Modulation of Intestinal Disorders by Claudin-2 and Occludin Via Canonical and Noncanonical Mechanisms



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Abstract The past 35 years, beginning with the identification of ZO-1 in 1986, have been an exciting time during which critical tight junction proteins were discovered. We have, however, only begun to define the mechanism by which tight junctions are regulated, their impact on health and disease, and noncanonical functions of individual tight junction-associated proteins. Here, we provide an overview of advances in understanding mechanisms of tight junction barrier regulation within the intestinal epithelium and discuss recent discoveries related to claudin-2 and occludin in greater detail. We anticipate that the next 35 years will yield major advances in fundamental understanding of tight junction protein interactions, regulation, and canonical and noncanonical functions that result in translational applications in which tight junction modulation is established as a therapeutic approach.

Keywords Intestinal permeability \cdot Intestinal barrier \cdot Tumor necrosis factor \cdot IL-13 \cdot IL-22 \cdot Inflammatory bowel disease \cdot Tight junction

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Abbreviations

CK2	Casein kinase 2
DSS	Dextran sulfate sodium
FRAP	Fluorescence recovery after photobleaching
MLC	Myosin II regulatory light chain
MLCK	Myosin light -chain kinase
TER	Transepithelial electrical resistance
TNBS	Trinitrobenzenesulfonic acid
TNF	Tumor necrosis factor

Introduction

The viability of multicellular life forms requires the presence of barrier-forming tissues that partition distinct compartments within the organism and separate the external environment from the internal milieu. In mammals, examples of surfaces that interact with the external environment include the skin as well as oral, gastrointestinal, and genitourinary mucosae. At each of these sites, the tight junction creates a selectively permeable seal that restricts flux across the paracellular space between cells. Tight junctions in the skin are nearly impermeant, and, in burn patients, the extent of epidermal barrier loss is a strong prognostic indicator of outcome. The gastrointestinal epithelium, particularly within the distal gut, interfaces with a dense and complex luminal microbiome that can include invasive pathogens as well as potentially harmful bacterial components and metabolites. Dysregulated systemic exposure to these luminal materials, which could lead to immune-mediated, metabolic, and infectious diseases, emphasizes the essential nature of the intestinal epithelial barrier. However, in contrast to the skin, the intestinal barrier must allow paracellular flux of water, ions, and nutrients. The charge and size selectivity of this barrier are determined by the tight junction, which is the rate-limiting step in paracellular flux.

Tight Junctions Are Regulated by Pharmacological and Physiological Stimuli

Farquhar's descriptions of the tight junction and apical junctional complex were remarkably consistent across samples and tissues [1]. Together with other data, this led many to conclude that tight junctions are static structures with unchanging permeabilities. Ussing's report [2], only 1 year after Farquhar's landmark publication, that application of an osmotic gradient could markedly increase paracellular flux across was interpreted as tight junction disassembly. Subsequent studies showing that osmotic gradients reduced [3], while phalloidin and proliferative stimuli increased [4, 5], the complexity of strand networks were generally dismissed as non-physiological. This changed in 1987 when Pappenheimer et al. published a series of papers demonstrating that increased intestinal epithelial tight junction permeability was a physiological response to Na⁺-nutrient cotransport [6–8]. These first studies showing physiological tight junction regulation also led to the discovery of "solvent drag" [8–10], a mechanism by which paracellular flux allows continued nutrient absorption despite saturation of transcellular transport pathways. Although initially controversial [11–15], subsequent in vitro and in vivo studies demonstrated that solvent drag occurs in many species, including humans [16, 17], as a result of increased paracellular amplification of transcellular transport in the renal tubule [21, 22].

