



ILDR1 is important for paracellular water transport and urine concentration mechanism

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Edited by David W. Russell, University of Texas Southwestern Medical Center, Dallas, TX, and approved April 6, 2017 (received for review January 18, 2017)

Whether the tight junction is permeable to water remains highly controversial. Here, we provide evidence that the tricellular tight junction is important for paracellular water permeation and that Ig-like domain containing receptor 1 (ILDR1) regulates its permeability. In the mouse kidney, ILDR1 is localized to tricellular tight junctions of the distal tubules. Genetic knockout of *Ildr1* in the mouse kidney causes polyuria and polydipsia due to renal concentrating defects. Microperfusion of live renal distal tubules reveals that they are impermeable to water in normal animals but become highly permeable to water in *Ildr1* knockout animals whereas paracellular ionic permeabilities in the *Ildr1* knockout mouse renal tubules are not affected. Vasopressin cannot correct paracellular water loss in *Ildr1* knockout animals despite normal effects on the transcellular aquaporin-2-dependent pathway. In cultured renal epithelial cells normally lacking the expression of *Ildr1*, overexpression of *Ildr1* significantly reduces the paracellular water permeability. Together, our study provides a mechanism of how cells transport water and shows how such a mechanism may be exploited as a therapeutic approach to maintain water homeostasis.

tight junction | water channel | nephrology | diuresis | diabetes insipidus

Tight junctions (TJs) are known to be leaky to ions and, thus, to constitute a paracellular pathway with ionic permselectivity similar to that of transmembrane channels (1). Whether the TJ is permeable to water, on the other hand, has been highly controversial. Jorge Fischberg noticed that the corneal endothelium retained over 60% of water permeability even when AQP1, the only aquaporin expressed by these cells, was knocked out by genetic deletion (2). Rosenthal et al. have demonstrated that, in cultured renal epithelium expressing the claudin-2 protein, transepithelial water permeability was significantly higher than in cells without its expression (3). Using an optical microscopic approach, Spring and coworkers have directly measured the velocity of water flow across the tight junction and have ruled out the existence of any significant transjunctional water flow, at least for the bicellular TJ (bTJ) (4). The tricellular tight junction (tTJ) is a specialized cell junction different in ultrastructure from that of the bTJ (5). Unlike the bTJ, which is made of claudin and occludin (6), the proteins making the tTJ include tricellulin and angulins (LSR/angulin-1, ILDR1/angulin-2, and ILDR2/angulin-3) (7, 8). Transgenic knockout (KO) of either tricellulin or *Ildr1* in mice causes hearing loss due to degeneration of mechanosensory cochlear hair cells in the organ of Corti (9, 10). Peculiarly, neither the endocochlear potential nor the paracellular permeability in the stria vascularis changed in these mutant animals. The permeation property of tTJ therefore remains a major mystery.

There are two forms of diabetes insipidus (DI): central (neurogenic) and nephrogenic. The nephrogenic DI (NDI) is caused by the inability of the kidney to reabsorb water. NDI is hereditary and has been linked to two genetic loci. The most common genetic cause (in 95% of cases) is an X-linked disorder mapped to the type-2 vasopressin receptor (AVPR2) gene in the X chromosome, mutation of which makes the collecting duct cells insensitive to vasopressin (11). The second cause is an autosomal recessive disorder linked to the aquaporin-2 (Aqp2) gene in

chromosome 12, and the sites of mutations were found to be functionally important in making the water permeation pore (12) or facilitating proper intracellular trafficking (13). Whether there may exist any previously unknown gene important for renal water reabsorption is a tantalizing question. Here, we have revealed that homozygous deletion of the ILDR1 gene, which encodes a TJ protein controlling the putative paracellular water pathway, can cause NDI-like phenotypes in the mouse kidney.

Results

The Gene Expression of ILDR1 in the Kidney. Whether the kidney expresses ILDR1 has not been determined before. We first performed microdissection on mouse kidneys to isolate each nephron segment, including the glomerulus, the proximal tubule (PT), the thin limb (TL) and the thick limb (TAL) of the loop of Henle, the distal convoluted tubule (DCT), and the collecting duct (CD), obeying a rigorous anatomic criterion (14) (Fig. S1A). Quantitative RT-PCR revealed that the distal nephron, from the TAL to the CD, expressed the highest levels of ILDR1 mRNA, which were over 58-fold higher than in the glomerulus and over 454-fold higher than in the PT (Fig. S1B). To reveal the cellular localization pattern of ILDR1 protein along the nephron, we performed colocalization analyses using molecular markers against each nephron segment. In WT mouse kidneys, ILDR1 proteins were detected in the tricellular junctions of renal tubules lacking the *Lotus tetragonolobus* lectin (a PT marker), but expressing the claudin-19 protein (a TAL marker) (15), the NCC protein (a DCT marker) (16), and the Aqp2 protein (a CD marker) (17) (Fig. 1). The ILDR1 antibody specificity was demonstrated by staining the ILDR1 knockout mouse kidney (Fig. S2). Thus, ILDR1 is expressed in the distal tubules of the kidney.

