-8), Ang 1-7, Ang IV (3-8), and Ang 1-5) were added to stabilized plasma samples at a concentration of 200 pg/mL and subjected to liquid chromatography tandem-mass spectrometry (LC-MS/MS)-based angiotensin quantification as described previously [22–25]. It was reported that similar outcomes are achieved using protease inhibitor-stabilized and equilibrated samples for angiotensin peptide quantification, thus both methods are representative of equilibrium hormone levels [25]. This technique was

used to evaluate RAAS hormones in plasma samples from SS^{WT} and SS^{Kcnj16−/−} rats fed each of four experimental diets: 1) LS LK (standard diet: 0.4% NaCl, 0.36% K⁺); 2) LS HK $(0.4\%$ NaCl, 1.41% K⁺); 3) HS LK (4% NaCl, 0.36% K⁺); and 4) HS HK (4% NaCl, 1.41% K^+). These analyses included six angiotensin metabolites (Ang I (1-10), Ang II (1-8), Ang III (2-8), Ang IV (3-8), Ang 1-7, and Ang 1-5), aldosterone levels, and ratios of different derivatives to give surrogate measurements of angiotensin-converting enzyme (ACE) activity (Ang II/Ang I), plasma renin activity (PRA, Ang I + Ang II), and the adrenal response to Ang II (AA2-Ratio).

Administration of inhibitors targeting RAAS and analysis of electrolytes.

To evaluate functional and physiological repercussions of altered RAAS signaling in $SS^{Kcnj16-/-}$ rats, we administered pharmacological inhibitors targeting aldosterone and angiotensin signaling in vivo. Spironolactone, a specific mineralocorticoid receptor (MR) antagonist, was used to block the action of aldosterone. Rats were given once daily IP injections of 50 mg/kg spironolactone for up to 10 days. This dose was chosen for maximal antagonism of MR in the ASDN while still safe and well tolerated in rats [26]. To prepare injections, spironolactone (S3378; Sigma-Aldrich) was dissolved in DMSO and ethanol at a 1:5 ratio and vehicle injections were performed (N=4) indicating no alteration to plasma electrolytes in control animals. A similar procedure was repeated to administer the angiotensin converting enzyme inhibitor (ACEi) captopril (C4042; Sigma-Aldrich), and angiotensin receptor blocker (ARB) losartan (PHR1602; Sigma-Aldrich), which were also given once daily IP (50 and 30 mg/kg/day, respectively in 0.9% saline). SSWT rats were used as a control strain in comparison with the effects of RAAS modulation in the SS^{Kcnj16−/−} strain.

To evaluate changes in renal ion transport, urine and blood electrolyte composition were assessed in SS^{WT} and $SS^{Kcnj16-/-}$ rats before and after administration of corresponding drugs. Rats were individually housed in metabolic cages to allow 24-hour urine collection at multiple time points. Control urine was collected before beginning any treatment. After the initial 24-hour control urine collection, rats received a daily dose of spironolactone or captopril/ losartan administered by IP injection. Standard 24-hour urine collection was repeated after the first administration and at the end of the protocol. Arterial blood samples were collected from the abdominal aorta under isoflurane anesthesia, and cortical renal tissue was harvested after flushing the blood from the kidneys. Electrolyte concentrations, pH, creatinine, and hemoglobin (ctHb) measurements from whole blood and 24-hour urine samples were determined by ABL800 FLEX blood gas analyzer (Radiometer, Copenhagen, Denmark) in SS^{WT} and SS^{Kcnj16−/−} rats (both LS LK and LS HK diets) before and after injections. For comparison, arterial blood was also analyzed from untreated $SS^{Kcnj16-/-}$ rats and SSWT control rats.

Histological and biochemical analyses.