Ex vivo studies of rodent mucosae showed that Na⁺-nutrient cotransport-induced tight junction permeability increases were associated with morphological changes including expansion of spaces between tight junction strands and condensation of perijunctional microfilaments [6]. This led to the hypothesis that actomyosin contraction might contribute to Na⁺-nutrient cotransport-induced tight junction regulation. However, methods available at that time precluded mechanistic analysis in intact mucosae. This obstacle was overcome by the development of a cell culture model of intestinal epithelial tight junction regulation in response to Na⁺-glucose cotransport [23]. Using a model facilitated the discovery that actomyosin contraction driven by myosin light chain kinase (MLCK)-mediated phosphorylation of myosin II regulatory light chain (MLC) was essential for physiological tight junction regulation. Moreover, enzymatic MLCK inhibition blocked Na⁺-glucose cotransport-induced tight junction regulation in cultured monolayers and isolated rodent mucosae [23]. Subsequent studies showed that Na⁺-glucose cotransport led to increased tight junction permeability and MLC phosphorylation in human intestinal mucosae [19]. Importantly, both in vitro and in vivo studies showed that these alterations were size selective in that mannitol (~7 Å diameter) permeability was increased, but there were no detectable changes in paracellular inulin (30-40 Å diameter) flux [20, 23]. Thus, MLCK activation by physiological stimuli is a canonical mechanism of epithelial tight junction regulation.

Tight Junctions Are Differentially Regulated by Inflammatory Stimuli

The discovery of MLCK as the mediator of Na⁺-glucose cotransport-induced permeability increases provided the first mechanistic insight into physiological tight junction regulation. This also created opportunities to discover mechanisms of pathophysiological barrier regulation. Of these, the most well-established is the discovery that MLCK also mediates tight junction regulation induced by proinflammatory cytokines, including TNF, LIGHT, and IL-1ß [24-26]. The efficacy of anti-TNF therapies and their ability to restore barrier function in patients highlights the clinical relevance of this discovery, although efficacy is more likely due to immunomodulation [27, 28]. Experimentally, barrier loss induced by TNF was completely reversed by a specific membrane-permeant inhibitor of MLCK (PIK). MLC phosphorylation, which was increased following TNF treatment, was also corrected by PIK [29]. PIK was, however, only able to restore barrier function in response to modest degrees of barrier loss [29], indicating that the effect was due to reversal of TNF-induced tight junction regulation rather than inhibition of TNFinduced apoptosis [30-32]. Moreover, PIK or genetic epithelial MLCK inhibition each prevented acute, TNF-induced barrier loss and diarrhea in vivo [33]. Further, epithelial MLC phosphorylation and MLCK expression were found to be increased in human inflammatory bowel disease and to correlate with disease activity [34]. Thus, TNF and more complex inflammatory stimuli hijack physiological mechanisms of tight junction regulation to increase intestinal permeability. There are, however, differences between tight junction regulation by physiological Na⁺glucose cotransport and that induced by TNF in that only TNF increases permeability of molecules larger than small sugars [23, 30]. Thus, although both are driven by MLCK, the barrier loss induced by TNF is qualitatively different from that induced by Na⁺-glucose cotransport. This may reflect a second signal induced by TNF, as TNF, but not Na⁺-glucose cotransport, triggers occludin endocytosis in vivo [33, 34].

Once claudins were discovered and antibodies became available, several groups began to assess patterns of expression in health and disease [35-38]. These demonstrated markedly increased claudin-2 expression during active inflammatory bowel disease and showed that IL-13, which is elevated in both ulcerative colitis and Crohn's disease, was able to induce claudin-2 expression in cultured intestinal epithelial monolayers [36, 37]. Further study showed that, although IL-13, like TNF, reduced transepithelial electrical resistance (TER), only TNF increased paracellular permeability to 4 kDa dextran (28 Å diameter) and reduced charge selectivity [39]. In contrast to TNF, IL-13-induced barrier loss, was due to a claudin-2mediated increase in paracellular Na⁺ permeability and was unaffected by PIK [39]. In vivo studies confirmed that IL-13 was sufficient to increase cation permeability, and subsequent studies demonstrated that IL-13 was unable to affect paracellular permeability of claudin-2 knockout mice, while transgenic claudin-2 overexpression was sufficient to replicate the effects of IL-13 in the absence of cytokine treatment [40]. These data demonstrated that the IL-13 and TNF modify paracellular permeability by different mechanisms, claudin-2 upregulation and MLCK activation, respectively, that have distinct functional effects.