Significance

The role of tight junction in water homeostasis is an important but understudied area. Here, we reveal a paracellular water permeation pathway made by the tricellular tight junction and regulated by its integral protein, known as *Ildr1*. The importance of paracellular water permeation is demonstrated here with emphasis of its potential role in causing urinary concentrating defects in the kidney. *Ildr1* and its related proteins, *Ildr2* and *Ildr3*, may play broader physiologic roles to establish the water homeostasis in other organs, such as the skin, the lung, and the intestine.

Author contributions: Y.G., N.H., M.B., and J.H. designed research; Y.G., N.H., A.S., S.M., and C.M. performed research; Y.G., N.H., M.B., and J.H. analyzed data; and Y.G., N.H., M.B., and J.H. wrote the paper.

The authors declare no conflict of interest.

This article is a PNAS Direct Submission.

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This article contains supporting information online at www.pnas.org/lookup/suppl/doi:10.1073/pnas.1701006114/-DCSupplemental.

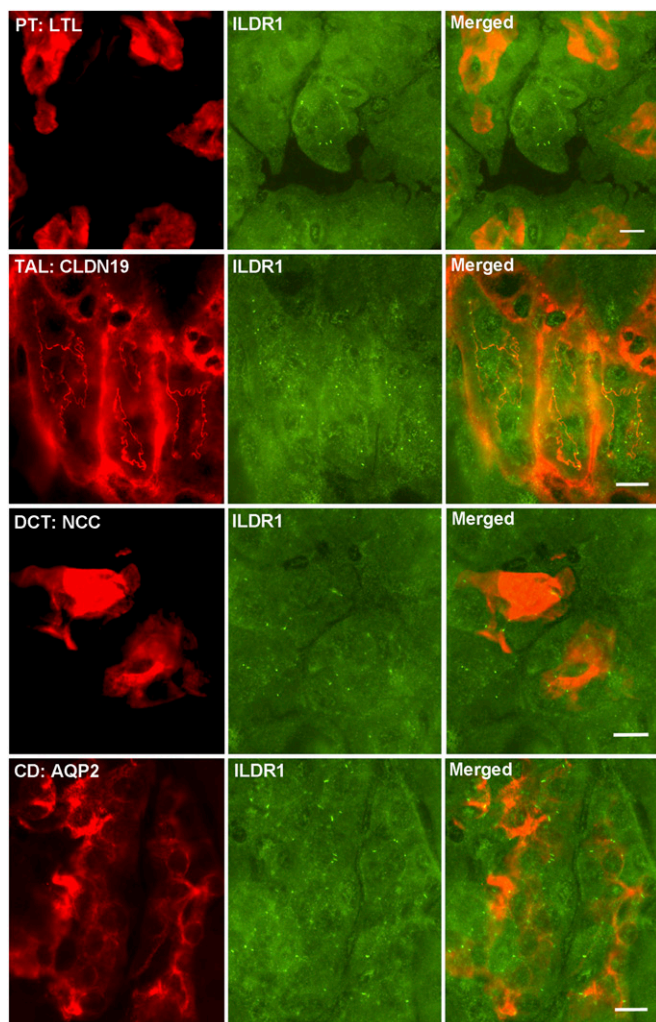


Fig. 1. ILDR1 protein localization in the kidney. Double staining of ILDR1 protein with a PT marker (LTL), with a TAL marker (CLDN19), with a DCT marker (NCC), and with a CD marker (Aqp2). (Scale bars: 10 μm .)

The Renal Phenotypes of ILDR1 KO Mice. ILDR1 KO mice have been generated, described, and validated before (18). The age-matched (4-mo-old) and sex-matched (male) ILDR1 KO mice (ILDR1^{-/-}) and their littermate control mice (ILDR1^{+/+}) were maintained on a regular salt diet (0.24% Na⁺ and 0.30% Cl⁻) with free access to water and subjected to 24-h metabolic cage analyses. The body weight (BW) was significantly lower in KO than in control animals (Table 1). The KO mice developed pronounced polyuria and polydipsia. Their urinary volume (UV) output was 5.8-fold higher than that of control mice (Table 1), and their water intake was 9.0-fold higher than that of control mice (Table 1). The significant urinary volume loss was accompanied by a 74% reduction in urinary osmolality (Table 1), indicating that the kidneys in KO animals were not able to retain water properly. The creatinine clearance (C_{Cr}) rates, as direct estimates of the glomerular filtration rate (GFR), were well-defended in KO animals (Table 1), suggesting that the urinary volume loss was not due to glomerular filtration rate differences. Because diuresis could be an effect of natriuresis or chloriguresis, we determined the kidney's ability to handle salt. The absolute urinary excretion rates for Na⁺ (E_{Na}) (Table 1) and Cl⁻ (E_{Cl}) (Table 1) were similar between KO and control animals. The fractional excretion rates (FE_{Na} , FE_{Cl}) for Na⁺ and Cl⁻ relative to renal creatinine metabolism rates and the plasma Na⁺ and Cl⁻

