

Appendix A

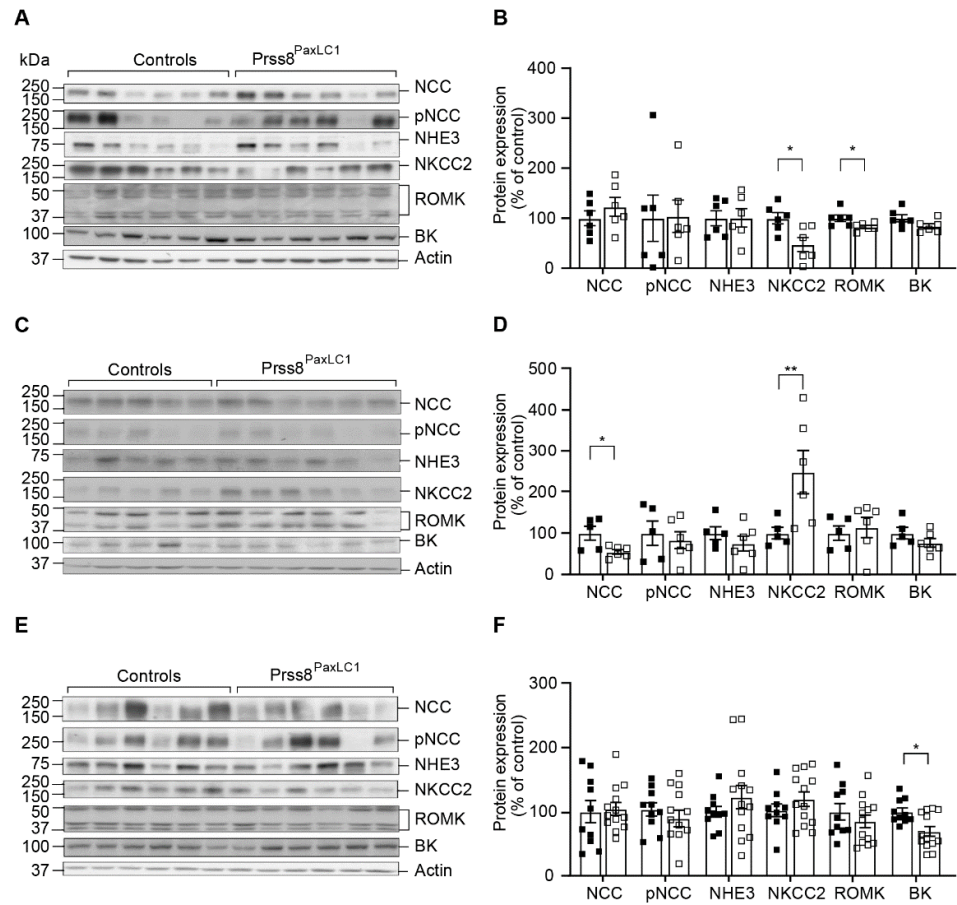


Figure S1. In *Prss8^{PaxLC1}* Ko mice, NKCC2 and ROMK protein expression levels were significantly reduced under standard, increased under high (only NKCC2) and not different on low Na⁺ diet. Western blot analyses of NCC, pNCC (pT53-NCC), NHE3, NKCC2, ROMK and BK on kidney lysates from control and *Prss8^{PaxLC1}* Ko mice under (A) standard, (C) high and (E) low Na⁺ diet; β -actin was used as loading control. (B,D,F) Quantification of data. Values are mean \pm SEM and datasets were analyzed by an unpaired two-tailed *t* test. *P* values <0.05 were considered statistically significant. **P* < 0.05, *P* < 0.01.**