To harvest kidneys for immunohistochemistry (IHC), rats were anesthetized with isoflurane and kidneys were flushed with PBS via aortic catheterization at 3 ml/min per kidney until blanched. Kidneys were formalin fixed (10% normal buffered formalin), paraffin embedded, sectioned (4 μm), and mounted on slides as previously described [9]. Hydroxysteroid 11-

β dehydrogenase 2 (HSD11β2) was visualized with HSD11β2 antibody (1:100; Abcam ab203132) and streptavidin-biotin immunohistochemistry. Images were created using Nikon Eclipse Ni-E microscope (20X/0.50 and 40X/0.75 objectives). For Western blot analysis, kidney cortical lysates were prepared as previously described [9]. Kidney tissue samples (15-25 mg) were pulse sonicated in Laemmli buffer with a protease inhibitor cocktail (Roche, Basel Switzerland) to achieve a final protein concentration of 25 mg/ml. The resulting supernatant was subjected to SDS-PAGE, transferred onto nitrocellulose membrane (Millipore, Bedford MA) and probed with antibodies to α-ENaC (1:1000, Stressmarq Biosciences Inc, Vancouver, cat# SPC-403D) or HSD11β2 (1:1000; Abcam ab203132), and visualized by enhanced chemiluminescence (Amersham Biosciences Inc, Piscataway, NJ). Antibody to β-actin was used as control.

Statistical methods.

Blood and urine measurements are presented in tables as mean ± SEM. RAAS hormone measurements are presented in boxplots with individual data points displayed. Bounds of boxes represent SEM, whiskers span three standard deviations, median is denoted by a horizontal line within each box, and mean is denoted by a square within each box. For statistical analysis of data, equal variance and normality (Shapiro-Wilk) tests were performed followed by two-way ANOVA to determine significant effects resulting from the knockout and dietary treatments. For data comparisons before and after drug administration, one-way repeated measures ANOVAs were used. Data which failed the normality test was analyzed using the Kruskal-Wallis one-way ANOVA on ranks followed by post hoc tests. P-values of less than 0.05 were considered significant.

Study approval.

All animal studies were conducted at Medical College of Wisconsin and protocols were approved by the MCW Institutional Animal Care and Use Committees (IACUC) and were performed in accordance with the standards set forth by the NIH Guide for the Care and Use of Laboratory Animals.

Results

SSKcnj16−/− rats exhibit broad elevations in plasma RAAS hormones.

To evaluate whether $K_{ir}5.1$ impacts overall RAAS balance, we used a highly sensitive mass spectrometry-based method to quantify equilibrium plasma hormone levels in SSWT and SS^{Kcnj16−/−} rats on a standard diet (LS LK; 0.4% NaCl, 0.36% K⁺). Angiotensin metabolites (Ang I, Ang II, Ang III, Ang IV, Ang 1-7, Ang 1-5) and aldosterone were dramatically elevated in SS^{Kcnj16-/-} compared to SS^{WT} rats (Figures 1 and 2C). Plasma levels of Ang II and aldosterone were increased by an order of magnitude in the knockout (371 vs. 3674 and 183 vs. 1878 pM Ang II and aldosterone in SS^{WT} and $SS^{Kcnj16-/-}$ rats, respectively). Clinical metrics and surrogate measures of enzyme activity were computed based on RAAS hormone quantification. As renin is the rate-limiting enzyme of the RAAS, plasma renin activity (PRA) is used clinically to estimate RAAS activation in patients and identify cases of essential hypertension. PRA, represented by the total quantity of Ang I and Ang II, was significantly elevated in SS^{Kcnj16−/−} rats compared to SS^{WT} rats (Figure 2B). Angiotensin

converting enzyme (ACE) activity, represented by the Ang II to Ang I ratio, was also elevated in the $Kcnj16$ (K_{ir}5.1) knockout (Figure 2A). However, the adrenal response to Ang II was not altered in the knockout strain (AA2 ratio; can be used clinically to detect primary aldosteronism and represented by the aldosterone to Ang II ratio; Figure 2D). There were no sex differences in measurements of angiotensin metabolites (Figure S3), aldosterone (Figure S4A), or surrogate functional parameters (PRA, ACE activity, and AA2 ratio; Figures S4B– D) in SS^{Kcnj16–/–} rats.

