Supplemental data

Plasma Potassium determines NCC Abundance in Adult Kidney-specific γENaC Knockout

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Supplemental Figure 1 Schema of the experimental protocols. Representative setup used to determine physiological parameters in control and $\gamma ENaC$ -deficient (Scnn1g^{Pax8/LC1}) mice placed either in standard (thin line) or metabolic cages (bold line) and induced by doxycycline (Dox) under standard diet (**A**), a short term high Na⁺ and low K⁺ diet (**B**), a short term low K⁺ diet (**C**), a K⁺-deficient diet +/- supplemented with a K⁺ chelator (kayexalate) (**D**), and (**E**) following a 6 days treatment with a K⁺-deficient diet at day 0 of Dox induction. Arrows indicate the end of the experiment, and the time point of serum and organ recovery.

Supplemental Figure 2 Physiological parameters measured following induction under standard diet in control and γ ENaC-deficient mice. Measurement of daily 24 hours (A) food (g/24h/gBW), and (B) water intake (ml/24h/g BW), (C) feces (g/24h/gBW), and (D) urine volume output (ml/24h/gBW), (E) Na⁺ and (F) K⁺ intake (mmol/24/gBW) in control and experimental (Scnn1g^{Pax8/LC1}) (each group, n=5) mice; *protocol A*. Values were normalized to the body weight. **P<0.01, ***P<0.001.

Supplemental Figure 3 Acute high Na⁺ and low K⁺ diets do not restore food and water intake, feces output and urinary volume excretion in γ ENaC-deficient mice. (A) Food intake (g/gBW), (B) water intake (ml/gBW), (C) feces output (g/gBW), and (D) urine volume (ml/gBW), in control (n=8) and experimental (Scnn1g^{Pax8/LC1}) (n=9) mice following doxycycline treatment upon standard diet (24-hour measurement) and high Na⁺/low K⁺ (6-hour measurement); *protocol B*. Values were normalized to the body weight. *P<0.05, **P<0.01, ***P<0.001.

Supplemental Figure 4 Acute low K⁺ diet does not normalize food and feces weight in γ ENaC-deficient mice. (A) Food (g/24h/gBW) and (B) water intake (ml/24h/gBW), (C) feces output (g/24h/gBW), and (D) urine volume (ml/24h/gBW) in control (n=6) and experimental mice (Scnn1g^{Pax8/LC1}) (n=6; *protocol C*). Values were taken normalized to the body weight. *P<0.05.

Supplemental Figure 5 No difference in protein abundances of NCC, pNCC, SPAK, and pSPAK in γ ENaC-deficient mice before Dox induction. Representative Western blot analysis (A) of NCC and pNCC, (D) SPAK and pSPAK on whole kidney protein extracts from Scnn1g control and deficient mice upon standard diet (each group, n=4). (B, E) Quantification of proteins and (C, F) ratio of (C) pNCC to total NCC abundances and (F) pSPAK to total SPAK abundances from the corresponding Western blot analyses. Protein levels were normalized to actin and expressed in percentage of control. Results are presented as mean \pm SEM and data were analyzed by unpaired t test. P values <0.05 were considered statistically significant.

Supplemental Figure 6 Scnn1g^{Pax8/LC1} mice stabilize their body weight upon K⁺deficient diet supplemented with sodium kayexalate. Body weight changes (Δ body weight, BW) of Scnn1g control treated with kayexalate (n=6) or without (n=3) and experimental mice treated with kayexalate (Scnn1g^{Pax8/LC1}, n=7) in percentage of initial BW upon standard Na⁺ and K⁺-deficient diet ± kayexalate following Dox induction (at day 0, *protocol D*). Results are presented as mean ± SEM and data were analyzed by unpaired t test. P values <0.05 were considered statistically significant. *P<0.05, **P<0.01, ***P<0.001. Supplemental Figure 7 Adult γ ENaC-deficient mice still present salt wasting under K⁺deficient diet with kayexalate treatment. (A) Body weight changes (Δ body weight, BW) of Scnn1g control treated (Scnn1g control with kayexalate, n=6) or without kayexalate (Scnn1g control without kayexalate, n=3) and experimental mice treated with kayexalate (Scnn1g^{Pax8/LC1} with kayexalate, n=7) in percentage of initial BW upon K⁺-deficient diet (*protocol D*). 24 hours urinary (B) Na⁺ and (C) K⁺ excretion (mmol/gBW), and 24 hours (D) Na⁺ and (E) K⁺ intake (mmol/gBW), (F) 24 hours food intake (g/gBW) and (G) water intake (ml//gBW), (H) feces output (g/gBW), and (I) urine volume (ml/gBW), in control without kayexalate (n=3) and with kayexalate (n=6), and experimental mice with kayexalate (n=7) (Scnn1g^{Pax8/LC1}) following doxycycline treatment and upon K⁺-deficient diet; *protocol D*. Values were normalized to the body weight. *P<0.05, **P<0.01, ***P<0.001.

Supplemental Figure 8 γ ENaC-deficient mice normalize plasma sodium and potassium concentration under K⁺-deficient diet with kayexalate treatment. Measurements of plasma (A) Na⁺ and (B) K⁺ concentrations (mmol/l) in Scnn1g control without (n=3) and with kayexalate (n=6) and experimental Scnn1g^{Pax8/LC1} mice with kayexalate (n=7) following doxycycline treatment upon K⁺-deficient diet; *protocol D*. Representative Western blot analyses of (C) total and (D) phosphorylated NCC and actin on kidney from control without (n=3) and with kayexalate (n=4) and experimental Scnn1g mice treated with kayexalate (n=4) upon K⁺-deficient diet (*protocol D*). (E and F) Quantification of proteins and (G) ratio of pNCC to total NCC abundances from corresponding Western blot analyses. Protein levels were normalized to actin and expressed in percentage of control. Results are presented as mean \pm SEM and data were analyzed by unpaired t test. P values <0.05 were considered statistically significant, *P<0.05, **P<0.01, ***P<0.001. Supplemental Figure 9 Increased urinary sodium loss in γ ENaC-deficient mice treated with K⁺-deficient diet supplemented with kayexalate. 24 hours fecal (A) sodium and (B) potassium concentration and (C) sodium and (D) potassium balance in Scnn1g control (n=6) and experimental (Scnn1g^{Pax8/LC1}) mice (n=7) following 20 day of doxycycline treatment and upon K⁺ -deficient diet supplemented with kayexalate; *protocol D*. The mean sodium or potassium intake (gray column) is compared with both urinary (white) and fecal (black column) sodium and potassium loss on K⁺-deficient diet complemented with kayexalate; *protocol D*. ***P<0.001.

Supplemental Figure 10 Loss of apical α and β ENaC subunit localization in cortical collecting duct cells from γ ENaC-deficient mice. Immunofluorescence detection of α -, β -, and γ ENaC in consecutive kidney sections from control or knockout (Scnn1g^{Pax8/LC1}) mice following two days of standard diet; *protocol A*; filled arrows, residual γ ENaC expressing cell that exhibits a apical membrane trafficking of the α - and β ENaC subunit (filled arrows) and of cells that exhibit a disturbed α ENaC subunit expression (open arrows).















Supplemental Figure 8



Supplemental Figure 9



