Supplementary information

Impaired NEPHRIN localization in kidney organoids derived from nephrotic patient iPS cells

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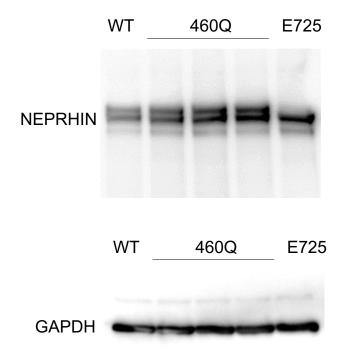
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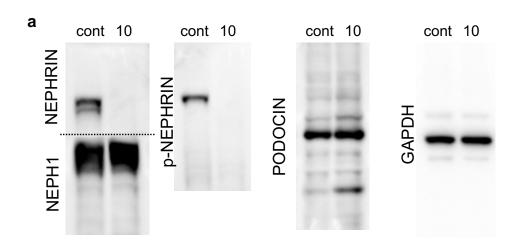
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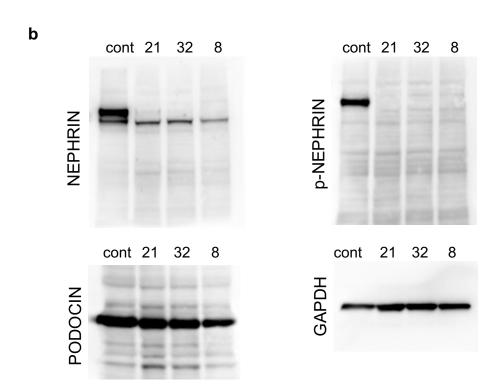


Supplementary Figure S1

The full-length western blots used for Figure 2

Western blotting analysis of HEK293 cells overexpressing wild-type (WT), R460Q mutant (Patient 3), and E725D mutant (Patient 0) NEPHRIN. GAPDH is used as a loading control.

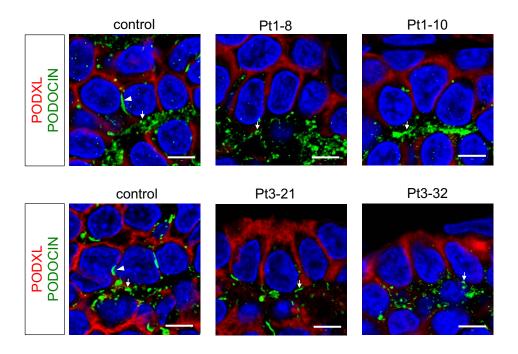




Supplementary Figure S2

The full-length western blots used for Figure 6

Western blot analyses of NEPHRIN, phosphorylated NEPHRIN (p-NEPHRIN), and PODOCIN in kidney organoids derived from Patient 1 (a) and 3 (b). The clone numbers are indicated at the top. GAPDH is used as a loading control. cont: control clones.

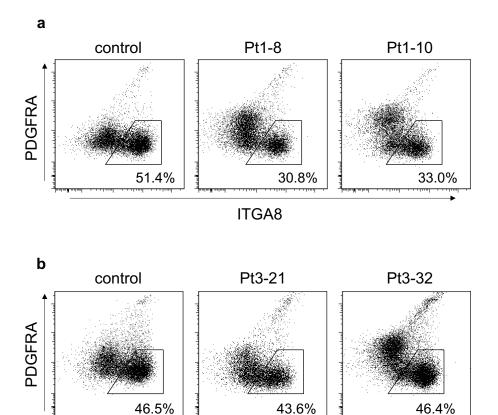


Supplementary Figure S3

Localization of PODOCIN is impaired in patient-derived podocytes

Section immunostaining of kidney organoids was performed with anti-PODOCIN and anti-PODXL antibodies (upper row: Patient 1; lower row: Patient 3). PODOCIN is undetectable in the lateral domains of the podocytes derived from both patients.

Arrowheads: lateral SD precursors; arrows: basal PODOCIN+ domains. Scale bars: $5 \mu m$.



ITGA8

Supplementary Figure S4

Induction of nephron progenitors from iPSCs

Induction was performed for the control clone (201B7), two Patient 1-derived clones (Pt1-8, Pt1-10), and two Patient 3-derived clones (Pt3-21, Pt3-32), and the formation of ITGA8+/PDGFRA- nephron progenitors was analyzed by flow cytometry on day 13. Representative data are shown.