Although RAAS values were compared between SS^{WT} and SS^{Kcnj16} and ST^{Kcnj16} LS LK diet, it is important to note that these rats were exposed to vastly differing plasma K⁺ levels (Figure S7), as we previously showed that SS^{Kcnj16–/−} rats are hypokalemic on the LS LK diet [9]. To control for this discrepancy in plasma K^+ , we performed the same RAAS measurements in SS^{Kcnj16−/−} rats fed a LS HK diet (0.4% NaCl, 1.41% K⁺). Although plasma K^+ was partially restored by dietary potassium supplementation, it was not enough to protect from hypokalemia, and corresponding plasma RAAS levels in $SS^{Kcnj16-/-}$ rats on the LS HK diet did not become comparable to SSWT rats (Figure S5). Ang II became further elevated, but aldosterone was unaffected on the LS HK diet compared to the LS LK diet (Figures 1B and 2C). Overall, the substantial differences in RAAS balance observed in the $SS^{Kcnj16-/-}$ rats persisted despite dietary K⁺ supplementation and augmentation of plasma K^+ .

Knockout of Kcnj16 altered the RAAS response to dietary Na+/K+ ratio.

Our previous work examined the effects of varying dietary Na^+ and K^+ content on blood pressure in SS^{WT} and SS^{Kcnj16-/-} rats, and found that K_{ir}5.1 was required for diet-specific blood pressure responses and salt-sensitive hypertension [9]. To determine whether the observed effect of $K_{ir}5.1$ on blood pressure occurred at least partially through diet-specific RAAS changes, we measured the RAAS response to systematically varying Na^+ and K^+ intake in SSWT and SSKcnj16−/− rats. Plasma RAAS hormones were quantified in SSWT and SS^{Kcnj16−/−} rats being fed each of four diets consistent with our previous blood pressure study: 1) LS LK (standard diet: 0.4% NaCl, 0.36% K⁺); 2) LS HK (0.4% NaCl, 1.41% K⁺); 3) HS LK (4% NaCl, 0.36% K⁺); and 4) HS HK (4% NaCl, 1.41% K⁺) diets.

Effect of HS diet on plasma RAAS hormones.—We and others reported that the administration of a HS LK diet elicits a robust increase in blood pressure, which is characteristic of the salt-sensitive hypertension phenotype in SSWT rats [9]. Plasma RAAS analysis in SS^{WT} rats showed that elevating dietary Na⁺ without altering K⁺ in the HS LK diet attenuated aldosterone but did not change Ang II levels compared to the standard LS LK diet (Figures 1, 2C, S1 and S2). Figures S1 and S2 show an expanded scale of the data from Figures 1 and 2C and represent a summary of angiotensin metabolite balance in SS^{WT} rats. None of the angiotensin metabolites evaluated in SS^{WT} rats were affected by the HS LK diet, except Ang 1-5 which was moderately elevated (Figure S1). As previously shown, the HS LK diet has vastly different physiological consequences in SS^{Kcnj16−/−} rats compared to SSWT rats. Specifically, the transition from a LS LK to a HS LK diet did not promote blood pressure elevation in these rats, but stimulated K^+ wasting and exacerbated hypokalemia, which resulted in rapid mortality of all SS^{Kcnj16-/−} rats within a few days [9].

Plasma RAAS quantification in $SS^{Kcnj16-/-}$ rats revealed that the switch to HS LK diet more than doubled Ang II (and all other angiotensin metabolites) and dramatically attenuated plasma aldosterone levels preceding HS-induced deaths (Figures 1B and 2C, respectively). In addition, HS LK treatment in SSWT increased ACE activity but did not alter PRA. Conversely, HS LK did not alter ACE2/ACE balance in SSKcnj16−/− rats but did elevate PRA. The HS LK diet decreased the AA2 ratio similarly in both SS^{WT} and $SS^{Kcnj16-/-}$ rats (Figure 2).

Capacity of dietary potassium to modulate the RAAS response on a HS diet.