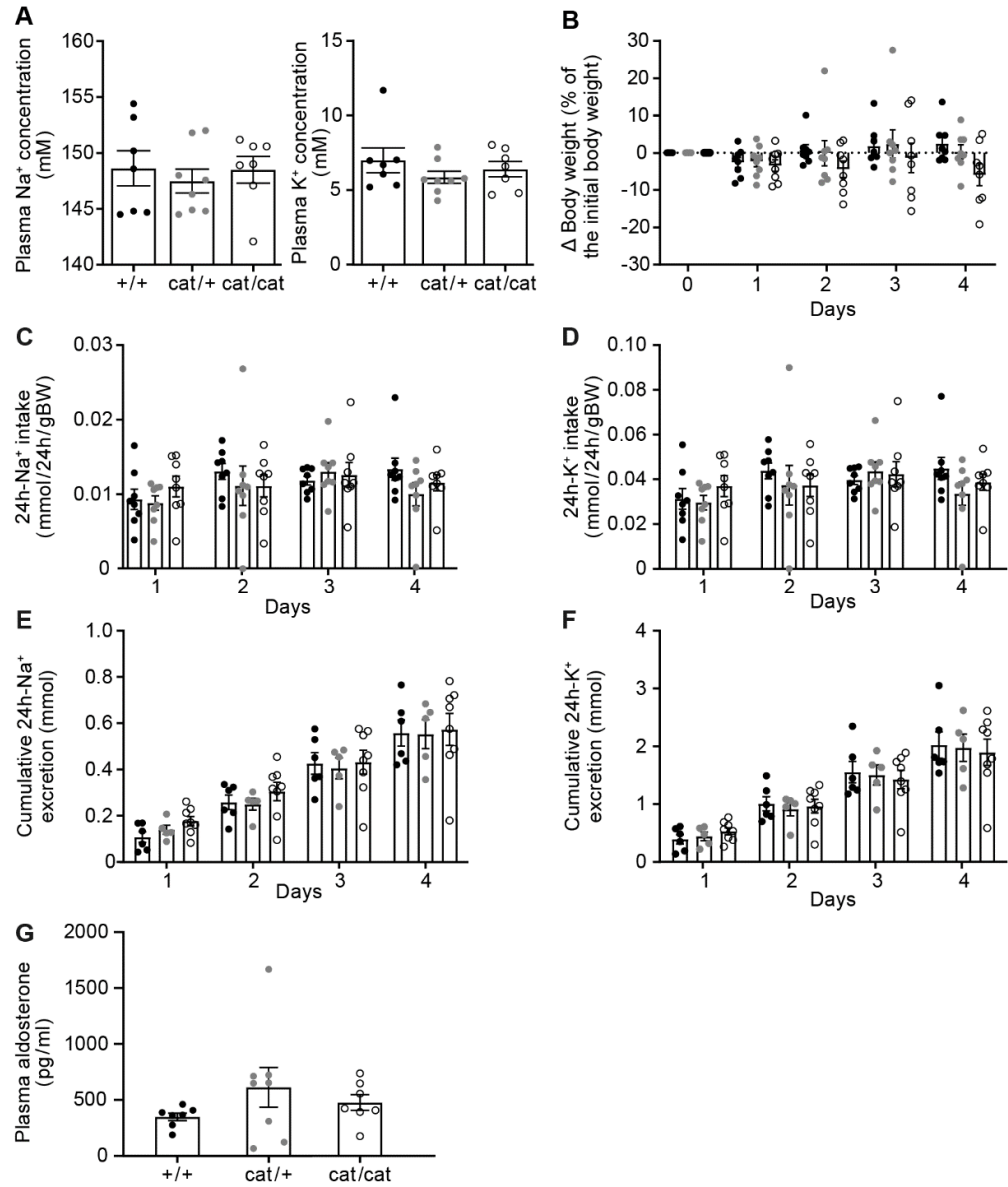


Figure S2. Under standard diet, heterozygous and homozygous mutant *Prss8*^{cat} mice displayed normal Na⁺ and K⁺ handling. (A) Plasma Na⁺ and K⁺ (mM) concentrations in control, heterozygous and homozygous *Prss8*^{cat} mice. (B) Body weight changes (expressed as percent of initial body weight), (C) 24h Na⁺ and (D) K⁺ (mmol/24h/gBW) intake and, (E) 24h cumulative Na⁺ and (F) K⁺ excretion (mmol/24h/gBW). (G) Plasma aldosterone levels. (+/+ n=7, *Prss8*^{cat/+} n= 8, *Prss8*^{cat/cat}, n=7). Results are presented as mean ± SEM. A and G were analyzed with a one-way ANOVA with a Tukey test. B-F were analyzed with a two-way ANOVA. *P* values <0.05 were considered statistically significant.

