

**Deletion of the serine protease CAP2/*Tmprss4* leads to dysregulated renal water
handling upon dietary potassium depletion**

Anna Keppner^{1,2,6}, Darko Maric^{1,2,6}, Chloé Sergi¹, Camille Ansermet¹, Damien De Bellis^{1,7,8},
Denise V. Kratschmar^{3,6}, Jérémie Canonica^{1,6,9}, Petra Klusonova^{3,6}, Robert A. Fenton⁴, Alex
Odermatt^{3,6}, Gilles Crambert⁵, David Hoogewijs^{2,6}, Edith Hummler^{1,6}

Supplementary Information

Figure S1. The daily urinary Na⁺ and K⁺ excretions are only moderately changed in CAP2/*Tmprss4* and nephron-specific GR knockout mice.

Figure S2. mRNA transcript expression of K⁺ transporting systems is unchanged in CAP2/*Tmprss4* knockout mice.

Figure S3. NKCC2, AQP3, AQP4 and AC6 mRNA expressions are modified in CAP2/*Tmprss4* knockout mice, independently of AVP and its receptors.

Figure S4. mRNA transcript expression of Na⁺ transporting systems is unchanged in CAP2/*Tmprss4* knockout mice.

Figure S5. Na⁺,K⁺-ATPase and total and phosphorylated NCC are unchanged in CAP2/*Tmprss4* knockout mice.

Figure S6. Plasma hormone levels and 11β-HSD2 mRNA expression in CAP2/*Tmprss4* knockout mice.

Figure S7. Protein expression of steroid hormone receptors in CAP2/*Tmprss4* knockout mice.

Tables S1-S4. Daily urinary Na⁺ and K⁺ excretions of CAP2/*Tmprss4* and Nr3c1^{Pax8/LC1} mice.

Uncropped Western blots parts 1 to 4

Supplementary figure legends

Figure S1. The daily urinary Na⁺ and K⁺ excretions are only moderately changed in CAP2/Tmprss4 and nephron-specific GR knockout mice. Daily urinary Na⁺ (A) and K⁺ (B) excretions (mmol/day) in CAP2/Tmprss4 wildtype (+/+), heterozygous mutant (Δ /+), and knockout (Δ / Δ) mice (n=6 per genotype) following 2 days under regular K⁺ and 4 days under low K⁺ diet. Daily urinary Na⁺ (C) and K⁺ (D) excretions (mmol/day) in control (white circles) and Nr3c1^{Pax8/LC1} (black circles) mice (n=4-6 per genotype) following 2 days under regular K⁺ and 4 days under low K⁺ diet. (n=4-6 per genotype). * p < 0.05.

Figure S2. mRNA transcript expression of K⁺ transporting systems is unchanged in CAP2/Tmprss4 knockout mice. (A-E) Relative mRNA transcript expression levels in kidneys from CAP2/Tmprss4 wildtype (+/+), heterozygous mutant (Δ /+), and knockout (Δ / Δ) mice (n=6 per genotype) under low K⁺ diet of (A) H⁺,K⁺-ATPase type 1 (HKA1), (B) renal outer medullary K⁺ channel 1 (ROMK1), (C) ROMK2, (D) big K⁺ channel 1 (BK β 1), (E) BK β 4, (F) vacuolar H⁺-ATPase subunit a4 (ATP6V0A4), and (G) subunit b1 (ATP6V1B1).

Figure S3. NKCC2, AQP3, AQP4 and AC6 mRNA expressions are modified in CAP2/Tmprss4 knockout mice, independently of AVP and its receptors. Relative mRNA transcript expression levels of (A) NKCC2, (B) AQP3 and AQP4, and (C) adenylate cyclase 5 (AC5), 6 (AC6) and the soluble form (sAC) in kidneys from CAP2/Tmprss4 wildtype (+/+), heterozygous mutant (Δ /+), and knockout (Δ / Δ) mice (n=4) under low K⁺ diet. (D) Plasma copeptin levels in CAP2/Tmprss4 wildtype (+/+), heterozygous (Δ /+), and knockout (Δ / Δ) under regular K⁺

diet (RK) and low K⁺ diet (LK) (n=6 per genotype). Relative mRNA expression levels of (E) vasopressin receptor 1a (*avpr1a*), and (F) *avpr2* in kidneys from *CAP2/Tmprss4* wildtype (+/+), white circles), heterozygous mutant (Δ /+, grey circles), and knockout (Δ / Δ , black circles) mice (n=4 per genotype) under low K⁺ diet. * p<0.05.

Figure S4. mRNA transcript expression of Na⁺ transporting systems is unchanged in *CAP2/Tmprss4* knockout mice. Relative mRNA transcript expression levels of (A) α -ENaC (*Scnn1a*), (B) β -ENaC (*Scnn1b*), and (C) γ -ENaC (*Scnn1g*) in kidneys from *CAP2/Tmprss4* wildtype (+/+, white circles), heterozygous mutant (Δ /+, grey circles), and knockout (Δ / Δ , black circles) mice (n=3) under low K⁺ diet. (D) Representative (cropped) immunoblots for α -ENaC (*Scnn1a*), β -ENaC (*Scnn1b*) and γ -ENaC (*Scnn1g*) in kidney lysates from *CAP2/Tmprss4* wildtype (+/+, white circles), heterozygous mutant (Δ /+, grey circles), and knockout (Δ / Δ , black circles) mice (n=3-4) under low K⁺ diet, and (E) corresponding protein quantifications for full-length (F) and cleaved (C) subunits. Actin was used as loading control. The membranes were first blotted against the different ENaC subunits, then stripped and blotted against actin. (F) Representative immunoblot for Na⁺-H⁺ exchangers 1 and 3 (NHE1 and NHE3) in kidney lysates from *CAP2/Tmprss4* wildtype (+/+) and knockout (Δ / Δ) mice under low K⁺ diet (n=4-6 per genotype). Actin was used as loading control. The membranes were cut, the lower part was blotted against actin, while the upper part was first blotted against NHE1, stripped, and then blotted against NHE3. (G,H) Corresponding protein quantification of (G) NHE1, and (H) NHE3. Full-length blots can be found at the end of this supplementary information file.

Figure S5. Na⁺,K⁺-ATPase and total and phosphorylated NCC are unchanged in *CAP2/Tmprss4* knockout mice. (A) Representative (cropped) immunoblots for Na⁺,K⁺-ATPase (NKA), NCC, and pT53-NCC in kidney lysates from *CAP2/Tmprss4* wildtype (+/+),

white circles), heterozygous mutant ($\Delta/+$, grey circles), and knockout (Δ/Δ , black circles) mice under low K^+ diet (n=3 per genotype). Actin was used as loading control. For NKA, the membrane was cut, the lower part was blotted against actin, while the upper part was blotted against NKA. For NCC and pT52-NCC, the membrane was cut, the lower part was blotted against actin, while the upper part was first blotted against pT53-NCC, stripped, and then blotted against NHE3. **(B-C)** Corresponding protein quantification of **(B)** NKA and **(C)** NCC and pT53-NCC. **(D-E)** Relative mRNA transcript expression levels of **(D)** WNK lysine deficient protein kinase 4 (WNK4), and **(E)** glucocorticoid-induced leucine zipper (GILZ) in kidneys from *CAP2/Tmprss4* wildtype (+/+, white circles), heterozygous mutant ($\Delta/+$, grey circles), and knockout (Δ/Δ , black circles) mice (n=3-4 per genotype) under low K^+ diet. Full-length blots can be found at the end of this supplementary information file.

