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Compromised regulation of the collecting duct ENaC activity in mice lacking AT_{1a} receptor

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Abstract

ENaC-mediated sodium reabsorption in the collecting duct (CD) is a critical determinant of urinary sodium excretion. Existing evidence suggest direct stimulatory actions of Angiotensin II (Ang II) on ENaC in the CD, independently of the aldosterone- mineralocorticoid receptor (MR) signaling. Deletion of the major renal AT₁ receptor isoform, AT_{1a}R, decreases blood pressure and reduces ENaC abundance despite elevated aldosterone levels. The mechanism of this insufficient compensation is not known. Here, we used patch clamp electrophysiology in freshly isolated splitopened CDs to investigate how AT_{1a}R dysfunction compromises functional ENaC activity and its regulation by dietary salt intake. Ang II had no effect on ENaC activity in CDs from $AT_{1a}R$ –/– mice suggesting no complementary contribution of AT_2 receptors. We next found that $AT_{1a}R$ deficient mice had lower ENaC activity when fed with low (<0.01% Na⁺) and regular (0.32% Na⁺) but not with high ($\sim 2\%$ Na⁺) salt diet, when compared to the respective values obtained in Wild type (WT) animals. Inhibition of AT₁R with losartan in wild-type animals reproduces the effects of genetic ablation of AT_{1a}R on ENaC activity arguing against contribution of developmental factors. Interestingly, manipulation with aldosterone-MR signaling via deoxycosterone acetate (DOCA) and spironolactone had much reduced influence on ENaC activity upon AT_{1a}R deletion. Consistently, $AT_{1a}R$ –/– mice have a markedly diminished MR abundance in cytosol. Overall, we conclude that $AT_{1a}R$ deficiency elicits a complex inhibitory effect on ENaC activity by attenuating ENaC Po and precluding adequate compensation via aldosterone cascade due to decreased MR availability.

Keywords

aldosterone; distal renal tubule; Mas; mineralocorticoid receptors; Na⁺ reabsorption

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1 INTRODUCTION

Sodium reabsorption by the kidney is central for maintaining sodium balance and setting blood pressure (Crowley & Coffman, 2012). Sodium transport in the distal renal tubule segments, including the collecting duct (CD), plays a major role in matching urinary salt excretion with day-to-day alterations in dietary NaCl intake (Pearce et al., 2015; Staruschenko, 2012). Activity of the epithelial Na⁺ channel (ENaC) on the apical plasma membrane of principal cells governs electrogenic Na⁺ influx at these sites (Pearce et al., 2015). Overactivation of ENaC contributes significantly to the pathology of salt- sensitive hypertension rodents and humans (Bhalla & Hallows, 2008; Hummler, 1999; Kakizoe et al., 2009). Likewise, mutations in genes encoding β - and γ -ENaC subunits resulting in gain-of-function of the channel underlie Liddle syndrome associated with high blood pressure (Hansson, Nelson-Williams, et al., 1995; Hansson, Schild, et al. 1995; Shimkets et al., 1994). Deficient ENaC activity leads to renal salt wasting and hypotension, as exemplified in patients with pseudohy- poladosteronism type I (Chang et al., 1996; Schild, 1996).