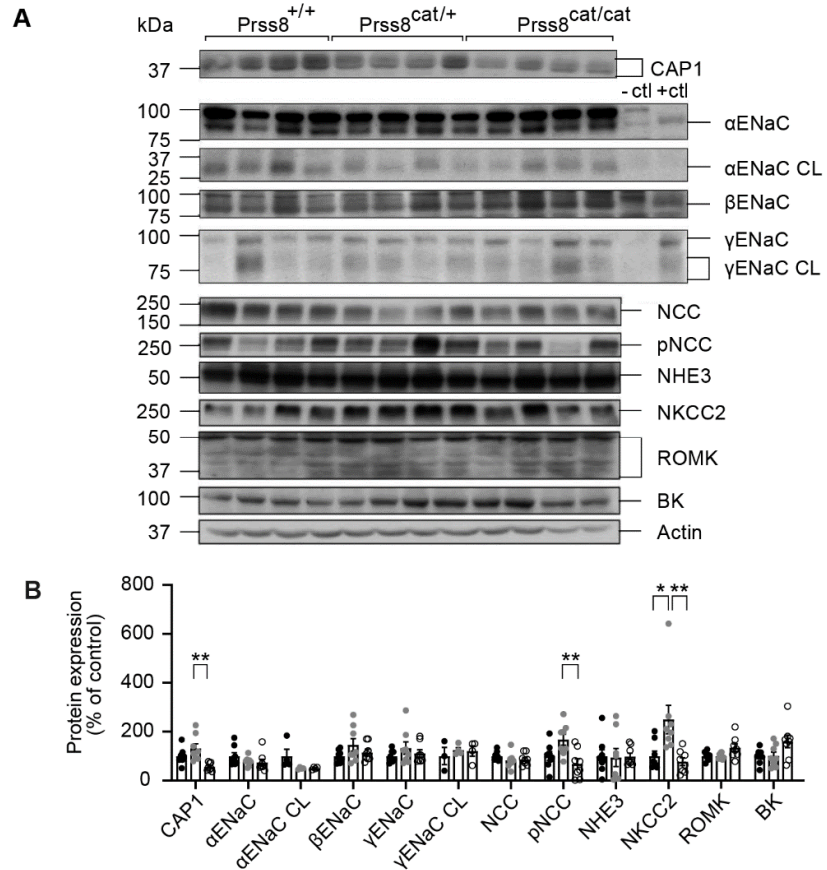


Figure S3. Under standard diet, heterozygous mutant *Prss8^{cat/+}* mice exhibited increased CAP1/*Prss8*, pNCC and NKCC2 protein expression levels. (A) Representative Western blot analyses of CAP1/*Prss8*, αENaC, αENaC CL (cleaved), βENaC, γENaC, γENaC CL (cleaved), βENaC, γENaC, γENaC CL (cleaved), NCC, pNCC (pT53-NCC), NHE3, NKCC2, ROMK and BK on kidney lysates from wild-type (+/+), heterozygous (*Prss8^{cat/+}*) and homozygous mutant *Prss8^{cat/cat}* (n=8 per genotype) littermates; β-actin was used as loading control. Kidney lysates from αENaC [9], βENaC and γENaC [7],[8] control and renal tubular-specific knockouts were used as negative (-ctl) and positive (+ctl) control. (B) Quantification of Western blots. Values are mean ± SEM and datasets were analyzed by a one-way ANOVA with a Tukey test. P values <0.05 were considered statistically significant. *P <0.05, **P <0.01.