Figure S6. Plasma hormone levels and 11 β -HSD2 mRNA expression in *CAP2/Tmprss4* knockout mice. Plasma levels (nM) of **(A)** progesterone, **(B)** testosterone, **(C)** androstenedione, **(D)** 11-dehydrocorticosterone, and **(E)** 11-deoxycorticosterone in *CAP2/Tmprss4* wildtype (+/+, white circles, n=6), heterozygous mutant ($\Delta/+$, grey circles, n=5), and knockout (Δ/Δ , black circles, n=5) mice under regular K^+ (RK) and low K^+ diet (LK). **(F)** Relative 11 β -HSD2 mRNA expression in *CAP2/Tmprss4* wildtype (+/+, white circles, n=6), heterozygous mutant ($\Delta/+$, grey circles, n=5), and knockout (Δ/Δ , black circles, n=5) kidneys under low K^+ diet. * p< 0.05, ** p< 0.01.

Figure S7. Protein expression of steroid hormone receptors in *CAP2/Tmprss4* knockout mice. **(A)** Representative (cropped) immunoblots for progesterone receptor α and β (PR α and PR β), androgen receptor (AR), and the different estrogen receptor α isoforms (ER iso1 55kDa, iso2 51kDa, iso3 46kDa, and iso4 35kDa) in kidney lysates from *CAP2/Tmprss4* wildtype (+/+),

white circles) and knockout (Δ/Δ , black circles) mice (n=4 per genotype) under low K⁺ diet; actin was used as loading control. **(B-D)** Corresponding protein quantifications of **(B)** PR, **(C)** AR, and **(D)** ER. *** p< 0.001. Full-length blots can be found at the end of this supplementary information file.

Uncropped Western blots. Part 1 contains the full-length original blots for main figures 3C, 4A, 4C, and 5C. Part 2 contains the full-length original blots for main figures 5E, 5G, 7B, 7D, and 7F. Part 3 contains the full-length original blots for supplementary figures S3D, S3F, and S4A. Part 4 contains the full-length original blots for supplementary figure S6A. For each blot, the cropped area is indicated by a black square. Molecular weights are indicated on each blot.

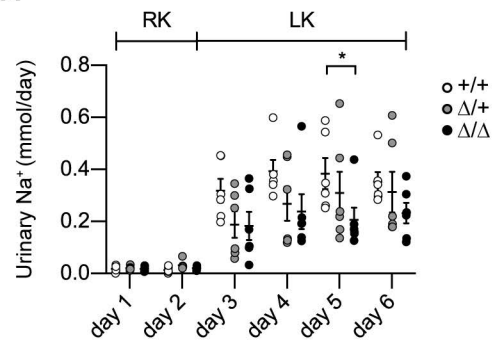
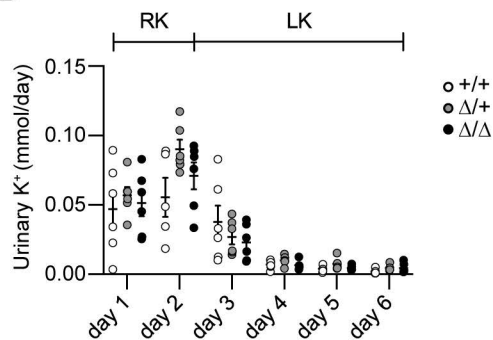
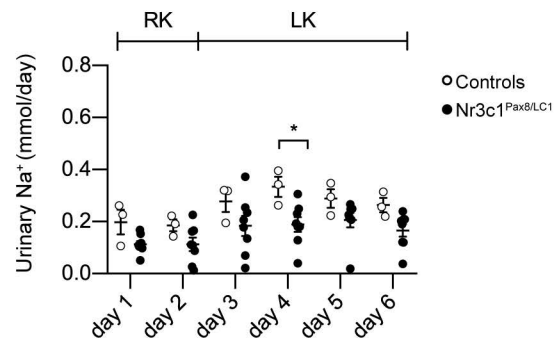
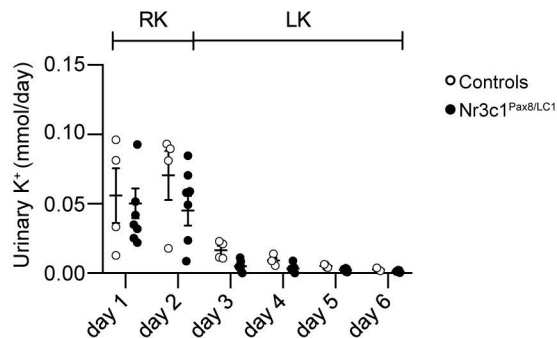
Figure S1**A****B****C****D**