ENaC is a critical end-effector of the renin-angiotensin- aldosterone system (RAAS). This allows inverse relation between ENaC activity in the distal tubule and dietary salt intake in order to fine-tune systemic Na⁺ balance largely caused by respective changes in circulating levels of aldosterone and Angiotensin II (Ang II) (Crowley & Coffman, 2012; Zaika, Mamenko, Staruschenko, & Pochynyuk, 2013). Through its binding to the mineralocorticoid receptors (MR) in principal cells, the principal ENaC activator, aldosterone exerts a plethora of stimulatory actions on the channel, including augmented synthesis, trafficking to the apical plasma membrane, etc. (Eaton, Malik, SaxeNa, Al-Khalili, & Yue, 2001; Masilamani, Kim, Mitchell, Wade, & Knepper, 1999; Pacha, Frindt, Antonian, Silver, & Palmer, 1993; Stockand, 2002). In addition to promoting aldosterone secretion from the zoNa glomerulosa of adrenal gland, Ang II is capable of directly increasing ENaC activity non-redundantly to aldosterone (Mamenko, Zaika, Ilatovskaya, Staruschenko, & Pochynyuk, 2012; Peti-Peterdi, Warnock, & Bell, 2002; Sun, Yue, & Wang, 2012). The physiological actions of Ang II are mediated by ATi and AT₂ receptors (ATiR and AT₂R, respectively). AT₁R is a product of two separate genes encoding two subtypes, Namely AT_{1a}R and AT_{1b}R in rodents, with the former being the only isoform expressed in the kidney (Ito et al., 1995). AT_{1a}R activation causes antinatriuresis and vasoconstriction, whereas AT₂R antagonize many of the biological effects of AT1aR and cause vasodilation, Natriuresis, and prostaglandin release (Berry, Touyz, Dominiczak, Webb, & Johns, 2001; Kaschina & Unger, 2003). Both receptors are abundantly expressed in the renal tubule, although AT_{1a}R expression predominates (Carey, Wang, & Siragy, 2000; Miyata, Park, Li, & Cowley, 1999; Ozono et al., 1997).

Mice with global $AT_{1a}R$ deletion develop urinary salt wasting and have lower blood pressure at the baseline, which is further exacerbated during the state of dietary salt deficiency (Ito et al., 1995; Oliverio, Best, Smithies, & Coffman, 2000). Interestingly, $AT_{1a}R$ –/– mice have intact regulation of aldosterone secretion during volume depletion, most likely due to the abundant expression of $AT_{1b}R$ isoform in the adrenal gland (Burson, Aguilera, Gross, & Sigmund, 1994). Despite this, $AT_{1a}R$ deficiency has profound effects on ENaC subunit expression. Decreased a-subunit abundance is paralleled by increased β and β subunit expression upon salt restriction, while high salt intake lowers α -subunit levels without any

changes to β and γ abundance (Brooks, Allred, Beutler, Coffman, & Knepper, 2002). Moreover, mice with targeted deletion of AT_{1a}R in principal cells only, while having a normal blood pressure independently of salt intake, fail to increase αENaC expression during chronic Ang II infusion (Chen et al., 2016). It is unknown why aldosterone-MR signaling is not potent to correct ENaC expression in the absence of AT_{1a}R. Furthermore, since all three subunits are equally essential to assemble into a functional ENaC, it is hard to predict how discrete and opposite changes in expression of ENaC subunits are translated into changes in ENaC activity during the state of AT_{1a}R deficiency.

In this study, we undertook direct measurements of ENaC activity with patch clamp in freshly isolated split-opened CDs to investigate how disruption of the stimulatory Ang II input compromises channel function during adaptations to dietary salt intake. We found that direct Ang II actions on single channel ENaC activity depend solely on $AT_{1a}R$. Deletion of the receptor had a moderate inhibitory effect during regular salt intake and this defect was exacerbated during dietary salt deficiency. Changes in ENaC activity were much reduced in $AT_{1a}R$ –/– mice upon stimulation and inhibition of MR receptors with DOCA and spironolactone, respectively. Furthermore, MR availability was significantly reduced in mutant mice on both low and high salt intake providing a mechanism for insufficient compensation by aldosterone. Overall, our results demonstrate indispensable role of Ang II - $AT_{1a}R$ mechanisms for adequate regulation of ENaC activity in the CD by dietary salt intake.

2 | MATERIALS AND METHODS

2.1 | Reagents and animals

All chemicals and materials were from Sigma (St. Louis, MO), VWR (Radnor, PA), and Tocris (Ellisville, MO) unless noted otherwise and were at least of reagent grade. For experiments, male C57BL/6J mice (Charles River Laboratories, Wilmington, MA) and B6.129P2-*Agtr1^{t-m1Unc}*/j (AT_{1a}R –/–, JAX strain #002682) 6–10 weeks old, were used. Animal use and welfare adhered to the NIH Guide for the Care and Use of Laboratory Animals following protocols reviewed and approved by the Animal Care and Use Committees of the University of Texas Health Science Center at Houston.