—To evaluate RAAS balance and the contribution of dietary K+ to modulate the renal response to Na⁺, plasma RAAS quantification was performed in SS^{WT} and SS^{Kcnj16-/-} rats fed a HS HK diet. The analysis revealed that supplementing the HS diet with HK caused a modest elevation in plasma Ang II in SSWT rats, but significantly reduced Ang II to levels comparable to the LS LK diet in SS^{Kcnj16-/−} rats (Figures 1B and S1B). Accordingly, PRA, which was elevated in SS^{Kcnj16−/−} rats on a HS LK diet, returned to baseline levels with the addition of HK. The HS HK diet also augmented the AA2 ratio and plasma aldosterone in SS^{Kcnj16−/−} rats, partially restoring it to LS LK levels. The HS HK diet did not change ACE activity in SS^{Kcnj16−/−} rats. The addition of HK to HS did not alter plasma aldosterone, AA2 ratio, PRA, or ACE activity in SS^{WT} rats (Figure 2).

Knockout of Kcnj16 altered balance between the renin-angiotensin system (RAS) axes.

To assess alterations in the balance between the classical and alternate RAS axes, plasma angiotensin effector peptides were grouped according to whether they primarily contribute to the classical (Ang II, Ang III, and Ang IV) or alternative (Ang 1-7 and Ang 1-5) RAS branch. We found that alternative branch peptides represented a greater proportion of the total angiotensin metabolites in SS^{Kcnj16-/-} rat plasma compared to SS^{WT} plasma (Figure 3). This finding is consistent with the $SS^{Kcnj16-/-}$ rat phenotype, which includes lower basal mean arterial pressure and reduced renal fibrosis [9]. Furthermore, the balance between the classical and alternative axes was markedly altered by Na⁺ and K⁺ intake in SS^{Kcnj16-/-} rats but remained relatively stable in SS^{WT} rats regardless of diet (Figure 3).

In $SS^{Kcnj16-/-}$ rats, angiotensin metabolites responded to dietary alterations distinctly according to whether the peptide belongs to the classical or alternative RAS branch. In the classical branch of the RAS, LS HK caused modest increases in Ang II, Ang III, and Ang IV, while the addition of HS in the HS HK diet caused these metabolites to return to control LS LK levels (Figure 1). In the alternate RAS branch, where Ang 1-7 is the main effector, Ang 1-7 and Ang 1-5 respond to LS HK with a modest decrease in the metabolite, which is sustained with the addition of high $Na⁺$ on the HS HK diet (Figure 1). Thus, unlike the classical RAS axis, the alternate axis is responsive to potassium intake but insensitive to sodium intake in SS^{Kcnj16–/−} rats.

Kcnj16 knockout increased renal ENaC and HSD11β**2 expression.**

To better understand the causes and effects of aberrant aldosterone signaling in SS^{Kcnj16−/−} rats, we assessed the expression of two proteins which are critically involved in the ASDN aldosterone signaling pathway: hydroxysteroid 11-beta dehydrogenase type 2 (HSD11β2)

and ENaC. It is established that aldosterone binds to the mineralocorticoid receptor (MR) to exert its effects on sodium transport in the ASDN through the stimulation of activity and expression of ENaC and NCC. Although MR is widely expressed in the kidney, it has equal affinity for both glucocorticoids and mineralocorticoids. The ASDN is the major site for aldosterone signaling due to co-expression of MR with HSD11β2 specifically in this nephron segment. HSD11β2 is an enzyme complex that oxidizes and inactivates glucocorticoids preventing their binding to MR, accounting for the specificity for MR to bind aldosterone in the ASDN. Deficiency in HSD11β2 has also been implicated in saltsensitive hypertension [27–29]. Western blot and IHC analysis revealed increased expression of HSD11β2 in the distal tubules of $SS^{Kcnj16-/-}$ rats compared to SS^{WT} controls (Figures 4A and C, respectively). Furthermore, downstream targets of aldosterone signaling were found to be elevated in SSKcnj16−/− rats compared to SSWT rats. Subunit expression of α -ENaC was significantly elevated in SS^{Kcnj16–/–} rats (Figure 4B) similar to increased expression of both total and phosphorylated form of NCC found previously and represent compensatory overexpression or sodium transporters in response to significant electrolyte wasting phenotype observed in $SS^{Kenj16-/-}$ rats [9].