levels (P_{Na} , P_{Cl}) were not altered in KO animals (Table 1). Compatible with well-preserved salt handling by the kidney, the mean blood pressure (BP) was not different between KO and control animals (Table 1), indicating that the urinary volume loss did not elicit a significant effect on the extracellular fluid volume (ECFV). Hypokalemia has been associated with polyuria due to down-regulation of Aqp2 in the collecting duct (19). No K⁺ deregulation was present in KO mice (Table 1), clearly distinguishing their phenotypes from Bartter syndrome or Gitelman syndrome. Hypercalcemia or hypercalciuria has also been associated with polyuria, presumably due to activation of the calcium-sensing receptor (CaSR) located in the basolateral membrane of the thick ascending limb or the luminal membrane of the collecting duct, respectively, and impairment of renal concentrating ability (20, 21). There was no difference in plasma Ca²⁺ levels between KO and control animals (Table 1). The urinary excretion rates (absolute excretion, E_{Ca} , and fractional excretion, FE_{Ca}), however, were significantly increased by 3.0- and 3.3-fold, respectively. Because KO animals excreted more water than calcium, the urinary calcium concentration was in fact lower in KO than in control animals. Thus, loss of function in ILDR1 can cause primary renal concentrating defect in mice.

Microperfusion of Thick Ascending Limb Tubules from ILDR1 KO Mice.

Since the late 1950s, the countercurrent multiplication hypothesis has been the widely accepted explanation for the generation of the osmolality gradient along the cortico-medullary axis, which in turn drives water reabsorption in the collecting duct through Aqp2 (22). This hypothesis holds that the osmotic pressure difference between the lumen and the interstitium generated by the thick ascending limb (TAL) of the loop of Henle is multiplied because the fluid flows in the opposite direction from the thin descending limb to the TAL. The driving force of this process depends upon the effective separation of salt reabsorption from water impermeation in the TAL, which creates the osmotic pressure difference that can then be multiplied. The TAL was previously considered water impermeable. We performed ex vivo perfusion experiments on microdissected TAL tubules from control and ILDR1 KO animals (Fig. 24). To record the water permeability across the microperfused tubule, a high molecular weight fluorescent dye (FITC-dextran, 150 kDa, 30 μM) was included in the perfusate (Table S1), which was too large to permeate through the epithelium. Then, the open end of the microperfused tubule was sealed, allowing stepwise increases in basolateral osmolality to generate an outward osmotic gradient that drives water to leave the tubule. The loss of water would be inversely proportional to increases in luminal fluorescence intensity (Fig. 2B). Consistent with the requirement of the countercurrent multiplication theory, the microperfused TAL tubules in control mice were impermeable to water, reflected by a nearly flat fluorescence intensity curve. The microperfused TAL tubules from KO mice, in contrast, showed stepwise increases in luminal fluorescence intensity (Fig. 2B). The fluorescence increment, measured as Δ fluorescence intensity per minute, reflected the water permeation velocity. As shown in Fig. 2C, the water permeation velocity for each 50 mOsm/kg step was below 0.01 in control TAL tubules whereas KO TAL tubules had a maximum of more than eightfold higher water permeation velocity. The inner medullary interstitial osmolality was significantly reduced in the KO mouse kidney (Table 1). Together, these results establish that deletion of ILDR1 in the kidney dissipates the cortico-medullary osmolality gradient due to increased water permeability in the TAL.

The Paracellular Ionic Permeability Remains Intact in ILDR1 KO Mice.

Our previous studies of claudins [e.g., claudin-16 and claudin-19 from an inherited human disease—familial hypomagnesemia and hypercalciuria with nephrocalcinosis (FHHNC) (OMIM 248190)] have