Distinct Tight Junction Pore and Leak Pathways

In parallel with the studies above, analyses of genetically modified MDCK monolayers showed that claudin-2 expression increased paracellular permeability of small molecules across pores with diameter of \sim 8 Å but had no effect on



Fig. 1 The intestinal barrier. The space between epithelial cells is sealed by selectively permeable tight junctions that allow paracellular flux by two distinct pathways. The pore pathway (left) is a size- and charge-selective, high-conductance pathway that accommodates molecules up to 6–8 Å in diameter. Upregulation of claudin-2, which forms paracellular channels, increases pore pathway permeability. The mucosal immune system can induce claudin-2 transcription via IL-13 or IL-22 secretion. The leak pathway (center) is a charge-nonselective, low-capacity macromolecular flux route that accommodates molecules up to 100–125 Å in diameter. TNF and IL-1 β are each able to increase leak pathway permeability by upregulating myosin light chain kinase (MLCK) expression and activity and recruiting the MLCK1 splice variant to the perijunctional actomyosin ring. This triggers removal of occludin by caveolar endocytosis and leads to increased leak pathway flux. A potential third route, the unrestricted pathway (right panel), is tight junction-independent, size-nonselective, and charge-nonselective. It is created by epithelial damage and can become the dominant route of intestinal barrier loss

paracellular flux of larger probes [41]. In contrast, knockdown of either ZO-1 or occludin increased paracellular permeability to larger macromolecules [42, 43]. Specifically, occludin knockdown increased flux across a pathway that accommodates macromolecules up to ~ 125 Å in diameter [43]. Thus, TNF treatment or occludin knockdown increased macromolecular permeability and reduced charge selectivity, while IL-13 treatment or claudin-2 overexpression increased cation selectivity without affecting macromolecular permeability. This confluence of observations led to the hypothesis that two distinct pathways, termed pore and leak, mediate paracellular flux (Fig. 1) [44, 45]. The pore pathway, exemplified by claudin-2 channels, is a high-conductance route that is both charge- and size-selective, with a maximum diameter of 6 to 8 Å. The low-capacity leak pathway is sizeselective, with an estimated maximum diameter of 100 to 125 Å and is not chargeselective. In contrast to the pore pathway, which is created by claudin-based, gated channels [46], the anatomy of the leak pathway has not been defined; some have suggested that leak pathway flux may occur primarily at tricellular tight junctions [47].

The increases in intestinal epithelial tight junction permeability induced by Na⁺nutrient cotransport to amplify transcellular nutrient absorption are chargenonselective, permit flux of molecules too large to traverse claudin channels, and are regulated by the cytoskeleton [8, 19, 23, 48]. These changes, therefore, reflect leak pathway flux. In an analogous fashion, claudin-2-mediated pore pathway flux allows paracellular Na⁺ absorption to amplify transcellular Na⁺ absorption in the renal proximal tubule [21]. Thus, defined flux across both pore and leak pathways is of physiological benefit.

Claudin-2: A Paracellular Cation and Water Channel

In normal human and rodent development, intestinal epithelial claudin-2 expression is highest in the immediate perinatal period, where it is expressed in villous and crypt epithelial cells [35, 49, 50]. Claudin-2 expression is then progressively down-regulated until, by weaning, expression is low and restricted to crypt epithelial cells [35, 49, 50]. Claudin-15, which also forms a paracellular cation and water channel [51–55], has a reciprocal temporal expression pattern and is present at low levels prior to weaning but is expressed at higher levels in both villous and crypt epithelia by adulthood [35, 49, 50].