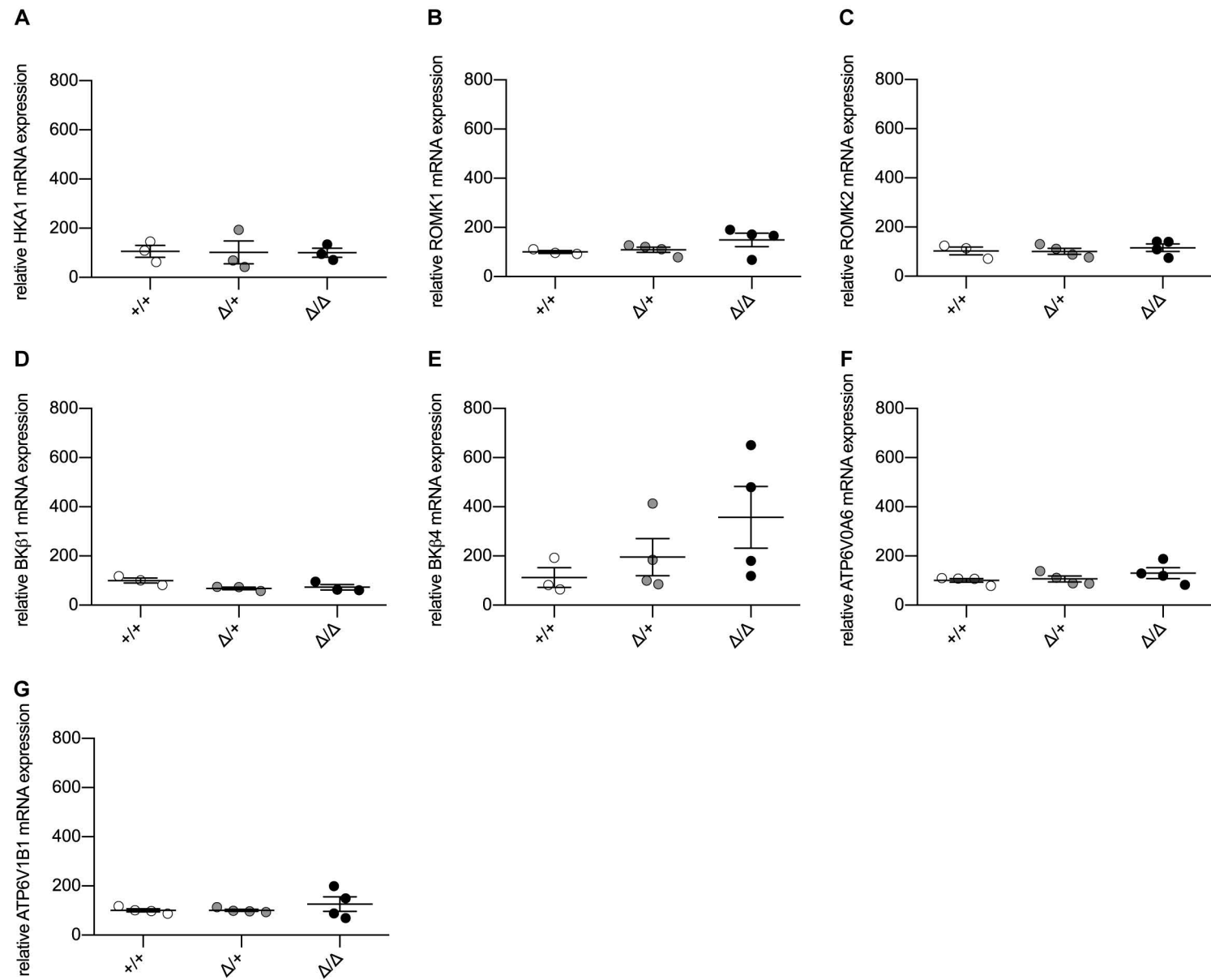
Figure S2

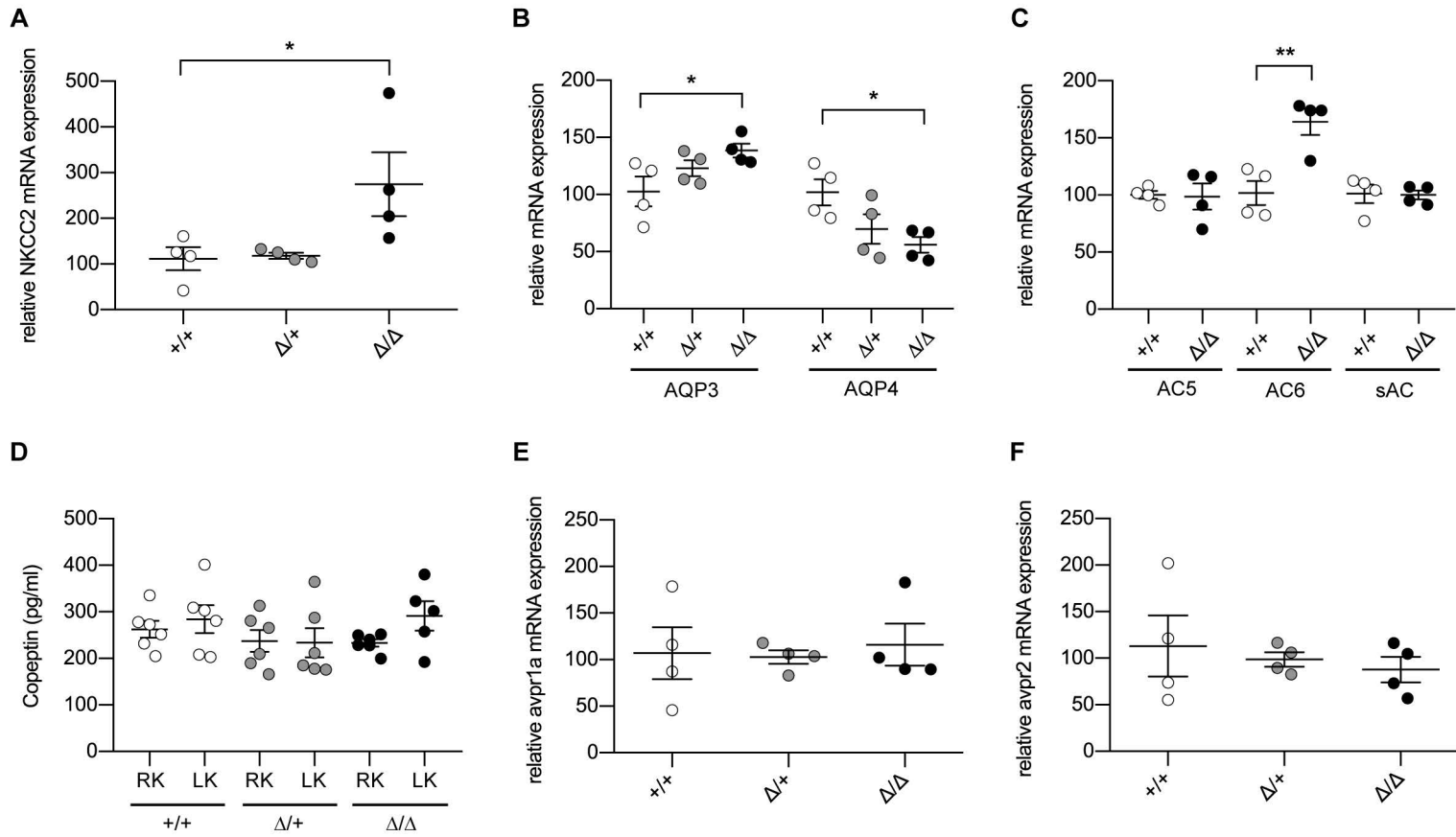
Figure S3

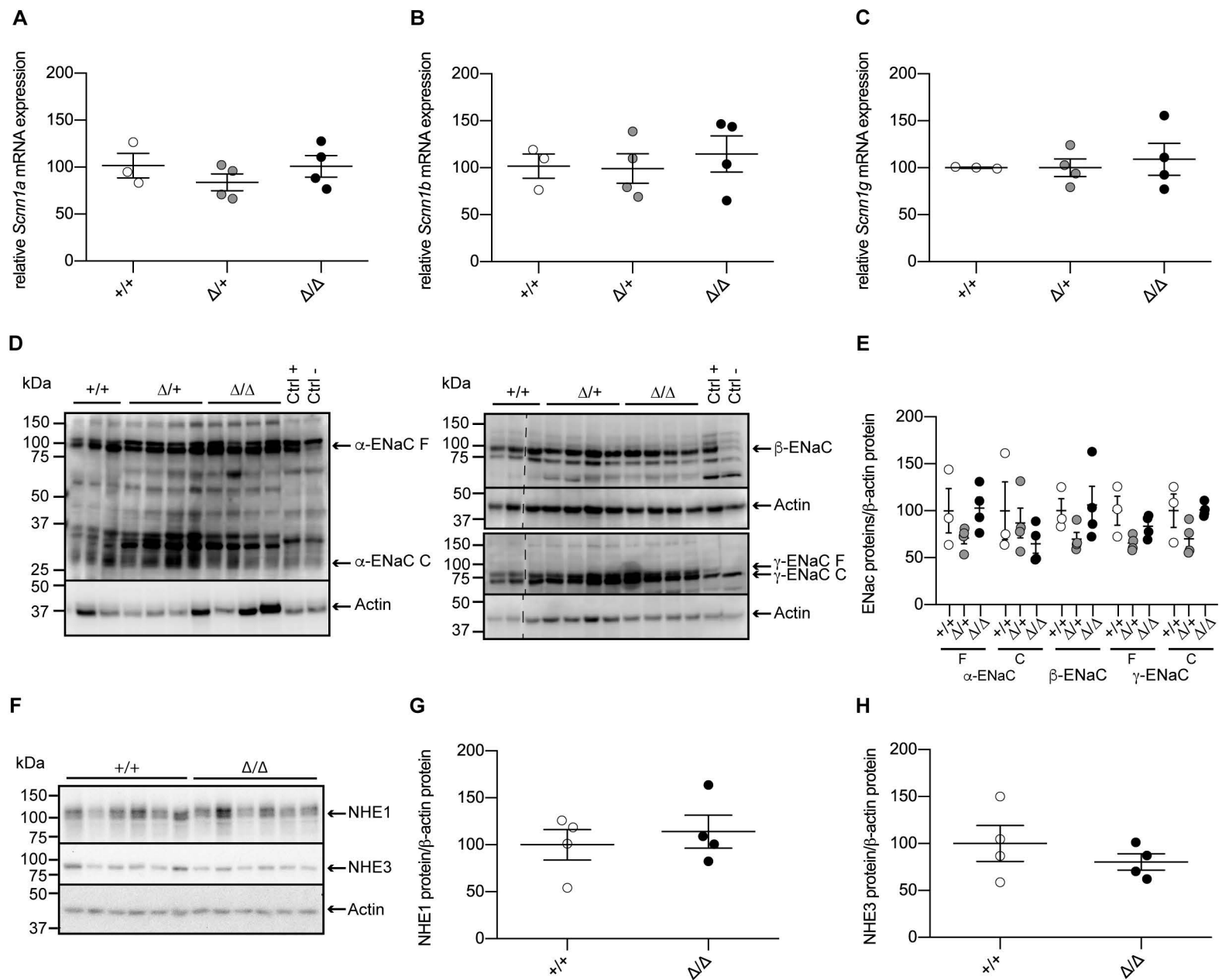
Figure S4

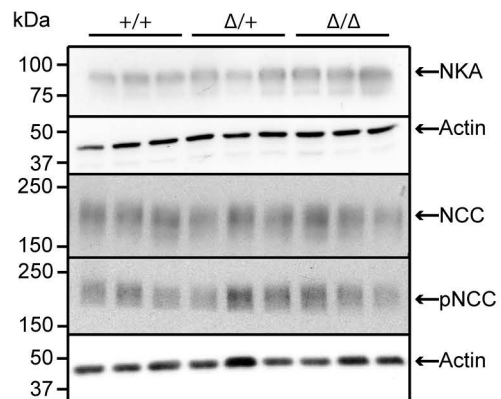