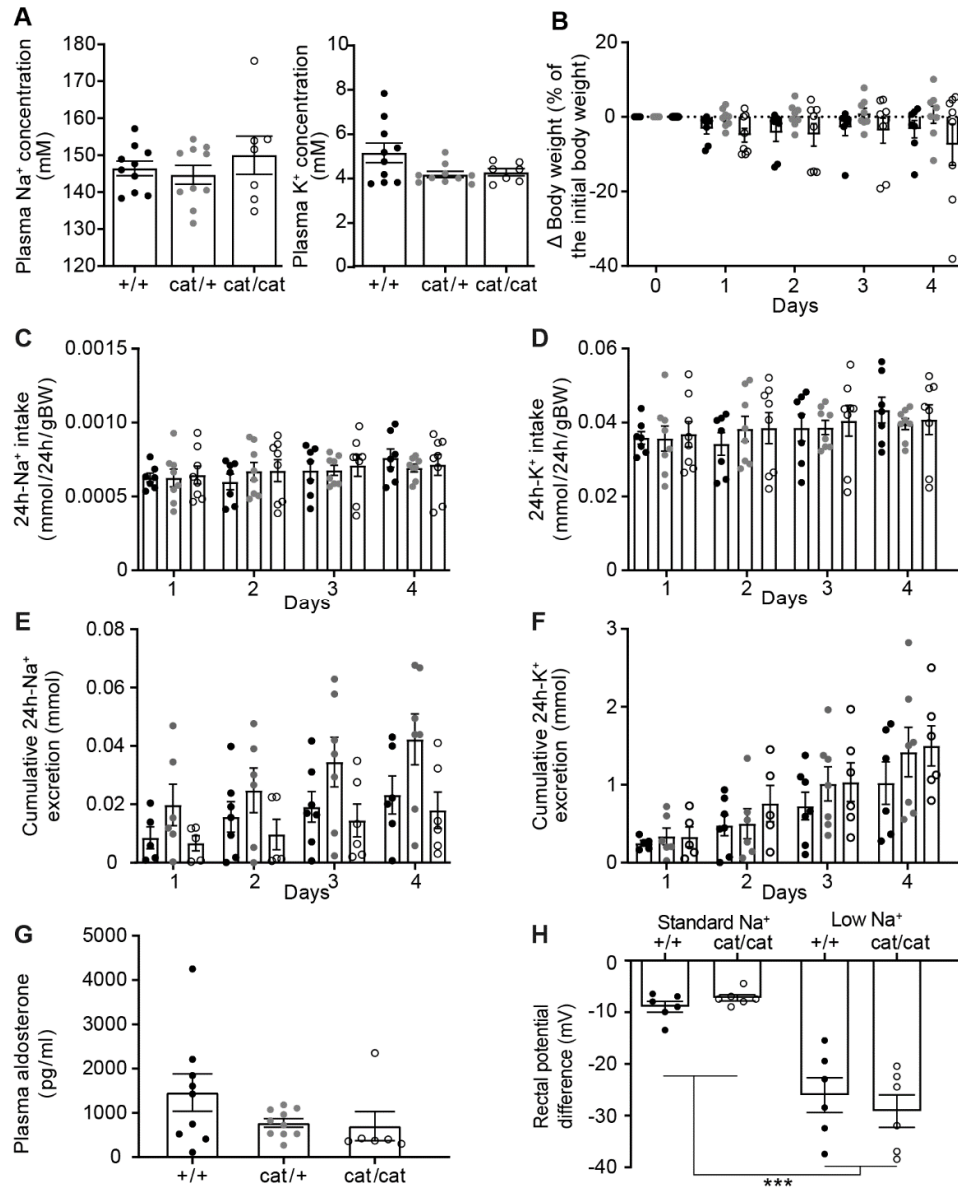


Figure S4. Under low Na⁺ diet, heterozygous (*Prss8^{cat/+}*) and homozygous mutant *Prss8^{cat/cat}* mice displayed normal Na⁺ and K⁺ handling. (A) Plasma Na⁺ and K⁺ (mM) concentrations in control, heterozygous and homozygous *Prss8^{cat}* mice (B) Body weight changes (expressed as percent of initial body weight) (C) 24h Na⁺ and (D) K⁺ (mmol/24h/gBW) intake and (E) 24h cumulative Na⁺ and (F) K⁺ excretion (mmol/24h/gBW). (G) Plasma aldosterone levels. (+/+ n=5-10, *Prss8^{cat/+}* n=6-10, *Prss8^{cat/cat}*, n=5-7). (H) Rectal ΔPD_{amil} (mV) measurements reflecting ENaC activity in wild-type (black) and homozygous mutant (white circle) *Prss8^{cat/+}* littermates in the distal colon following standard Na⁺ (left panel) and low Na⁺ diet (right panel). Results are presented as mean ± SEM, A,G,H were analyzed with a one-way ANOVA with a Tukey test. B-F were analyzed with a two-way ANOVA. P values <0.05 were considered statistically significant. ***P <0.001