Table 1. Renal metabolism in ILDR1 KO and control mice

Physiologic parameters	Group		Significance (n = 10)
	Control	KO	
Weight, g	34.27 ± 1.48	27.26 ± 0.61	<i>P</i> < 0.01
UV, μL/24 h per g	49.47 ± 5.73	287.80 ± 38.22	<i>P</i> < 0.01
Water Intake, mL/24 h per g	0.14 ± 0.02	1.26 ± 0.12	<i>P</i> < 0.01
Osm, Osm/kg H ₂ O	1.46 ± 0.12	0.39 ± 0.05	<i>P</i> < 0.01
Inner medullary Osm, Osm/kg H ₂ O	0.76 ± 0.08	0.32 ± 0.07	<i>P</i> < 0.01 (n = 5)
C _{Cr} , mL/24 h per g	4.52 ± 0.83	3.84 ± 0.57	n.s.
Mean BP, mmHg	103.4 ± 2.2	98.6 ± 2.3	n.s.
P _{Ca} , mmol/L	2.12 ± 0.05	2.09 ± 0.04	n.s.
P _{Na} , mmol/L	145.45 ± 1.78	149.72 ± 1.25	n.s.
P _{Cl} , mmol/L	107.22 ± 0.57	108.68 ± 0.59	n.s.
P _K , mmol/L	3.58 ± 0.29	3.82 ± 0.44	n.s.
FE _{Ca} , %	0.70 ± 0.21	2.34 ± 0.59	<i>P</i> < 0.05
E _{Ca} , μmol/24 h per g	0.056 ± 0.018	0.167 ± 0.039	<i>P</i> < 0.05
FE _{Na} , %	0.93 ± 0.18	1.13 ± 0.22	n.s.
E _{Na} , μmol/24 h per g	5.14 ± 0.78	5.39 ± 0.97	n.s.
FE _{Cl} , %	1.25 ± 0.25	1.36 ± 0.28	n.s.
E _{Cl} , μmol/24 h per g	5.01 ± 0.70	5.32 ± 1.04	n.s.
FE _K , %	20.36 ± 4.21	30.74 ± 8.82	n.s.
E _K , μmol/24 h per g	2.61 ± 0.27	3.13 ± 0.49	n.s.

E_K, absolute excretion rate of K⁺; FE_K, fractional excretion rate of K⁺; n.s., not significant; P_{Ca}, plasma concentration of Ca⁺⁺; P_K, plasma concentration of K⁺.

established that the tight junction, in particular the bicellular tight junction, is responsible for paracellular cation permeation (23). To verify that changes in tTJ permeability were not byproducts of changes in bTJ, we recorded the electrophysiological profile of microperfused TAL tubules from control and ILDR1 KO mice, including the transepithelial voltage (V_{te}), the transepithelial resistance (R_{te}), the equivalent short-circuit current (I_{sc}), and the permeability ratio (P_{Na}/P_{Cl}), as described before (24). As shown in Fig. 2 D–G, none of these parameters differed between KO and control animals, indicating that the bTJ permeabilities to Na⁺ and Cl[−] were not affected by tTJ alteration. Further immunostaining of ZO-1, as an indicator of bTJ integrity, and claudin-16 and -19, as indicators of bTJ functionality, and NKCC2 as an indicator of transcellular ionic permeability, showed no difference at all between KO and control TAL tubules (Fig. S3). The tTJ integrity, assessed by tricellulin localization, remained intact in KO TAL tubules (Fig. S4). Together, these results indicate that the gross tight junction barrier function is not affected by the deletion of ILDR1 in tricellular junction.

Vasopressin Cannot Correct Paracellular Water Loss Despite Normal Transcellular Water Response. From Fig. 1, we learned that ILDR1 is also expressed in the collecting duct. In microperfused collecting duct tubules (Fig. 3A), luminal fluorescence intensity was significantly higher in KO than in control animals when they were given free access to water (Fig. 3B), indicating that the paracellular water pathway also existed in the collecting duct. The water permeation velocity, as determined by luminal fluorescence increment, was approximately threefold higher in KO than in control collecting duct tubules (Fig. 3C). To verify that the water loss in the collecting duct was due to the paracellular pathway but not through the transcellular pathway, we ascertained the gene expression (Fig. S5A), the protein abundance (Fig. S5B), and the membrane trafficking (Fig. S5C) of the major aquaporin in the collecting duct—aquaporin 2 (Aqp2) from KO and control mouse kidneys. Compared with the control level, none of these above parameters significantly differed in the KO kidneys. To determine whether vasopressin regulation of water permeation remained intact in the collecting duct, we treated the microperfused collecting duct tubules *in vitro* with a 50-nM dose of vasopressin. In both control and KO collecting duct tubules, vasopressin elicited a sharp increase in

luminal fluorescence intensity (Fig. 3B). The maximal water permeation velocity after vasopressin stimulation was not significantly different between KO and control collecting duct tubules (Fig. 3C). To address whether vasopressin effect on urinary concentration in live KO animals may be compromised, we infused vasopressin to both groups of animals at the rate of 0.6 μg·day^{−1}·kg BW^{−1} via s.c. pump for 24 h. As shown in Fig. 3D, although vasopressin treatment significantly increased the urinary osmolality in control and KO mice, the KO mouse urine remained significantly more hypotonic than the control mouse urine. Finally, we asked whether vasopressin secretion might be affected in the KO animal during water restriction. The baseline plasma vasopressin levels were similar between control and KO mice (Fig. 3E). Water deprivation for 12 h significantly increased the plasma vasopressin levels in control and KO animals by 2.0- and 2.3-fold, respectively (Fig. 3E). The urinary osmolality was increased in both groups of animals after water deprivation (Fig. 3F). Nevertheless, the KO mouse urinary osmolality was still significantly lower than that of the control mouse urine (Fig. 3F). Together, these results suggest that the renal water loss in ILDR1 KO mouse kidney is specifically due to the paracellular pathway defect.