Targeted pharmacological RAAS inhibition revealed that amplified aldosterone signaling is required for survival of SSKcnj16−/− rats.

To better understand the physiological implications of the aberrant RAAS signaling and differences between angiotensin and aldosterone pathways, we used spironolactone, captopril, and losartan in vivo to inhibit MR, ACE, and angiotensin receptor, respectively. Blocking aldosterone signaling with spironolactone caused mortality in all SS^{Kcnj16−/−} rats (Figure 5A) without significantly altering their low plasma K^+ levels (Table 1). Survival of SS^{WT} rats was unaffected by spironolactone treatment and plasma electrolyte measurements reveal a typical response to a potassium-sparing diuretic (Table 1 and Figure S5). As our previous studies have shown that potassium supplementation improved the cardio-renal phenotype and survival of $SS^{Kcnj16-/-}$ rats [9], we predicted that increased dietary potassium intake would compensate for hypokalemia in SS^{Kcnj16-/−} rats and restore spironolactone-mediated potassium-sparing diuretic effects observed in SSWT rats. Potassium supplementation in SS^{Kcnj16-/-} rats dramatically changed their response to spironolactone, showing changes in plasma electrolytes that were more similar to SSWT rats than to SS^{Kcnj16−/−} rats on the standard LS LK diet (Table1 and Figure S5). Like SS^{WT} rats, LS HK fed SS^{Kcnj16-/−} rats did not experience mortalities with spironolactone treatment. Table 1 summarizes plasma electrolytes in SSWT and SSKcnj16−/− rats (fed either LS LK or LS HK diets) injected with spironolactone. Changes in urinary electrolyte excretion following spironolactone administration are shown in Figure S6. In contrast to spironolactone, inhibiting Ang II signaling with captopril (ACEi) or losartan (ARB) had no effect on survival of LS LK fed $SS^{Kenj16-/-}$ rats (Figure 5B). Unlike the effects of K⁺ supplementation, neither captopril nor losartan were able to prevent SS^{Kcnj16-/−} rats' mortalities when switched to the HS LK diet. Changes in urinary electrolyte excretion with daily captopril or losartan administration in SS^{WT} and SS^{Kcnj16-/-} rats are shown in Table S1.

Discussion

Basolateral $K_{ir}4.1/K_{ir}5.1$ channel is responsible for setting the resting membrane potential and the transepithelial voltage in the distal nephron and collecting ducts and are thereby major determinants of fluid and electrolyte distribution [4, 30]. Importantly, this channel have been predicted to contribute to renal control of blood pressure and have been implicated in salt-sensitive hypertension [9]. However, the specific mechanisms and extent of this regulatory role remains unclear. The impact of $K_{ir}4.1/K_{ir}5.1$ channel activity on blood pressure control has previously been explained by its ability to alter the kidneys' pressure natriuresis response through its control over sodium reabsorption and fluid balance [10]. RAAS is well established for its role in regulating blood pressure, electrolyte homeostasis and fluid volume. We hypothesized that $K_{ir}5.1$ exerts its control over blood pressure both through its established role in determining K^+ homeostasis and *via* an additional regulatory role of RAAS signaling in ASDN. We utilized a knockout of Kcnj16 on the Dahl SS rat background to investigate the relationship between $K_{ir}5.1$ and the RAAS in the sensitivity of blood pressure to the dietary Na^+/K^+ ratio.