2.2 | Research diets and treatments

To examine effects of salt intake, animals were provided diets containing nominally free (<0.01% Na⁺, TD.90228), regular (0.32% Na⁺, 7012), and high (-2% Na⁺, TD.92034) salt for 1 week. All diets were nutritionally balanced and were purchased from Envigo (Madison, WI). Spironolactone USP (30 mg/kgBW; Amneal Pharmaceutical) and losartan (10 mg/kgBW) were added to drinking water for 7 days. As necessary, mice were injected with Deoxycorticosterone acetate (DOCA) for three consecutive days (2.4 mg/injection/ animal) prior to the experimentation similarly to what we have done previously (Mamenko, Zaika, Doris, & Pochynyuk, 2012; Mamenko et al., 2013, 2017).

2.3 | Tissue isolation

The procedure for isolation of the CDs suitable for electrophysiology followed previously published protocols (Mamenko, Zaika, Doris, et al., 2012; Mamenko et al., 2013; Mironova, Bugay, Pochynyuk, Staruschenko, & Stockand, 2013; Prieto et al., 2017). Briefly, mice were sacrificed by CO₂ administration followed by cervical dislocation and kidneys were removed immediately. Kidneys were cut into thin slices (<1mm) with slices placed into icecold physiological saline solution (PSS) containing (in mmol/L) 150 NaCl, 5 KCl, 1 CaCl₂, 2 MgCl₂, 5 glucose and 10 HEPES (pH 7.35). CDs were visually identified by their morphological features (pale color; coarse surface and, in some cases, bifurcations) and were mechanically isolated from kidney slices by micro-dissection using watchmaker forceps under a stereomicroscope. Isolated CDs were attached to 5×5 mm cover glasses coated with poly-L-lysine. A cover-glass containing a CD was placed in a perfusion chamber mounted on an inverted Nikon Eclipse Ti microscope and perfused with PSS at room temperature. CDs were split-opened with two sharpened micropipettes, controlled with different micromanipulators, to gain access to the apical membrane. The tubules were used within 2 hr of isolation. For each experimental condition, CDs from at least four different mice were analyzed.

2.4 Single channel recordings in isolated collecting ducts

ENaC activity in principal cells was determined in cell-attached patches on the apical membrane made under voltage-clamp conditions ($-V_p = -60$ mV) using standard procedures (Mamenko, Zaika, Doris, et al., 2012; Mamenko, Zaika, Ilatovskaya, et al., 2012; Mamenko et al., 2013; Prieto et al., 2017). Current recordings were made in a permanently perfused bath (1.5 ml/min). Recording pipettes had resistances of 8-10 megaOhms. Typical bath and pipette solutions were (in mmol/L): 150 NaCl, 5 mM KCl, 1 CaCl₂, 2 MgCl₂, 5 glucose and 10 HEPES (pH 7.35); and 140 LiCl, 2 MgCl₂ and 10 HEPES (pH 7.35), respectively. Gap-free single channel current data from gigaOhm seals were acquired and analyzed with Axopatch 200B (Molecular Devices, Sunnyvale, CA) patch clamp amplifier interfaced via a Digidata 1440 (Molecular Devices) to a PC running the pClamp 10.4 suite of software (Molecular Devices). Currents were low-pass filtered at 100 Hz with an eightpole Bessel filter (Warner Instruments, Hamden, CT). Events were inspected visually prior to acceptance. ENaC activity was analyzed over a span of 60-120 s for each experimental condition. Using previously described analysis (Mamenko, Zaika, Ilatovskaya, et al., 2012), we can reliably (p < 0.05) estimate the maximal number of functional ENaC in a patch using this time span. Channel activity in individual patches, defined as NP₀, was calculated using the following equation: NP_o = $(t_1 + 2t_2 + ... + nt_p)$, where N and P_o are the number of ENaC in a patch and the mean open probability of these channels, respectively, and t_n is the fractional open time spent at each of the observed current levels. P_0 was calculated by dividing NP_o by the number of active channels within a patch as defined by all-point amplitude histograms. To estimate total ENaC activity (fNP₀) under a particular physiological condition, we normalized NP_{o} to the frequency of observing patches with at least one active channel (f = number of patches with active channels/total number of patches). To assess functional ENaC expression for each experimental condition, the mean of number of active channels within a patch was corrected to f. For representation, current traces were corrected for a slow baseline drifts as necessary.