The Colonic Water Loss in ILDR1 KO Mice. Because the fold of increase in water intake volume was higher than that in urinary output volume (Table 1), we asked whether there might exist any route of water loss in addition to the kidney. The intestine plays important roles in water absorption or secretion. We first addressed where the ILDR1 protein was localized in the intestine. The mouse small intestine (including the duodenum, the jejunum, and the ileum) was without any detectable expression of ILDR1 protein (Fig. S6). The colonic epithelium, by contrast, expressed the ILDR1 protein exclusively in the tricellular tight junction (Fig. 4A). Then, we analyzed the fecal content of age-matched (3-mo old) and sex-matched (male) ILDR1 KO and control mice in metabolic cages for 24 h. The fecal water volume excreted in the stool was 2.1-fold higher in KO than in control animals (Fig. 4B). Neither the dry stool weight (Fig. 4C) nor the food intake level (Fig. 4D) differed between KO and control animal groups. Knowing that colon inflammation may also cause excessive water loss, we performed histologic examination on age- and sex-matched ILDR1 KO and control mouse

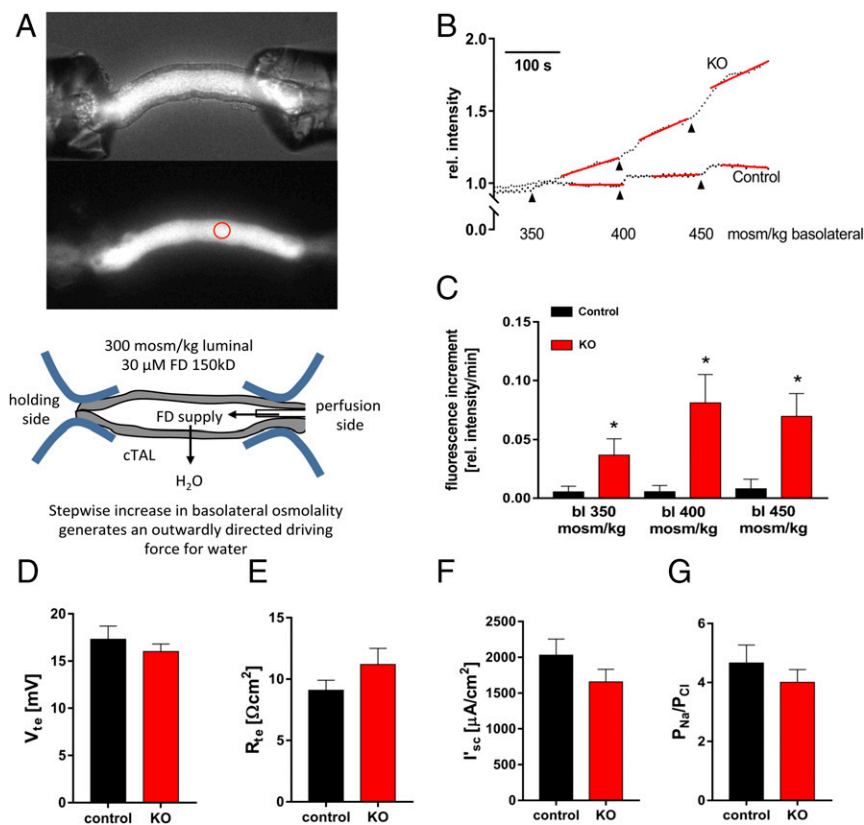


Fig. 2. Ex vivo microperfusion of thick ascending limb from ILDR1 KO kidney. (A) Light microscopy demonstration of the microperfusion system with live TAL tubules in perfusion with 150 kDa FITC-dextran in the lumen. The luminal fluorescence intensity was measured continuously at the focal point depicted by the red circle. (B) Real-time luminal FITC fluorescence intensity recording after stepwise increases in basolateral osmolality. (C) Fluorescence increment values as a direct measure of water permeation velocity in KO and control TAL tubules. * $P < 0.05$, $n = 6$ animals, data are given as mean \pm SEM of nine and six perfused TAL. (D) Transepithelial voltage V_{te} measurement in KO and control TAL tubules. (E) Transepithelial resistance R_{te} measurement in KO and control TAL tubules. (F) Equivalent short-circuit current I'_{sc} measurement in KO and control TAL tubules. (G) Paracellular permeability ratio P_{Na}/P_{Cl} in control and KO TAL tubules. (D–G) Data are shown of 10 to 14 individual TAL tubules, $n = 6$ animals.

colons. Similar to the control mouse colon, the KO mouse colon was morphologically normal and showed no sign of leukocyte infiltration or epithelial necrosis (Fig. S7). Together, these results indicate that the paracellular water pathway may have a universal role in many epithelia including the colonic and the renal epithelia.