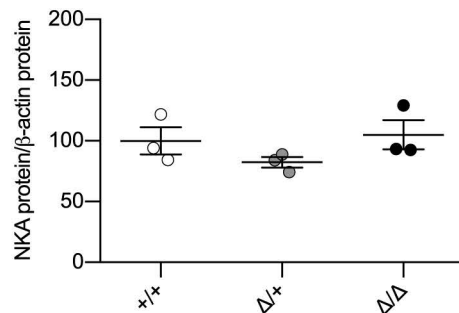
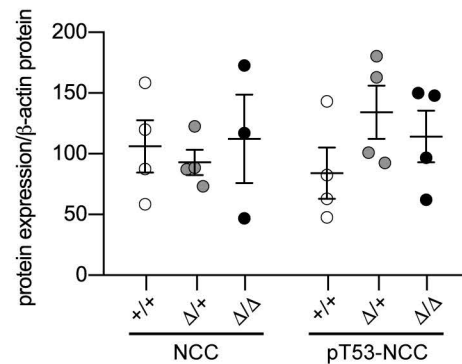
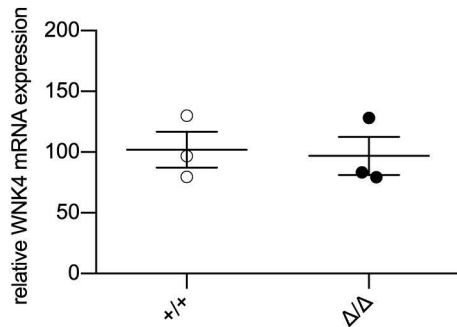
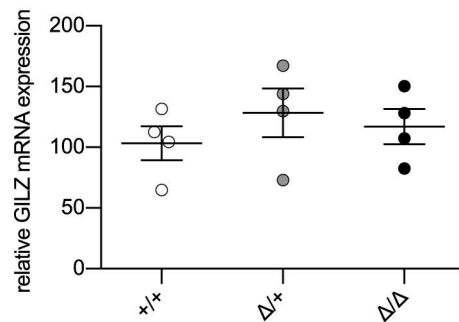
Figure S5**A****B****C****D****E**

