SUPPLEMENTARY FIGURES ONLINE

Consequences of SPAK inactivation on Hyperkalemic Hypertension caused by *WNK1* mutations: evidence for differential roles of WNK1 and WNK4.

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Supplementary Figure S1. The expression and phosphorylation of NKCC2 are decreased in *WNK1*^{+/FHHt}:*SPAK*^{243A/243A} mice.

Upper panel: Representative immunoblots of NKCC2 and phosphorylated NKCC2 performed on the membrane-enriched fractions of the renal cortex of mice of each genotype. As pNKCC2 antibody is also able to recognize pNCC, we loaded a protein sample extracted from a *NCC*^{-/-} mouse kidney (in blue, last lane of the pNKCC2 blot) to identify the band corresponding specifically to pNKCC2.

Lower panel: Densitometric analysis. NKCC2 abundance and phosphorylation are decreased in $WNK1^{+/FHHt}$: $SPAK^{243A/243A}$ and $SPAK^{243A/243A}$ mice. In contrast to NCC, there is no difference between these two groups. Number of animals: 4 control, 5 $WNK1^{+/FHHt}$, 4 $WNK1^{+/FHHt}$: $SPAK^{243A}$: 243A and 4 $SPAK^{243A:243A}$ male mice. Values are means \pm s.e.m. ^a p<0.05 vs. controls. ^b p<0.05 vs. $WNK1^{+/FHHt}$ mices (Kruskal-Wallis one-way analysis of variance, followed by Mann-Whitney test). Immunoblot images are full-lenght images.



Supplementary Figure S2. WNK4 expression is decreased in $WNK1^{+/FHHt}$: $SPAK^{243A:243A}$ mice. Upper panel: Representative immunoblot of WNK4 performed on protein extracted from the renal cortex of mice of each genotype. Lower panel: Densitometric analysis. WNK4 abundance is decreased in $WNK1^{+/FHHt}$: $SPAK^{243A/243A}$ and $SPAK^{243A/243A}$ mice compared to control mice. Number of animals: 4 control, 5 $WNK1^{+/FHHt}$, 4 $WNK1^{+/FHHt}$: $SPAK^{243A:243A}$ and 4 $SPAK^{243A:243A}$ male mice. Values are means ± s.e.m. ^a p<0.05 vs. controls. ^b p<0.05 vs. $WNK1^{+/FHHt}$ mice. The immunoblot image is a full-lenght image.



Supplementary Figure S3. Full-lenght iamges of the NCC and pNCC immunoblots, corresponding to Figure 2.



Supplementary Figure S4. Full-lenght iamges of the pendrin and γENAC immunoblot, corresponding to Figure 3.



Supplementary Figure S5. Total SPAK, total OSR1, pSPAK/pOSR1 (S-motif) and Cab39 full-lenght immunoblot images.



Supplementary Figure S6. Representative Coomassie gel staining. Preliminary 10% SDS-PAGE gels were run and stained with Coomassie blue to confirm equality of loading in each lane.





L = length of the ROI

M25% = Mean intensity of pixels from 0 (apical membrane) to L/4

M100% = Mean intensity of all pixels

Apical OSR1 = M25% / M100%

Supplementary Figure S7. Schematic description of the method used to quantify OSR1 apical staining. A detailed description is provided in the Material and Methods section.