ILDR1 Inhibits Transepithelial Water and Solute Permeability. Rosenthal et al. previously have demonstrated that a putative paracellular water channel—claudin-2—mediates transepithelial water permeation in MDCK-II cells (3). We found that endogenous

ILDR1 expression in MDCK-II cells was below the detection limit by immunoblotting or immunofluorescent labeling (Fig. S8). Transfection via retrovirus generated stable and homogenous overexpression of ILDR1 protein in the tricellular tight junctions of MDCK-II cells (Fig. S8). The claudin-2 protein expression or localization was not affected by the ectopic expression of ILDR1 (Fig. S8). Using a semiperfused diffusion chamber system (Materials and Methods), we measured the transepithelial water permeability in MDCK-II cells expressing ILDR1. A high molecular mass fluorescent dye (FITC-dextran, 150 kDa, 1 μM) was

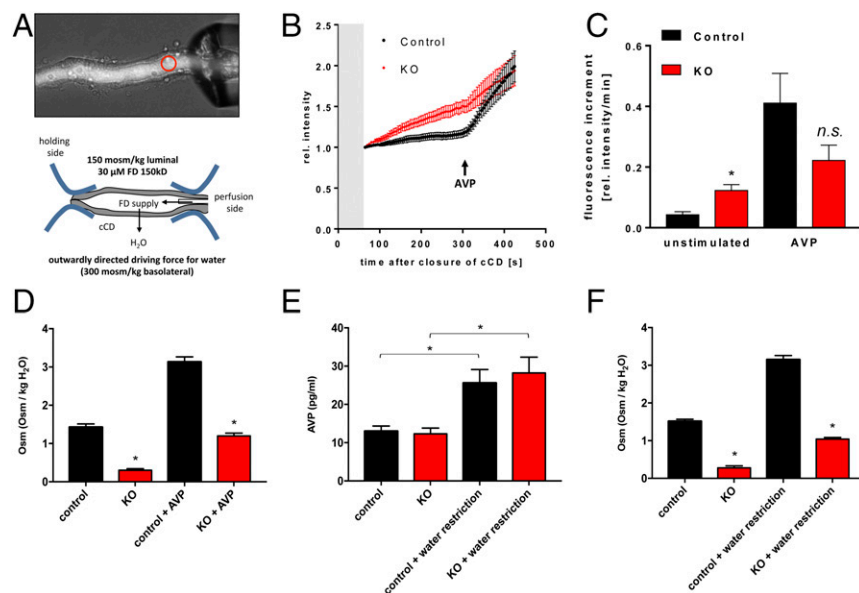


Fig. 3. Ex vivo microperfusion of collecting duct from ILDR1 KO kidney. (A) Light microscopy demonstration of the microperfusion system with live collecting duct tubules in perfusion with 150 kDa FITC-dextran in the lumen. The luminal fluorescence intensity was measured continuously at the focal point depicted by the red circle. (B) Real-time luminal FITC fluorescence intensity recording after a luminal fluid exchange (gray area) to lower osmolality. AVP was added at 300 s. (C) Fluorescence increment values as a direct measure of water permeation velocity in KO and control collecting duct tubules in the absence or presence of vasopressin. * $P < 0.05$, $n = 6$ animals, data are given as mean \pm SEM of 10 and 8 CDs. (D) The 24-h urinary osmolality levels in KO and control animals in the absence or presence of vasopressin infusion. * $P < 0.05$, $n = 5$ animals. (E) Plasma vasopressin levels in KO and control animals with free access to water or under 12-h water deprivation. * $P < 0.05$, $n = 5$ animals. (F) The 24-h urinary osmolality levels in KO and control animals with free access to water or under 12-h water deprivation. * $P < 0.05$, $n = 5$ animals.