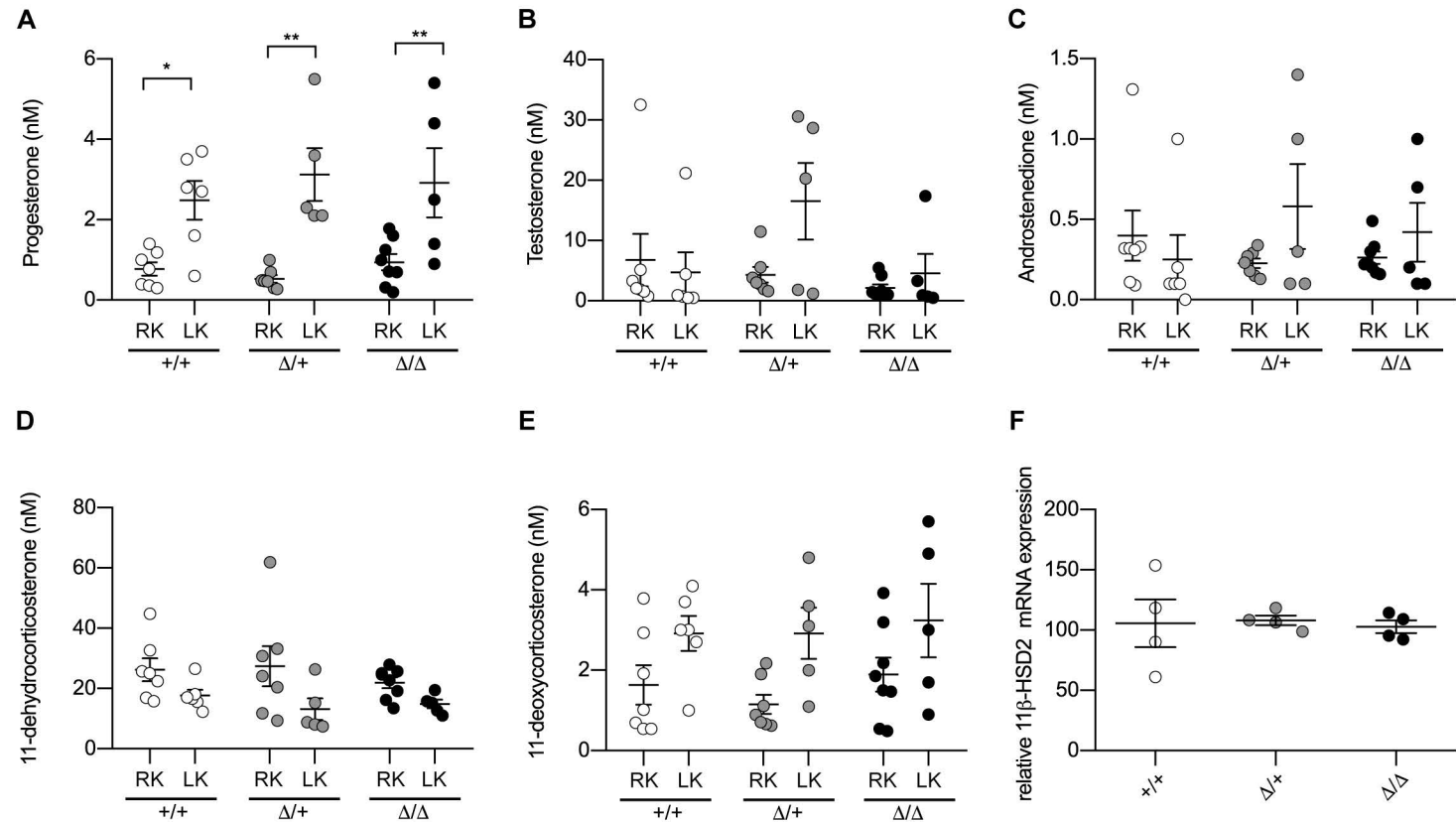
Figure S6

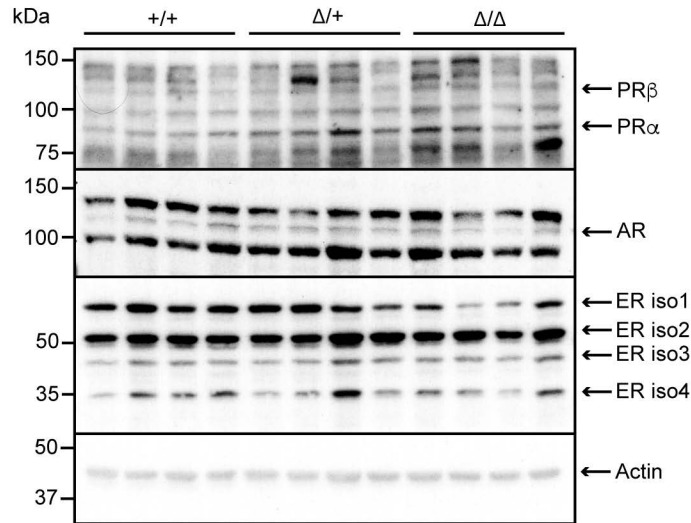
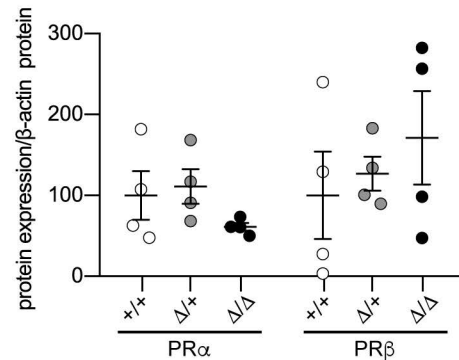
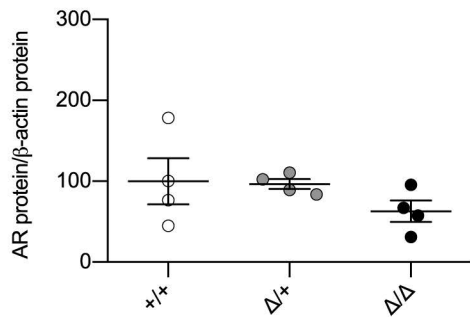
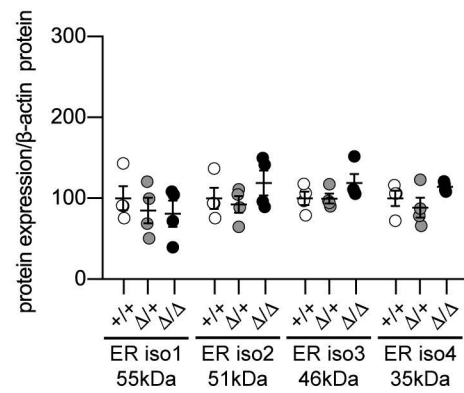
Figure S7**A****B****C****D**

Table S1 : Daily urinary Na⁺ excretion (mmol/day) for CAP2/*Tmprss4* mice after 2 days under regular K⁺ diet (days 1 and 2) and 4 days (days 3-6) under K⁺-deficient diet.

	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6
+/+ 1	0.03	0.01	0.22	0.36	0.54	0.36
+/+ 2	0.01	0.01	0.45	0.60	0.59	0.53
+/+ 3	0.03	0.02	0.20	0.30	0.31	0.20
+/+ 4	0.01	0.01	0.45	0.34	0.26	0.24
+/+ 5	0.01	0.01	0.30	0.38	0.35	0.30
+/+ 6	0.03	0.03	0.28	0.38	0.25	0.18
Δ/+ 1	0.02	0.07	0.30	0.45	0.65	0.61
Δ/+ 2	0.03	0.02	0.35	0.46	0.44	0.50
Δ/+ 3	0.01	0.03	0.25	0.32	0.22	0.19
Δ/+ 4	0.02	0.03	0.06	0.13	0.24	0.22
Δ/+ 5	0.02	0.03	0.08	0.12	0.14	0.18
Δ/+ 6	0.02	0.02	0.09	0.13	0.17	0.18
Δ/Δ 1	0.01	0.03	0.36	0.57	0.44	0.40
Δ/Δ 2	0.01	0.01	0.03	0.21	0.17	0.37
Δ/Δ 3	0.02	0.02	0.10	0.12	0.15	0.13
Δ/Δ 4	0.03	0.02	0.33	0.19	0.19	0.23
Δ/Δ 5	0.03	0.02	0.11	0.14	0.13	0.12
Δ/Δ 6	0.02	0.02	0.17	0.19	0.16	0.22

Table S2 : Daily urinary K⁺ excretion (mmol/day) for CAP2/*Tmprss4* mice after 2 days under regular K⁺ diet (days 1 and 2) and 4 days (days 3-6) under K⁺-deficient diet.

	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6
+/+ 1	0.06	0.05	0.02	0.03	0.03	0.04
+/+ 2	0.02	0.05	0.03	0.02	0.02	0.05
+/+ 3	0.07	0.07	0.01	0.02	0.02	0.03
+/+ 4	0.01	0.02	0.01	0.01	0.00	0.01
+/+ 5	0.03	0.03	0.01	0.01	0.01	0.01
+/+ 6	0.07	0.06	0.03	0.02	0.01	0.02
Δ/+ 1	0.05	0.06	0.03	0.01	0.04	0.05
Δ/+ 2	0.08	0.07	0.02	0.03	0.03	0.05
Δ/+ 3	0.04	0.04	0.01	0.02	0.02	0.03
Δ/+ 4	0.06	0.08	0.03	0.03	0.05	0.03
Δ/+ 5	0.06	0.07	0.02	0.02	0.03	0.05
Δ/+ 6	0.05	0.07	0.02	0.02	0.01	0.03
Δ/Δ 1	0.07	0.06	0.01	0.01	0.02	0.06
Δ/Δ 2	0.03	0.03	0.01	0.02	0.02	0.03
Δ/Δ 3	0.05	0.05	0.02	0.01	0.01	0.03
Δ/Δ 4	0.08	0.07	0.02	0.01	0.03	0.03
Δ/Δ 5	0.07	0.06	0.01	0.01	0.02	0.03
Δ/Δ 6	0.06	0.06	0.02	0.01	0.01	0.04

Table S3 : Daily urinary Na⁺ excretion (mmol/day) for Nr3c1^{Pax8/LC1} mice after 2 days under regular K⁺ diet (days 1 and 2) and 4 days (days 3-6) under K⁺-deficient diet.

	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6
Control 1	0.33	0.22	0.32	0.40	0.35	0.31
Control 2	0.26	0.19	0.19	0.26	0.22	0.30
Control 3	0.11	0.14	0.32	0.34	0.29	0.26
Control 4	0.23	0.07	0.36	0.29	0.24	0.16
Control 5	0.14	0.10	0.35	0.31	0.31	0.24
Control 6	0.15	0.12	0.28	0.34	0.29	0.29
Nr3c1 ^{Pax8/LC1} 1	0.12	0.13	0.20	0.19	0.23	0.18
Nr3c1 ^{Pax8/LC1} 2	0.12	0.11	0.21	0.23	0.23	0.20
Nr3c1 ^{Pax8/LC1} 3	0.10	0.09	0.07	0.20	0.22	0.19
Nr3c1 ^{Pax8/LC1} 4	0.11	0.10	0.21	0.19	0.21	0.19
Nr3c1 ^{Pax8/LC1} 5	0.10	0.12	0.13	0.18	0.21	0.12
Nr3c1 ^{Pax8/LC1} 6	0.05	0.11	0.18	0.18	0.25	0.16

Table S4 : Daily urinary K⁺ excretion (mmol/day) for Nr3c1^{Pax8/LC1} mice after 2 days under regular K⁺ diet (days 1 and 2) and 4 days (days 3-6) under K⁺-deficient diet.