2.5 | Western blotting

Immediately after dissection, kidneys were frozen in liquid nitrogen and stored there for further use. Prior to experimentation the kidneys were homogenized in three volumes of icecold hypotonic lysis buffer containing 50 mmol/L Tris, 1% Triton X-100, 5 mmol/L EDTA (pH = 7.4) supplemented with 1mM PMSF and 2 mg/ml protease inhibitor cocktail (Complete mini, Roche Diagnostics, Germany). Homogenates were centrifuged at 1,000g for 15 min at +4C to separate cytosolic fraction. Protein concentration in the Homogenates was determined with a Bradford assay. The samples (25 µg/lane) were separated on 9% polyacrylamide gels at 150 V for 1hr 15 min and transferred to nitrocellulose membranes at 100 V for 1 hr 45 min. The membranes were blocked for 1 hr at RT in 5% nonfat milk in TBS-T (150 mmol/L NaCl, 50mmol/L Tris-HCl pH = 7.4, 0.1% Tween 20). Subsequently the membranes were probed with anti-MR primary rabbit antibodies (1:1000, ab64457, Abcam, Cambridge, UK) followed by peroxidase-conjugated goat anti-rabbit secondary antibodies (1:20000, Bio-Rad, Hercules, CA) for 1 hr at RT. The membranes were re-probed with anti-β-actin (1:5000, ab8227, Abcam) primary rabbit antibodies and peroxidaseconjugated goat anti-rabbit secondary antibodies (1:20000, Bio-Rad). When total MR expression was assessed, MR band intensities were normalized to the intensities of the corresponding β -actin bands.

2.6 | Statistics

All summarized data are reported as mean \pm SEM. All statistical comparisons were made using one-way ANOVA. A *p*-value of less than 0.05 was considered significant.

3 | RESULTS

3.1 | AT_{1a} receptors (AT_{1a}R) are central in regulation of ENaC activity by Angiotensin cascade

We and others reported previously that Ang II exhibits non-redundant stimulatory actions on ENaC activity beyond those elicited by the classical aldosterone-MR cascade (Mamenko, Zaika, Ilatovskaya, et al., 2012; Mamenko et al., 2013). Consistently, acute application of Ang II (500 nM) increased single channel ENaC activity in freshly isolated split-opened CDs from WT mice (Figure 1a, upper trace). Figure 1b contains the summary graph of stimulatory Ang II actions on ENaC in paired patch clamp experiments. We further took advantage of mice lacking AT_{1a} receptors, as a model of the defective Ang II signaling. Ang II (500 nM) had no measurable effect on single channel ENaC activity in freshly isolated split-opened CDs, as shown on a representative continuous patch clamp experiment in Figure 1a (bottom trace) and the summary graph in Figure 1b. This also argues against a complementary role of AT₂R in controlling ENaC function by Ang II cascade. Furthermore, application of Ang 1–7 (500 nM) did not affect ENaC activity in Wild type (WT) mice, as shown in the summary graph in Figure 1c suggesting no contribution of the alternative (vasoprotective) branch of the RAAS and specifically Mas receptors in regulation of ENaCmediated Na⁺ reabsorption in the CD. Overall, our results in Figure 1 provide a direct evidence of the exclusive role of AT_{1a}R in conveying stimulatory Ang II signal to ENaC.

3.2 | Decreased basal ENaC activity and blunted response to dietary salt restriction in $AT_{1a}R_{-}/_{-}$ mice

We next investigated how disruption of Ang II signal to ENaC in AT_{1a}R -/- mice affects channel function upon variations in dietary salt intake. As shown in representative current traces in Figure 2 (top panels) and the summary graph in Figure 3a, ENaC activity in split-opened CDs is significantly lower in mutant mice kept on regular (0.32% Na⁺) salt intake. Detailed analysis of single channel ENaC activity revealed that this difference was due to reduced open probability (P_o , Figure 3b), while the average number of active channels per patch (fN) remained similar (Figure 3c).