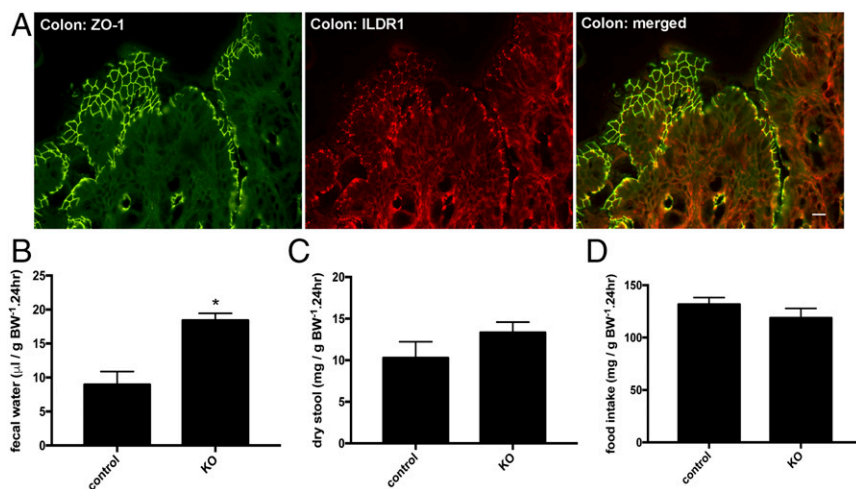


Fig. 4. ILDR1 gene expression and physiology in the colon. (A) Double staining of ILDR1 protein with a tight junction marker (ZO-1) in the mouse colon. (Scale bar: 10 μm .) (B) Fecal water volume in KO and control animals. $*P < 0.05$, $n = 6$ animals. (C) Dry stool weight in KO and control animals. $n = 6$ animals. (D) Food intake levels in KO and control animals. $n = 6$ animals.

included in the apical solution. Then, the basolateral perfusate osmolality was increased by 100 mOsm/kg H₂O to generate an apico-basal osmotic gradient. The increase in apical fluorescence intensity would be inversely proportional to the water loss. The apical fluorescence increment, measured as Δ fluorescence intensity per minute, reflected the water permeability velocity. Compared with the baseline MDCK-II level, ILDR1 expression significantly decreased the water permeation velocity by 35% (Fig. 5A). The transepithelial permeability of fluorescein (332 Da) was reduced by 73% of the baseline MDCK-II level in ILDR1-expressing cells (Fig. 5B) whereas neither the paracellular Na⁺ permeability (P_{Na}) (Fig. 5C) nor the paracellular Cl⁻ permeability (P_{Cl}) (Fig. 5D) was significantly altered by ILDR1 expression. Together, these results demonstrate that ILDR1 regulates paracellular water and solute permeability with no effect on paracellular ionic permeability.

Discussion

Permeability Property of the Tricellular Junction. In freeze fracture replica electron micrographs, the bTJs are discontinuous at tricellular contacts (25, 26). The tTJ is composed of three pairs of TJ strands arranged vertically and known as the central sealing element (5). Unlike the bTJ, which contains ion channels of 4 to 7 Å in diameter (1), the tTJ is predicted to create a paracellular pathway with much a larger diameter—~10 nm surrounded by the central sealing element (5). Such a large paracellular pathway would be deemed suitable for free water (~3 Å in diameter) permeation as well as for permeation of other solutes such as fluorescein (~9 Å in diameter). The water permeability of tTJ is likely to have important roles in many epithelia. Notably, the combined water excretion through the kidney and the colon accounted for only 20% of the increased water intake by the ILDR1 KO animal. Insensible water loss through the skin or the lung may also be increased in the absence of the ILDR1 protein. Judging from angulin-2/ILDR1's barrier function, other angulins may regulate the tTJ permeability in a similar way to ILDR1. For example, angulin-1/LSR is important in regulating the blood-brain barrier permeability. Deletion of LSR in brain vasculature increased the tTJ permeability to biotin but not to albumin (27). The organization of the tricellular junction has been an interesting but understudied topic. By studying ZO-1-deficient MDCK cells, Choi et al. have proposed a model that the bicellular junctions are anchored “end-on” to the tricellular junctions (28). Compatible with this theory, knockdown of tricellulin in Eph4 cells rendered bicellular junctions discontinuous (7). Molecular interaction assays revealed that tricellulin could *cis*-interact with claudins to facilitate the anchoring of bTJ onto tTJ

(29). It is not known whether ILDR1 may interact with claudins; e.g., claudin-2 to regulate their permeabilities. In mouse kidneys and in MDCK-II cells, manipulation of ILDR1 expression seemed to exert no effect on bTJ structure and function, which would suggest that ILDR1 is a tTJ-specific modulator. The barrier function of ILDR1 may derive from the *trans*-interaction between its extracellular immunoglobulin-like domain, in a scenario similar to a class of bTJ protein—JAMs (30).

Paracellular Water Permeability and Urine Concentration Mechanism.