	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6
Control 1	0.10	0.09	0.02	0.01	0.00	0.03
Control 2	0.08	0.09	0.01	0.01	0.01	0.02
Control 3	0.03	0.08	0.02	0.01	0.00	0.02
Control 4	0.03	0.02	0.01	0.01	0.01	0.02
Control 5	0.05	0.06	0.01	0.01	0.01	0.02
Control 6	0.06	0.05	0.01	0.01	0.01	0.02
Nr3c1 ^{Pax8/LC1} 1	0.05	0.05	0.01	0.01	0.00	0.01
Nr3c1 ^{Pax8/LC1} 2	0.04	0.06	0.00	0.00	0.00	0.01
Nr3c1 ^{Pax8/LC1} 3	0.03	0.05	0.00	0.00	0.01	0.01
Nr3c1 ^{Pax8/LC1} 4	0.02	0.04	0.01	0.00	0.00	0.01
Nr3c1 ^{Pax8/LC1} 5	0.03	0.05	0.00	0.00	0.00	0.01
Nr3c1 ^{Pax8/LC1} 6	0.07	0.04	0.00	0.00	0.00	0.02

Fig. 3C

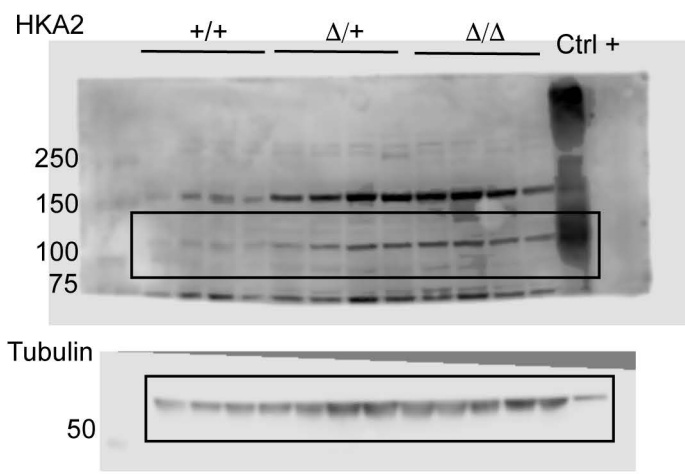


Fig. 4A

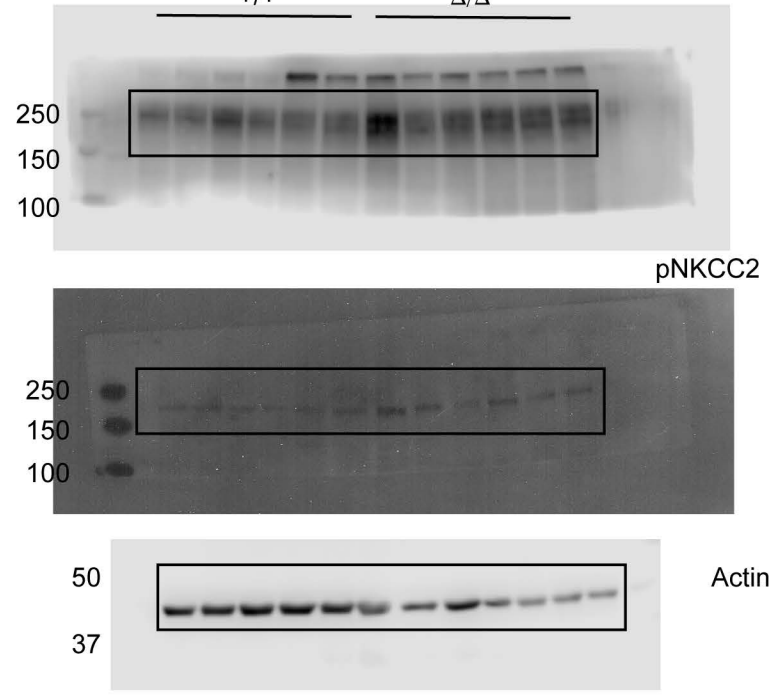


Fig. 4C

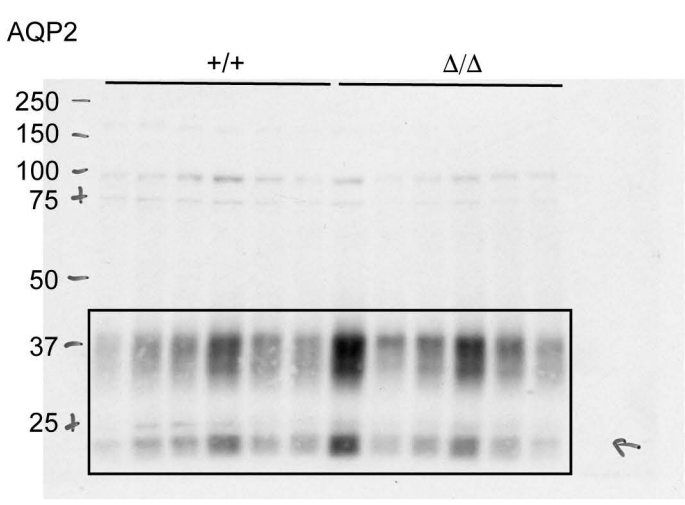
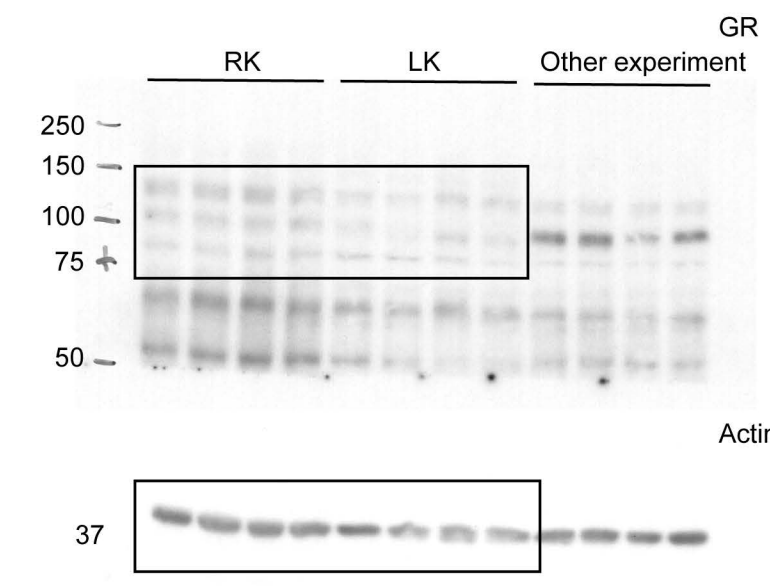