Although the reasons for developmental regulation of claudin-2 expression have not been defined, it may reflect differences in nutrient demand, which is greatest during the rapid growth that characterizes the neonatal and pre-weaning periods. This is best understood by recognizing that expression of claudin-2 or claudin-15 is required for ongoing Na⁺-nutrient cotransport; mice lacking either claudin survive while those lacking both die of malnutrition by 3 weeks of age [56]. It is also notable that paracellular Na⁺ conductance is significantly greater in neonatal mice, relative to adults [49]. Thus, it may be that claudin-2 is able to more effectively conduct Na⁺ than claudin-15. This has not, however, been demonstrated experimentally. In the context of inflammatory stimuli, including disease, intestinal epithelial claudin-2 expression is upregulated, while claudin-15 expression is largely unchanged (Fig. 2) [36, 39, 40, 50, 57–60]. This could therefore represent a compensatory mechanism that restores nutrient absorption in the face of mucosal damage. However, the observation that claudin-2 is not upregulated in claudin-15-deficient mouse intestine [61] suggests that alternative explanations should also be considered.

Claudin-2-Dependent Fluid Efflux Promotes Pathogen Clearance

One major limitation of most in vivo studies of permeability is that the probe or probes used do not allow discrimination between changes in pore, leak, and damageinduced (unrestricted pathway) permeability. Lactulose and mannitol, the most



Fig. 2 Regulation of claudin-2, claudin-15, and occludin expression in healthy and diseased adult human intestine. Claudin-2 (green) is primarily expressed in the crypts of the small intestine and colon, and this is increased in inflammatory disease. In contrast, expression of claudin-15 (green), which also forms paracellular cation-selective pores, is present throughout the crypt-villus axis and is unchanged in disease. Occludin (green) expression is reduced in inflammatory disease. E-cadherin (magenta), ZO-1 or γ -actin (red), and nuclei (blue) are shown for reference. Bar, 50 µm; 20 µm (insets)

extensively used probes in human studies, are both able to traverse leak and unrestricted pathways. Mannitol is too large to be accommodated by claudin-2 channels in vitro, but it remains to be determined if it can cross the pore pathway in vivo [41]. In mouse models, most investigators simply use 4 kDa FITC-conjugated dextran. Serum recovery of this probe can reflect increased permeability across both leak and unrestricted, but not pore, pathways [62]. Experimental data suggest that greater than two-fold increases in 4 kDa dextran flux are most likely to reflect unrestricted pathway activation as consequence of epithelial damage [62]. To overcome this limitation, we established a method using creatinine (6 Å diameter), 4 kDa dextran (28 Å diameter), and 70 kDa dextran (120 Å diameter), to measure flux across pore, leak, and unrestricted pathways, respectively [60, 62]. This approach, which has been validated by other groups [63], readily distinguishes between barrier defects induced by *C. rodentium* infection, TNF, and DSS [62].

Using the three-probe in vivo permeability assay, we discovered that creatinine flux is uniquely increased 2 days after *C. rodentium* infection in mice [60]. This correlated with increased epithelial expression of claudin-2, but not other tight junction proteins, and elevated mucosal IL-22 levels [60]. Analyses using recombinant IL-22 demonstrated that this member of the IL-10 superfamily is responsible for the increased intestinal epithelial claudin-2 transcription and translation seen after *C. rodentium* infection.

In order to understand the impact of claudin-2 upregulation in *C. rodentium* infectious colitis, we took advantage of claudin-2 knockout and claudin-2 transgenic mice [60]. The very low claudin-2 expression observed in healthy adults was similar to the absence of claudin-2 in knockout mice, while the increased expression in claudin-2 transgenic mice was comparable to levels detected at the peak of *C. rodentium* infection [60]. The genetically modified mice therefore model the extremes of claudin-2 expression in wild-type mice.