Consistent with our hypothesis, the knockout of $Kcnj16$ ($K_{ir}5.1$) resulted in extensive elevations in aldosterone and angiotensin metabolites in SS^{Kcnj16−/−} rats compared to SS^{WT} rats (Figures 1 and 2), indicating the direct crosstalk between K_{ir}5.1 function and RAAS balance. Although ACE activity and PRA were both elevated in $SS^{Kcnj16-/-}$ rats, AA2 ratio remained unchanged, indicating that the knockout does not alter Ang II-dependent aldosterone production by the adrenal glands. Physiologically, such an excess in plasma Ang II and aldosterone should correspond with blood pressure elevations, but $SS^{Kcnj16-/-}$ rats have chronically low blood pressure [9]. This clearly suggests that $K_{ir}5.1$ is critical not only to maintain plasma RAAS balance, but also to mediate functional RAAS responses in the kidney required for appropriate blood pressure adjustments.

Although Ang II and aldosterone were similarly elevated in SS^{Kcnj16−/−} rats, their dietspecific responses and in vivo consequences were distinct. Increased aldosterone in $SS^{Kcnj16-/-}$ rats was partially attenuated with increased Na⁺ intake (HS LK), but elevations in Ang II became further increased (Figure 2C and 1B, respectively). Increased dietary K⁺ is known to have beneficial effects in modulating salt-sensitive hypertension by altering renal $Na⁺$ handling. We previously reported that supplementing the HS challenge with HK (HS HK diet) attenuated but did not abolish salt-induced blood pressure elevations in SSWT rats, and completely prevented the salt-induced mortalities in SS^{Kcnj16−/−} rats [9]. Notably, SS^{Kcnj16-/-} rats showed no HS-induced blood pressure response, indicating a loss of the salt-sensitive phenotype. Furthermore, targeted pharmacological inhibition of Ang II and aldosterone revealed that the amplified aldosterone, but not Ang II signaling is critical for the survival of $SS^{Kcnj16-/-}$ rats (Figure 5). Although specific mechanisms remain unclear, our combined results suggest that RAAS regulation by $K_{ir}4.1/K_{ir}5.1$ channels is mediated by aldosterone rather than angiotensin signaling in the kidney.

Our results suggest that $K_{ir}4.1/K_{ir}5.1$ channels' critical role as potassium sensors in the distal nephron is required to dynamically alter the renal response to aldosterone dependent on present environmental conditions to maintain blood pressure and electrolyte homeostasis.

Sometimes referred to as "the aldosterone paradox," high plasma aldosterone can be induced by either hyperkalemia or hypovolemia, but to vastly different effects [31]. In hyperkalemic conditions, aldosterone signaling in the ASDN results in net K^+ secretion to restore K^+ homeostasis without affecting blood pressure, while in hypovolemia aldosterone causes Na⁺ retention, volume restoration and increases blood pressure. Herein, we found that without contribution from $K_{ir}5.1$, aldosterone loses this flexibility, suggesting a crucial regulatory role of $K_{ir}4.1/K_{ir}5.1$ channels is required for the dynamic physiological effects of aldosterone. Although more work needs to be done to delineate a precise mechanism, this study, paired with the work of others, elucidates a promising potential mechanism. In states of hyperkalemia, high plasma K^+ inhibits $K_{ir}4.1/K_{ir}5.1$ channels, depolarizing the basolateral membrane which results in less electroneutral NaCl reabsorption through NCC in the early DCT (DCT1) and consequently higher $Na⁺$ delivery to the ASDN. This, along with aldosterone, increases $Na⁺$ reabsorption through ENaC, driving $K⁺$ secretion through apical channels [32]. The outcome is K^+ secretion without Na⁺ retention and thus, no increase in blood pressure. However, if high aldosterone is a result of hypervolemia, the opposite mechanism should occur. Increased $K_{ir}4.1/K_{ir}5.1$ activity hyperpolarizes the membrane NCC activation in DCT1, reducing Na⁺ delivery to the ASDN resulting in net Na⁺ retention without K⁺ secretion. In SS^{Kcnj16-/-} rats' hypovolemic conditions, aldosterone should functionally increase blood pressure without K^+ wasting. However, this cannot occur in SS^{Kcnj16−/−} rats because this mechanism is dependent on K^+ sensing by basolateral $K_{ir}4.1/K_{ir}5.1$ channels. Accordingly, deletion of $K_{ir}5.1$ has been shown to result in chronic depolarization of the basolateral membrane [33], which would eliminate aldosterone's ability to respond dynamically to extracellular potassium, effectively forcing aldosterone to permanently function as if in hyperkalemic conditions. The resulting chronic K^+ secretion without $Na⁺$ retention provides an elegant explanation for how such high plasma aldosterone, which is classically associated with hyperkalemia and increased blood pressure, can be paired with hypokalemia and low blood pressure in SS^{Kcnj16-/-} rats. Dysfunction of this mechanism may also contribute to the loss of the salt-sensitive phenotype in $SS^{Kcnj16-/-}$ rats as well as aldosterone's apparent insensitivity to dietary K^+ modulation.