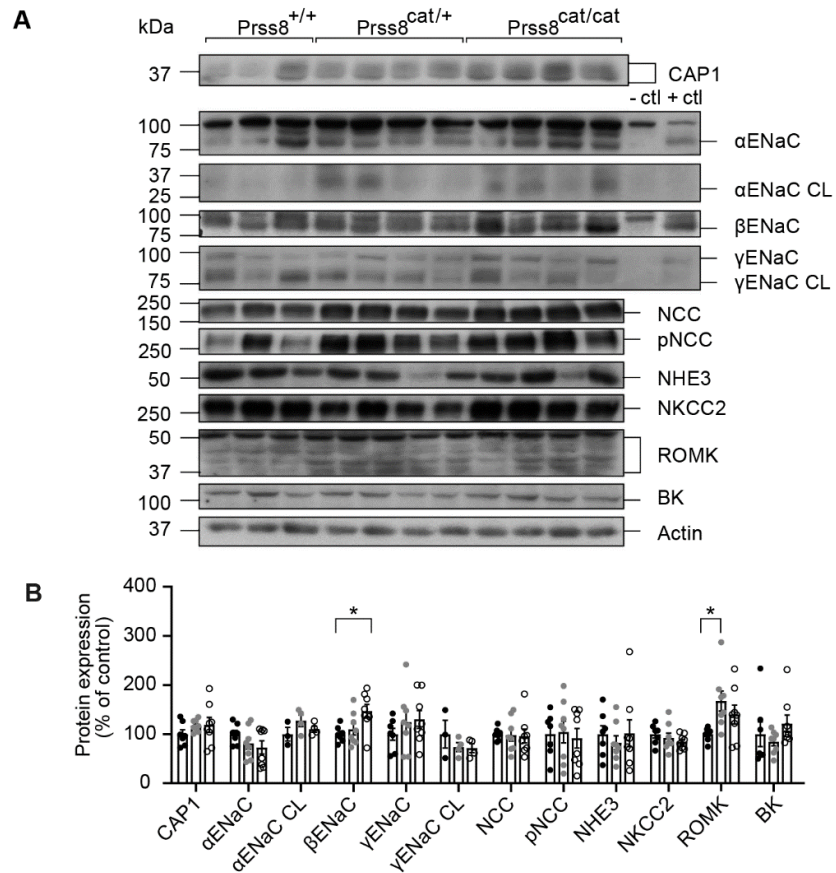


Figure S5. Under low Na⁺ diet, homozygous mutant *Prss8^{cat/cat}* mice showed increased βENaC protein expression levels. (A) Representative Western blot analyses of CAP1/*Prss8*, αENaC, αENaC CL (cleaved), βENaC, γENaC, γENaC CL (cleaved), NCC, pNCC (pT53-NCC), NHE3, NKCC2, ROMK and BK on kidney lysates from wild-type (+/+), heterozygous (*Prss8^{cat/+}*) and homozygous mutant *Prss8^{cat/cat}* (n=8 per genotype) littermates; β-actin was used as loading control. Kidney lysates from αENaC [9], βENaC and γENaC [7],[8] control and renal tubular-specific knockouts were used as negative (-ctl) and positive (+ctl) control. (B) Quantification of Western blots. Values are mean ± SEM and datasets were analyzed by a one-way ANOVA with a Tukey test. *P* values <0.05 were considered statistically significant. **P*<0.05