We further compared ENaC activity in $AT_{1a}R$ –/– and WT mice subjected to a low (<0.01% Na⁺) and high (~2% Na⁺) salt intake for one week. Dietary sodium restriction increased ENaC activity in both animal strains (Figure 2, bottom panel). However, mutant mice had consistently lower total ENaC activity (Figure 3A) and P_o (Figure 3b) compared to the respective values in WT mice under these conditions. In contrast, dietary salt loading negated all differences in ENaC activity and open probability between $AT_{1a}R$ –/– and WT mice (Figure 3). Overall, we concluded that disruption of Ang II signaling blunts regulation of ENaC activity in the CD by dietary salt intake due to significantly lower P_o changes.

3.3 | Inhibition of AT₁R with losartan decreases ENaC activity similarly to the receptor deletion

We next compared the effects of $AT_{1a}R$ deletion and receptor antagonism in WT mice on ENaC function to exclude putative compensatory effects due to chronic gene editing. For this, WT animals were fed with low, regular and high salt diets and concomitantly supplied with AT_1R blocker losartan (10 mg/kgBW) in drinking water. We similarly observed a reduced ENaC activity in animals kept on regular and low but not high salt intake (Figure 4a). The effect was attributed again to the reduced P_o (Figure 4b), the number of functional ENaC channels was moderately lower in mice on the regular salt regimen (Figure 4c). Overall, we concluded the $AT_{1a}R$ deletion and AT_1R blockade elicit comparable reduction of ENaC activity suggesting minimal compensation. In addition, Figures 3 and 4 suggest a critical role of Ang II signaling in controlling ENaC activity during regular and low salt intake.

3.4 | AT_{1a}R deletion compromises regulation of ENaC by aldosterone-MR cascade

We next tested the effect of maximal activation of MR receptors after repetitive DOCA injections on ENaC function in WT and $AT_{1a}R$ –/– mice. As expected, we detected a marked increase in (fNP_o) from 0.38 ± 0.04 to 0.71 ± 0.11 in WT, which was blunted in $AT_{1a}R$ –/– mice: 0.25 ± 0.03 to 0.29 ± 0.04 , not statistically significant (Figure 5a). We further probed basal activity of MR-aldosterone axis in $AT_{1a}R$ knockouts with MR antagonist, spironolactone (30 mg/kgBW). As summarized in Figure 5b, spironolactone failed to produce a significant inhibitory effect on ENaC fNP_o under regular salt intake decreasing total channel activity from 0.26 ± 0.03 to 0.18 ± 0.05 . Impaired response of MR signaling cascade to DOCA despite its low basal activity is indicative of reduced MR abundance. Indeed, Western blot analysis detected a markedly reduced expression of cytosolic MR in $AT_{1a}R$ –/– kidney Homogenates, as compared to the WT mice on both low

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salt (<0.01% Na⁺) and high salt (~2% Na⁺) diets (Figure 6a). At the same time, we observed an upregulation of MR levels upon dietary salt restriction in both strains (Figure 6b). This is consistent with a greater inhibitory effect of spironolactone on single channel ENaC activity in AT_{1a}R –/– mice kept on a low salt diet, where ENaC activity was significantly diminished from 0.52 ± 0.08 to 0.31 ± 0.08 (Figure 5b). Overall, we concluded that AT_{1a}R deletion compromises regulation of ENaC by aldosterone cascade during variations in dietary salt intake at least partially due to reduction in MR availability in the cytosol for stimulation by aldosterone.

4 | DISCUSSION

In this study, we employed $AT_{1a}R$ –/– mice as a model deficient of Ang II regulation of ENaC to investigate how this affects regulation of channel activity by dietary salt intake and aldosterone-MR signaling. We report here that $AT_{1a}R$ deletion diminishes ENaC activity at the baseline and blunts upregulation of channel activity by low salt diet. Detailed patch clamp analysis in split-opened CDs revealed that the defect was primarily attributed to altered ENaC gating, Namely decreased open probability, P_{o} . In addition, we found that the lack of $AT_{1a}R$ also interferes with stimulatory input to ENaC from aldosterone-MR cascade.