Polyuria with the production of an excessive amount of isotonic urine, known as isosthenuria, can be seen as a type of DI phenotype derived from an impaired water dilution mechanism in the loop of Henle. The two extreme states of water homeostasis that the kidney can handle are depicted in Fig. S9. Under both circumstances, the TAL has to dilute the luminal fluid and to fuel the countercurrent multiplication mechanism. To produce concentrated urine (Fig. S9, *Left*), the collecting duct has, in a second step, to allow water permeation and therefore equilibration of the luminal osmolality to the basolateral osmolality

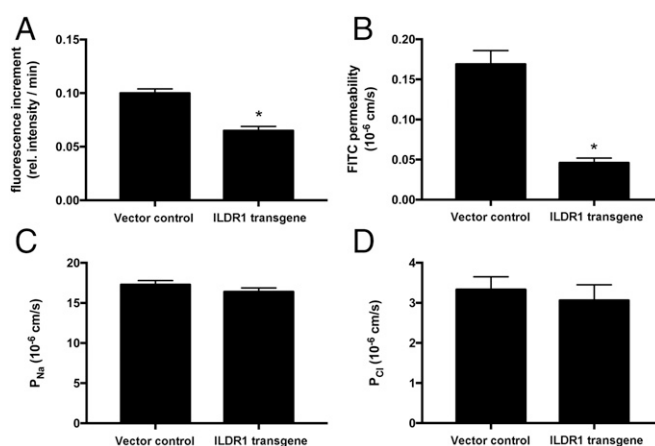


Fig. 5. In vitro recording of ILDR1 function in MDCK-II cells. (A) Transepithelial water permeability in MDCK-II cells expressing an empty vector control or ILDR1 transgene, as determined by the fluorescence increment. $*P < 0.05$, $n = 5$ clones. (B) Transepithelial FITC permeability in MDCK-II cells expressing control or ILDR1 transgene, as determined by the flux assay. $*P < 0.05$, $n = 5$ clones. (C and D) Paracellular Na⁺ permeability (P_{Na} , C) and paracellular Cl⁻ permeability (P_{Cl} , D) in MDCK-II cells expressing control or ILDR1 transgene, as determined by the transepithelial electrical resistance (TER) and dilution potential (DP) measurements. $n = 5$ clones.

(water will be transported away continuously by the vasa recta, also part of the countercurrent mechanism). This process is under the tight control of vasopressin and is defective in DI. Urine produced by DI patients therefore resembles the dilute urine produced from the end of the TAL. If the water permeability of the TAL is increased, as is the case in ILDR1 KO mice, then the TAL is not diluting the luminal fluid properly to values below plasma tonicity, no hypotonic urine can be produced, and the interstitial osmotic gradient dissolves. The end result is no urinary concentration beyond isosthenuria. In mice, vasopressin challenge may still increase the urea transport in the collecting duct, which will partially increase the interstitial osmolality in the inner medulla and concentrate the urine to a certain degree. In the TAL, we also investigated paracellular ionic permeability properties, in addition to paracellular water permeability, and could not detect any difference although ions could likely permeate through the leakier tTJ. The bTJ already confers a huge ionic permeability, considering the typical meandering pattern of the bTJ in the segment. Some additional tTJ ionic permeability would be invisible under these conditions, a finding shown in a similar way in MDCK-II cell culture. In contrast, paracellular water permeability was clearly visible in the otherwise water-impermeable TAL. The increased water permeability across the TAL tight junction would suggest that the luminal fluid in the TAL remains isotonic to the serum. Generation of the lumen-positive diffusion potential (V_{te} in the TAL), however, depends upon the NaCl concentration difference between the lumen and the peritubular space (equal to the serum) (31). V_{te} , serving as a major driving force for the reabsorption of divalent cations (32), is therefore expected to diminish in the ILDR1 KO mouse

TAL, causing increases in renal calcium excretion. The elevated luminal tonicity in the KO mouse TAL would also be expected to activate the tubuloglomerular feedback (TGF) mechanisms leading to a reduction in the GFR. Although our initial estimation of GFR based upon the creatinine clearance rate did not reveal a significant reduction, more accurate recording approaches, such as the single nephron glomerular filtration rate (SNGFR) measurement, will be needed to interrogate the real-time TGF response in each individual nephron.

In conclusion, we provide insights into tricellular junction permeability by showing that ILDR1 can function as a water barrier in the tricellular tight junction. Manipulating tricellular junction permeability may represent a mechanism to regulate epithelial water homeostasis.

Materials and Methods

All mice were bred and maintained according to the Washington University animal research requirements, and all procedures were approved by the Institutional Animal Care and Use Committee. *SI Materials and Methods* includes additional topics on animal metabolic studies, analyses of inner medullar osmolality, molecular cloning and retrovirus production, renal tubule perfusion, and transepithelial water permeability measurements.

ACKNOWLEDGMENTS. We thank Dr. Rodger Liddle and Dr. Rashmi Chandra (Duke University) for kindly providing the ILDR1 knockout mouse model and the ILDR1 antibody; Dr. Lushan Zhou and Dr. Lane Baker (Indiana University) for assistance with electrophysiological recording; and Dr. George Jarad for assistance with mouse intestinal dissection. This work was supported by National Institute of Diabetes and Digestive and Kidney Diseases Grant R01DK084059 and American Heart Association Grant 17SDG33410806.

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