Although the results of this study indicate that $K_{ir}5.1$ is likely most directly involved in aldosterone signaling, alterations in RAS balance and diet-specific responses of angiotensin peptides are likely very relevant to the SS^{Kcnj16−/−} phenotype. The overall function of the RAS is dependent on the balance between the 'classical' and 'alternative' axes, which are known to have opposing physiological effects. The classical axis produces vasoconstriction, inflammation, and increases in blood pressure, while the alternative axis causes vasodilation and is anti-inflammatory. We found that peptides of the alternative RAS branch (Ang 1-7 and Ang 1-5) were proportionally more represented in SSKcnj16−/− rats than in SSWT rats on all diets. This may contribute to low blood pressure in SS^{Kcnj16−/−} rats. Also, balance between the classical and alternative RAS was more affected by alterations in diet composition in SS^{Kcnj16−/−} rats compared to SS^{WT} rats. Specifically, in SS^{Kcnj16−/−} rats, the alterative axis was responsive to dietary potassium but not sodium intake, which could partially explain their corresponding loss of the SS hypertensive phenotype.

Overall, we conclude that $K_{ir}5.1$ is critical for specific RAAS responses to alterations in the dietary Na^+/K^+ ratio, for RAAS hormones to exert their effects on blood pressure, and for

maintaining appropriate balance between the classical and alternative RAS axes. The genetic knockout of $K_{ir}5.1$ caused an apparent uncoupling of the RAAS from renal ion transport in the ASDN, eliminating blood pressure responsiveness to Na^{+}/K^{+} intake, resulting in the previously reported resolution of SS hypertensive phenotype. This study supports $K_{ir}4.1/$ $K_{ir}5.1$'s involvement in the aldosterone paradox, as the knockout $K_{ir}5.1$ resulted in loss of dynamic control of aldosterone function in the ASDN, effectively blinding aldosterone to environmental conditions, compromising its role in blood pressure homeostasis. We propose Kir5.1 as a key protein in regulating plasma RAAS hormones and facilitating physiological responses to RAAS signaling for the maintenance of homeostasis.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Clinical Perspectives:

- Renal K_{ir}5.1 containing channels play a role in blood pressure modulation in response to dietary salt. The renin-angiotensin-aldosterone system (RAAS) also has a prominent role in diet-specific blood pressure alterations. However very little is known about the capacity for these channels to modulate plasma RAAS hormones or the potential involvement of $K_{ir}5.1$ in RAAS signaling in the kidney.
- We found that genetic knockout of *Kcnj16* to eliminate K_{ir}5.1 channels resulted in profound elevations in plasma RAAS hormones and altered the response to dietary sodium and potassium.
- Understanding the relationship between RAAS signaling and renal K_{ir}5.1 channels in diet-specific blood pressure responses may elucidate novel therapeutic targets or strategies for a subset of hypertensive patients.