Using direct measurements of single channel activity, we found that AT_{1a}R ablation precludes regulation of ENaC by Ang II (Figures 1a and 1b). In addition to promoting its anti-Natriuretic actions via AT1R, Ang II is also a potent ligand for AT2R, which generally oppose the effects mediated by AT₁R (Berry et al., 2001; Kaschina & Unger, 2003). AT₂R deficiency exacerbates Ang II-dependent antinatriuresis (Siragy, Inagami, Ichiki, & Carey, 1999), suggesting AT_2R effects of tubular sodium reabsorption. However, this seems to be associated rather with effects in proximal tubule having high levels of AT₂R and specifically the sodium hydrogen antiporter 3, NHE-3 and Na⁺, K⁺ ATPase (Carey, 2017). We and others showed a dominant role of AT₁Rin regulation of sodium reabsorption in the distal tubule, whereas application of AT₂R agonist, CGP42112 had no effect on ENaC (Mamenko, Zaika, Ilatovskaya, et al., 2012; Peti-Peterdi et al., 2002; Sun et al., 2012). This is in agreement with the current finding that Ang II has neither stimulatory nor inhibitory effects on ENaC, when $AT_{1a}R$ are deleted (Figure 1). It should be noted though that we used exclusively males in our patch clamp studies. Substantial differences in expression/function of distal tubule sodium transporters between males and females at the baseline as well as in response to salt loading have been reported (Gillis & Sullivan, 2016; Veiras et al., 2017). Specifically, female rats have increased thiazide-sensitive sodium chloride cotransporter (NCC) and ENaC abundance to partially compensate for the diminished volume reabsorption in the proximal tubule (Veiras et al., 2017). While these differences were more pronounced in rats than in mice, further careful examination of sex polymorphism in Ang II-dependent regulation of ENaC is granted. In this regard, AT₂R expression and its hypotensive effects are more pronounced in females (Hilliard et al., 2011).

Recent studies also provided a circumferential evidence of a stronger hypotensive and Natriuretic effect of azilsartan, an AT₁R antagonist that was linked to augmented renal ACE2 mRNA and reduced α ENaC pointing to an upregulation of the alternative (vasoprotective) branch of the renin-angiotensin-aldosterone system, the ACE2/Ang-(1–

7)/Mas cascade in addition to AT_1R blockade after antagonist treatment (Iwanami et al., 2014). To our disappointment, we did not observe a measurable inhibitory effect of Ang 1–7 in split- opened CDs (Figure 1C) suggesting no apparent role of Mas receptors in regulation of ENaC activity in the distal tubule at least in mice on a regular salt intake.

We found that deletion of AT_{1a}R blunts salt-sensitivity of ENaC in the CD, explicitly, the dynamic range in which ENaC activity is changed during variations of dietary NaCl from high to low (Figures 2 and 3). ENaC activity was similar between salt loaded Wild type and mutant mice when circulating Ang II levels are low, whereas channel activity was significantly reduced in AT_{1a}R -/- mice during dietary salt restriction (associated with high Ang II levels) indicating lower CD sodium reabsorption and higher urinary sodium excretion in this case. This could potentially contribute to the worsened hypotensive phenotype in these mice on low NaCl diet (Oliverio et al., 2000). On the contrary, Coffman's group reported recently that mice with targeted deletion of AT_{1a}R in principal cells, while exhibiting a blunted hypertensive response to Ang II infusions, have blood pressure indistinguishable from controls on all tested salt intakes (Chen et al., 2016). Since the employed cre-lox strategy does not guarantee the complete receptor ablation and no positive functional control was provided, it is possible that the residual receptor expression is sufficient to modulate ENaC activity during variations in dietary salt intake. In contrast, we showed no effect of Ang II on ENaC justifying no functional AT_{1a}R in the current mutant model. Alternatively, intact AT_{1a}Rin intercalated cells of the CD might play a compensatory role by stimulating electroneutral NaCl reabsorption (Leviel et al., 2010) upon AT_{1a}R deletion in principal cells.