Fig. 5C



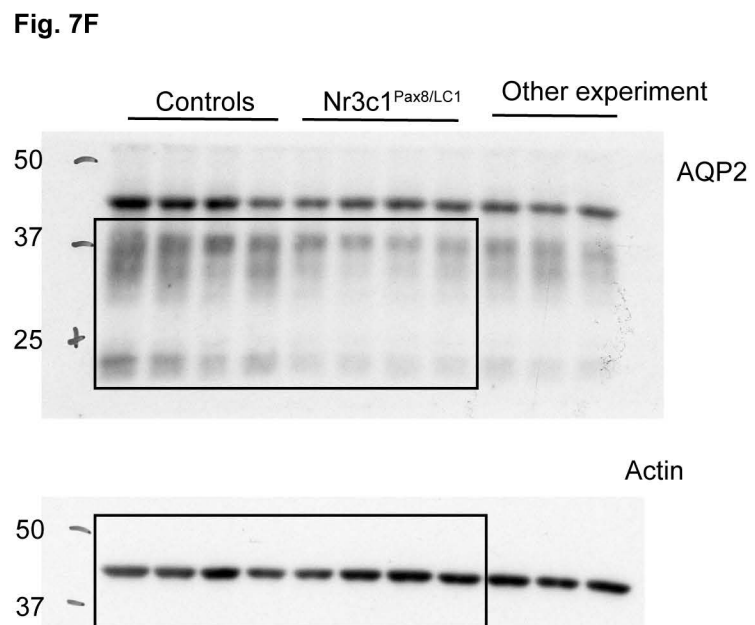
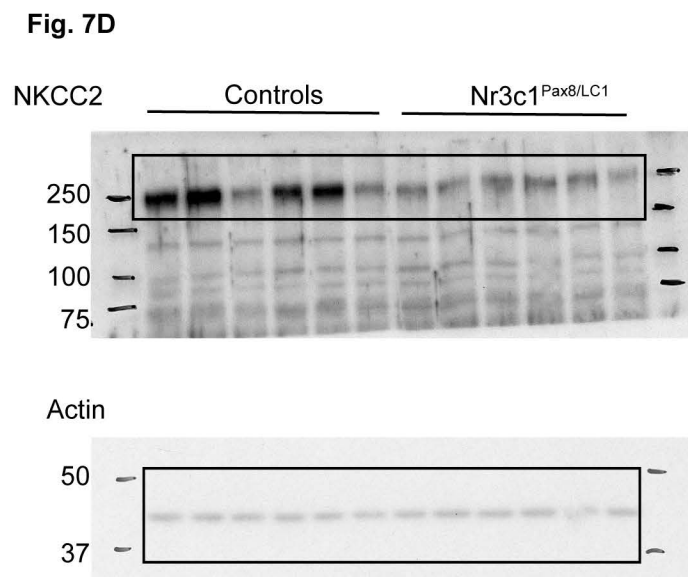
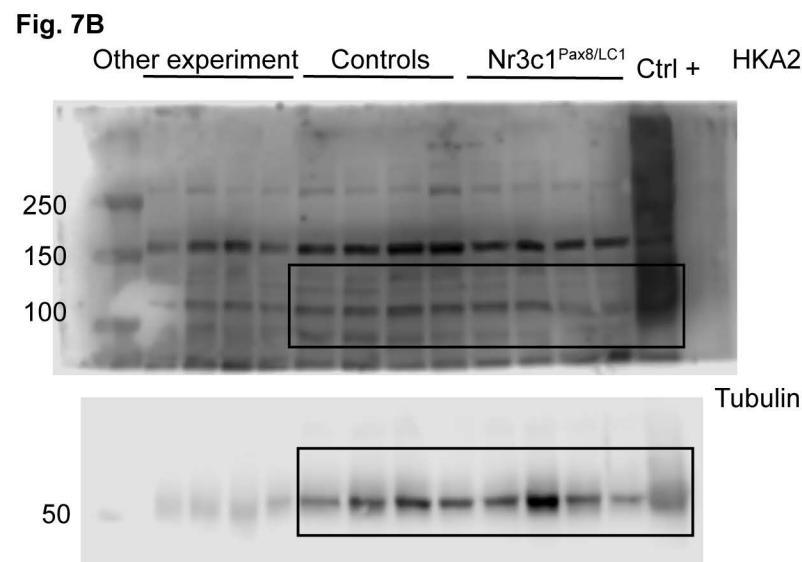
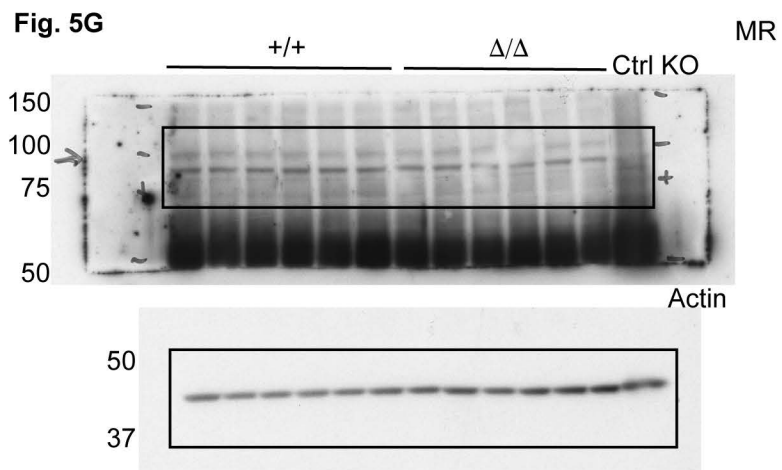
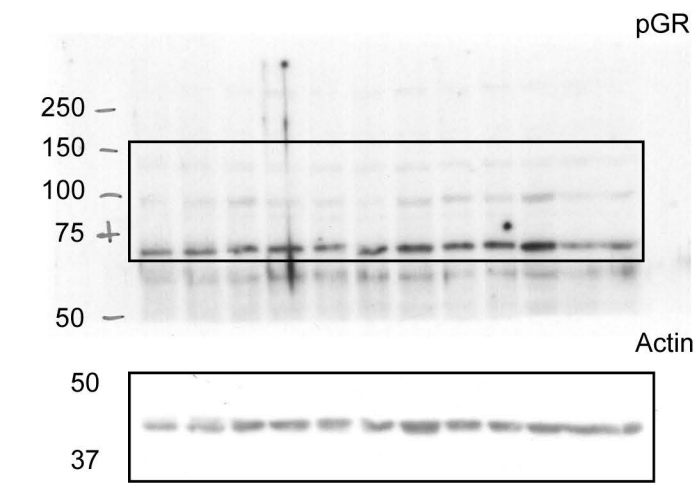
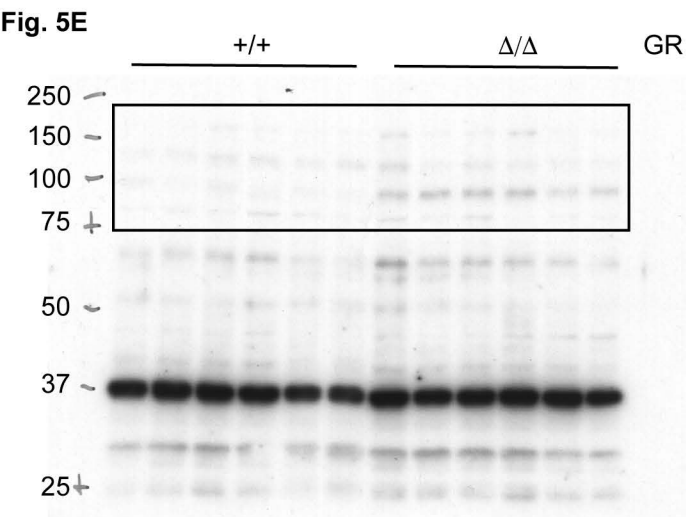