Figure 1. Quantification of equilibrium angiotensin metabolites in SSKcnj16−/− and SSWT rats. Ang I (1-10) (**A**), Ang II (1-8) (**B**), Ang III (2-8) (**C**), Ang 1-7 (**D**), Ang IV (3-8) (**E**), and Ang 1-5 (**F**) were quantified from SS^{Kcnj16-/- (N=5-10 males) and SS^{WT} (N=4-7 males)} plasma samples collected from rats being fed one of 4 diets. In the boxplots, black data points denote rats fed a low salt, low potassium diet (LS LK, 0.4% NaCl and 0.36% K⁺); green indicates a low salt, high potassium diet (LS HK, 0.4% NaCl and 1.41% K⁺); red indicates a high salt, low potassium diet (HS LK, 4% NaCl and 0.36% K⁺); and blue indicates a high salt, high potassium diet (HS HK, 4% NaCl and 1.41% K⁺). Please see

Figure S1 for expanded scale of angiotensin derivatives in SS^{WT} rats. Measurements are shown in pM. Bounds of boxes represent SEM, whiskers span 3 standard deviations, median is denoted by a horizontal line, and mean is denoted by a square within each box. p -values are given for each graph.

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Figure 2. Equilibrium aldosterone and surrogate RAAS measures in SSWT and SSKcnj16−/− rats. RAAS components were quantified using a mass spectrometry-based analysis of plasma samples collected from $SS^{Kcnj16-/-}$ (N=5-10 male) and SS^{WT} (N=4-7 male) rats fed one of 4 diets as described in Figure 1. (**A**) ACE activity indicated by the ratio of plasma Ang II to Ang I. (**B**) Plasma renin activity (PRA) represented by the combined content of Ang I and Ang II in pM. (**C**) Plasma aldosterone in pM (please see Figure S2 for expanded scale of aldosterone level in SSWT rats). (**D**) The ratio of aldosterone to Ang II (AA2 ratio). Bounds of boxes represent SEM, whiskers span 3 standard deviations, median is denoted by a horizontal line, and mean is denoted by a square within each box. p-values are given for each in the figure graph.

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Classical vs. Alternative Plasma Angiotensin Peptides

Figure 3. Angiotensin peptides of the classical versus alternative RAS axis.

The proportion of classical axis related peptides (Ang II, Ang III, and Ang IV) and alternative axis related peptides (Ang 1-7 and Ang 1-5) represented in plasma from SS^{WT} and SS^{Kcnj16−/−} rats. Proportions shown represent data compiled from raw angiotensin metabolite measurements depicted in Figure 1 (N=4-7 male SS^{WT} and N=5-10 male SS^{Kcnj16−/−} rats). Diets are indicated on the y-axis and the percentage of plasma angiotensin metabolites designated to the classical RAS axis versus the alternative RAS axis is represented on the x-axis.

Figure 4. ENaC and HSD11β**2 expression in SSWT and SSKcnj16−/− rats.**

(A) Western blot analysis of HSD11β2 expression in kidney tissue from SSWT and SS^{Kcnj16-/-} rats on the standard LS LK diet (0.4% NaCl and 0.36% K⁺; N=5 males per group). (**B**) Western blot quantification of α-ENaC protein in kidney tissue from SSWT and SS^{Kcnj16−/−} rats on a LS LK diet (N=5 males per group). *p*-values are given for each graph. (**C**) Representative immunohistochemical staining of HSD11β2 protein in SSWT and SS^{Kcnj16−/−} rats at 20x and 40x magnification. Scale bars are 20 µm.

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Figure 5. Survival of SSWT and SSKcnj16−/− rats following pharmacological RAAS modulation. (**A**) Kaplan-Meier survival probability of SSWT rats on a LS LK diet (0.4% NaCl and 0.36% K⁺; N = 7), SS^{Kcnj16-/-} rats on a LS LK diet (0.4% NaCl and 0.36% K⁺; N = 17 total, 10 males and 7 females), and SSKcnj16−/− rats on a LS HK diet (0.4% NaCl and 1.41% K⁺; N = 6) were given IP injections of 50 mg/kg spironolactone daily for 10 days. (**B**) Kaplan-Meier survival probability of $SS^{WT} (N=3$ males per ACEi and ARB treatment group) and $SS^{Kcnj16-/-}$ rats (N=4 males per ACEi and ARB treatment group) injected with captopril or losartan (daily IP injections of 50 and 30 mg/kg, respectively) 8 days on a LS KL diet followed by 5 days on a HS LK diet.

Table 1.

Effect of spironolactone on plasma electrolytes.

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