We also report here that the observed defect in ENaC activity in the CD upon deletion of $AT_{1a}R$ is due to decreased channel open probability on regular and low salt intake (Figures 2 and 3). This is consistent with previous observations that Ang II is potent to increase ENaC P_{o} and this effect persists in DOCA- and spironolactone-treated animals, suggesting its aldosterone-independent Nature (Mamenko, Zaika, Ilatovskaya, et al., 2012; Sun et al., 2012). We also previously showed that this Ang II-dependent regulation of ENaC P_o may contribute to regulation of channel activity by dietary salt intake. Thus, inhibition of MR signaling with spironolactone precluded salt- dependent regulation of the number of active ENaC measured with patch clamp in split-opened CDs, but failed to affect changes in open probability (Mamenko et al., 2013). In agreement, we observe that the changes in P_0 are substantially reduced in $AT_{1a}R$ knockouts, while we did not detect significant changes in the average number of active channels per patch between mutant and Wild type mice (Figure 3c). Recall, $AT_{1a}R$ –/– animals retain the ability to increase aldosterone secretion in response to volume depletion most likely due to contribution of unaffected AT_{1b}R isoform in the adrenal gland (Burson et al., 1994). Interestingly, Western blot detection of ENaC in the kidney of $AT_{1a}R$ –/– mice showed reduced abundance of α , but increased abundance of cleaved, most active form of y (Brooks et al., 2002). Since 1:1:1 stoichiometry for $\alpha:\beta:\gamma$ is required to assemble into a functional ENaC (Stockand, Staruschenko, Pochynyuk, Booth, & Silverthorn, 2008), this should lead to the overall reduction in the channel numbers with a scarcity being a limiting factor. Since we did not observe differences in the number of active channels between WT and mutant mice, the most likely explanation is increased channel residence on the apical plasma membrane in $AT_{1a}R$ –/– driven by actions of aldosterone.

Indeed, we previously observed cumulative inhibitory effects of MR inhibitor spironolactone and AT₁R blocker losartan on ENaC activity in the CD (Mamenko et al., 2013). We also found that inhibition of AT₁R with losartan recapitulated the major phenotype with regard to decreased ENaC activity in the CD (Figure 4). This argues against major contribution of the developmental factors associated with receptor deficiency. This is consistent with a previous report showing no significant differences in body and kidney weights in AT_{1a}R –/– and Wild type mice suggesting that the receptor is not essential for the normal organogenesis of the kidney (Oliverio et al., 1998).

One of the important results is that $AT_{1a}R$ deficiency cannot be properly compensated by aldosterone-MR cascade. We show that maximal stimulation of MR with DOCA has little stimulatory effect on ENaC activity in $AT_{1a}R$ –/– mice (Figure 5a). This is consistent with reduced availability of the receptors in cytosol, detected by Western blot (Figure 6). While to a much lower extent, dietary salt restriction increases the number of available MR in mutants resulting in the upregulation of ENaC activity. This observation is corroborated by a greater effect of MR inhibitor spironolactone on ENaC activity during this condition (Figure 5b). While the exact mechanism of this MR deficiency requires further investigation beyond the scope of the current manuscript, Ang II was shown to potentiate aldosterone binding to MR in intercalated cells of the CD by removing a specific S843 phosphorylation precluding ligand binding (Shibata et al., 2013). This was proposed to account for discrete adaptive responses to hypokalemia (when only aldosterone is present) and hypovolemia (both aldosterone and Ang II are elevated). Furthermore, inhibition of Ang II pathway with angiotensin converting enzyme (ACE) inhibitor, captopril reduced cardiac fibrosis not only by decreasing Ang II production but also by attenuating the aldosterone-signaling pathway by decreasing the expression of MR receptors (de Resende, Kauser, & Mill, 2006). In summary, our findings support the overarching view that cooperative but yet non-redundant actions of aldosterone and Ang II signaling cascades are necessary to regulate electrolyte transport in principal and intercalated cells of the CD during variations in dietary intake. With respect to ENaC function, deficiency of Ang II signal cannot be properly compensated by aldosterone-MR cascade, as demonstrated in the current study.