Fig. S4D

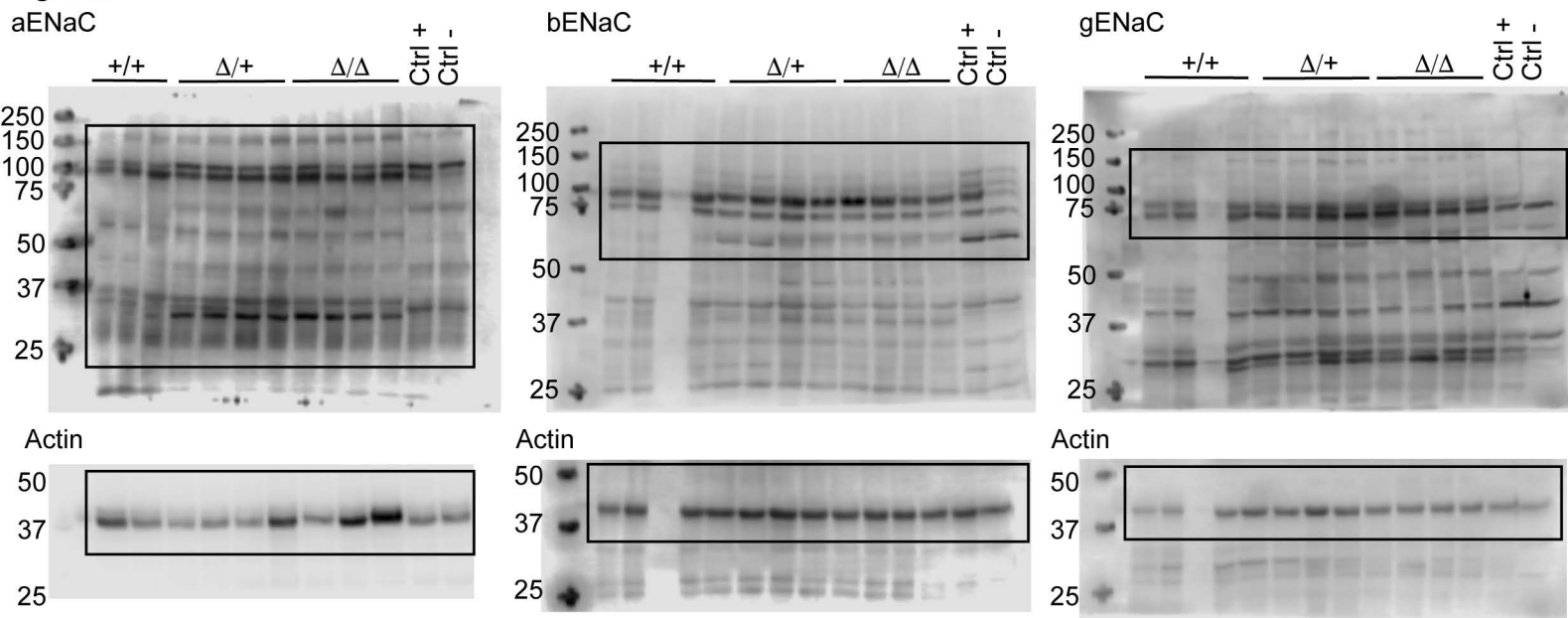


Fig. S4F

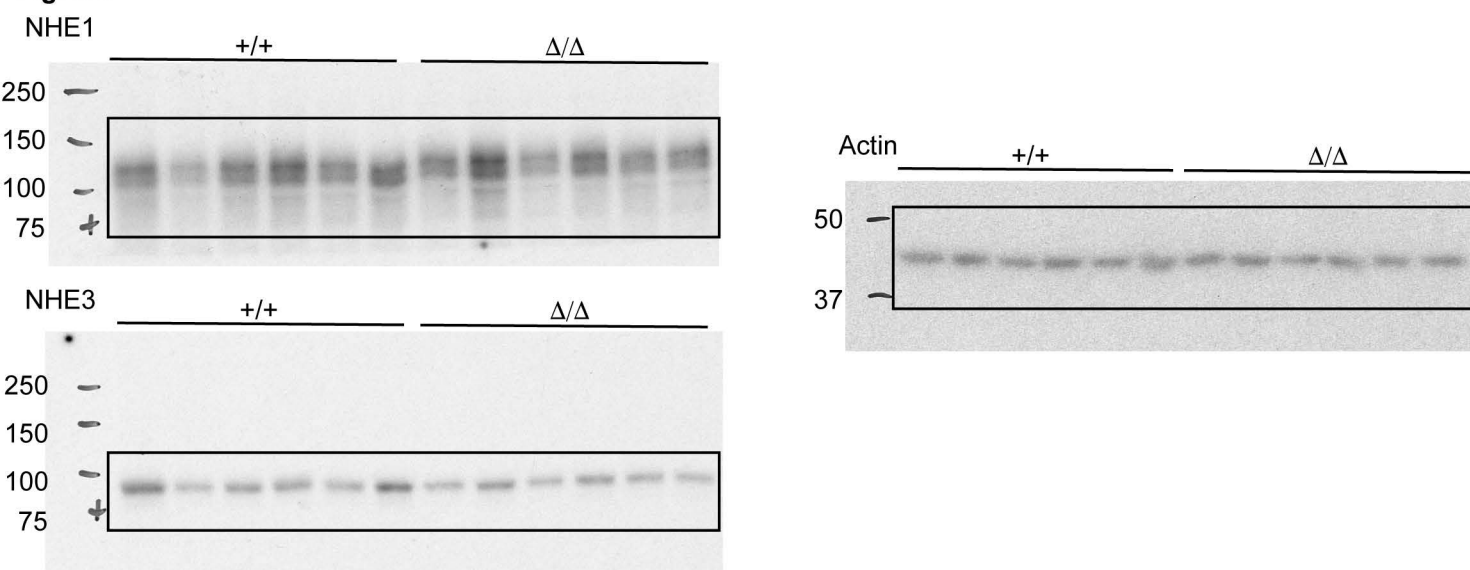
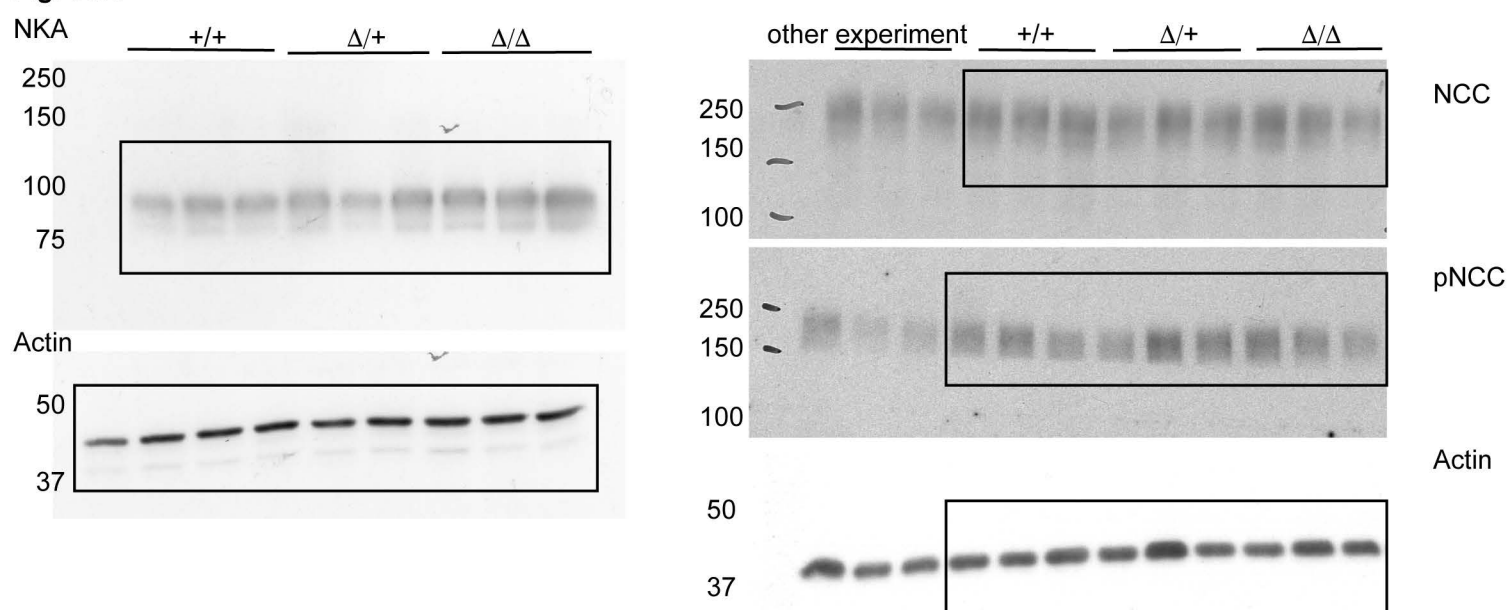


Fig. S5A



Uncropped Western blots part 4

Fig. S7A