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FIGURE 1.

AT 1aRs are central for regulation of ENaC activity in the CD by Angiotensin II. (a) Representative continuous current traces from cell-attached patches monitoring ENaC activity in the control and during application of 500 nM Ang II (shown with a bar on the top) in a split-opened CD from WT (upper trace) and AT1aR -/- (bottom trace) mice. The patches were held at a test potential of $V_h = -V_p = -60$ mV. Inward currents are downward. Dashed lines indicate the respective current state with a c denoting the closed state. (b) Summary graph of ENaC Po changes in response to Ang II from paired patch clamp experiments in WT and AT1aR -/- mice similar to that shown in Figure 1a.*significant increase versus WT control. (c) Summary graph of ENaC Po changes in response to stimulation of Mas receptors with 500 nM Ang 1–7 in split-opened CDs from Wild type mice



FIGURE 2.

ENaC activity is reduced in AT_{1a}R -/- mice. Representative current traces of ENaC activity in split-opened CDs isolated from Wild type (left column) and AT_{1a}R -/- (right column) mice kept on control (0.32% Na⁺, top row) and low (<0.01% Na⁺, bottom row) salt intake for 1 week. Dashed lines indicate the respective current state with o_i indicating the number of simultaneously opened channels; *c* denotes the closed non-conducting state

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FIGURE 3.

AT_{1a}R –/– deletion reduces ENaC salt-sensitivity. Summary graphs of averaged total ENaC activity, fNP_o (a); ENaC open probability, P_o (b); and number of active channels fN (c) in wild type (light grey) and AT_{1a}R –/– (dark gray) mice kept on low (<0.01% Na⁺), control (0.32% Na⁺), and high (~2% Na⁺) salt diet for one week, respectively. At least four different animals were tested for each condition. *significant decrease versus respective wild type values

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FIGURE 4.

Inhibition of AT1R with losartan reduces ENaC activity on low and control salt intake. Summary graphs of averaged total ENaC activity, fNP_o (a); ENaC open probability, P_o (b); and number of active channels fN (c) in Wild type mice kept on low (<0.01% Na⁺), control (0.32% Na⁺), and high (~2% Na⁺) salt diets for one week in the absence (light gray) and presence (dark gray) of concomitant treatment with AT₁R antagonist losartan (10 mg/kgBW) in drinking water, respectively. At least four different animals were tested for each condition. *significant decrease versus respective untreated conditions



FIGURE 5.

AT1aR deletion impairs regulation of ENaC activity by mineralocorticoid receptor (MR) signaling. (a) Summary graph of averaged total ENaC activity, fNP_o in Wild type and AT_{1a}R -/- mice in the control and after systemic Deoxycorticosterone acetate (DOCA) injections for 3 consecutive days to maximally stimulate MR. *significant increase versus Wild type control; **significant decrease versus Wild type + DOCA. (b) Summary graph comparing averaged total ENaC activity, fNP_o in AT1aR -/- mice kept on low (<0.01% Na⁺) and control (0.32% Na⁺) salt intake in the absence (light gray) and presence (dark grey) of concomitant treatment with MR antagonist spironolactone (30 mg/kgBW) in drinking water, respectively. *significant decrease versus AT_{1a}R -/- low salt



FIGURE 6.

Decreased MR availability in $AT_{1a}R$ –/– mice during different salt intake. (a) Representative Western blot from whole kidney lysates (cytosolic fraction) of WT and $AT_{1a}R$ –/– mice (each line reflects individual animal) kept on low (<0.01% Na⁺) and high (~2% Na⁺) salt intake for 1 week simultaneously probed with anti-MR and anti- β actin antibodies. (b) Summary graph comparing MR expression normalized to the respective intensity of β -actin in WT and $AT_{1a}R$ –/– mice on the same conditions as described in Figure 6a. *significant decrease versus respective Wild type values