The severity of C. rodentium colitis was exacerbated by claudin-2 knockout, as indicated by increased histopathology scores, increased mucosa-associated C. rodentium, prolonged C. rodentium shedding, and markedly elevated mucosal TNF, IL-1β, IL-6, and IL-22 (Fig. 3) [60]. The observation that fecal water and Na⁺ were increased in healthy claudin-2 transgenic mice suggested that the greater severity of disease in claudin-2 knockout mice might be secondary to reduced water and Na⁺ efflux. Consistent with this hypothesis, induction of mild osmotic diarrhea by adding polyethylene glycol to the drinking water reduced histopathology scores, mucosa-associated C. rodentium, fecal C. rodentium shedding, and mucosal TNF, IL-18, IL-6, and IL-22 levels such that all measures of disease were similar in polyethylene glycol-treated claudin-2 knockout, transgenic, and wild-type mice [60]. These data suggest that the primary means by which claudin-2 upregulation accelerates pathogen clearance and reduces infectious enterocolitis severity is by promoting paracellular water and Na⁺ efflux [60]. Because epithelial cell turnover is accelerated in C. rodentium-induced colitis, this cannot be simply due to bacterial washing from the epithelial surface but is more likely to reflect disruption of the cycle by which newly formed epithelial cells are infected (Fig. 3a). The increased fecal water resulting from claudin-2 overexpression likely also dilutes and reduces local DSS concentrations to attenuate mucosal damage, thereby explaining the protection fromm DSS-induced injury observed in claudin-2 transgenic mice [64]. Thus, claudin-2 increases luminal water efflux to promote pathogen clearance and dilute offending chemical agents in infectious and chemical colitis, respectively. Intestinal claudin-2 upregulation and the water efflux that follows can therefore be considered a primitive form of innate immunity.



Fig. 3 IL-22 upregulates epithelial claudin-2 to drive diarrhea and enteric pathogen clearance. (a) IL-22 released in response to enteric infection upregulates claudin-2 expression to enhance paracellular Na⁺ and water efflux and promote pathogen clearance. Here, polyethylene glycol (PEG) is shown as an inducer of osmotic diarrhea. (b) *C. rodentium* (red) and nuclei (blue) 11 days after infection. Colonization of crypt spaces in claudin-2 knockout mice is prevented by PEG treatment. Bar, 50 µm. (c) Mucosa-associated *C. rodentium* numbers are reduced in *Cldn2^{Tg}* and increased in *Cldn2^{-/-}*, relative to wild-type (*Cldn2^{+/+}*), mice. PEG treatment eliminated differences between genotypes. (d) Histopathology was increased in *Cldn2^{-/-}* mice and reduced by PEG treatment. (e and f) Mucosal TNF and IL-22 were markedly greater in *Cldn2^{-/-}* mice relative to wild-type or *Cldn2^{Tg}* mice, at day 11 after infection in the absence, but not presence, of PEG treatment. (Data from Tsai et al. Cell Host Microbe, 2017, with permission)

Claudin-2 Exacerbates Immune-Mediated Colitis

Although claudin-2 upregulation reduces severity of infectious and chemical colitis, it is not clear if it represents a beneficial adaptation or contributes to progression of immune-mediated colitis. To address this, claudin-2 knockout and transgenic mice were crossed onto an immunodeficient *Rag1* knockout background and studied using the adoptive T-cell transfer colitis model [40, 65]. In contrast to infectious and chemical colitis models, immune-mediated disease severity was increased by transgenic claudin-2 expression and reduced by claudin-2 knockout, as measured by weight loss, disease activity index, cytokine production, T-cell infiltration, and histopathology scores (Fig. 4) [40]. Conversely, the same measures showed that claudin-2 knockout ameliorated immune-mediated colitis [40]. It therefore appears that claudin-2 upregulation is a maladaptive response to mucosal immune activation in the absence of offending exogenous agents, such as pathogenic bacteria or toxic chemicals. Further study is needed to understand how claudin-2 modifies mucosal immune activation and whether this contributes to the relative protection afforded by claudin-2 overexpression in *C. rodentium* colitis.

Despite reduced disease severity, survival after T-cell transfer was reduced in claudin-2 knockout mice. This was unexpected and, in most cases, followed an unusual clinical pattern. Wild-type and claudin-2 transgenic mice displayed progressive increases in disease activity and weight loss over weeks. In contrast, claudin-2 knockout mice appeared well until they became acutely ill in the few days before death. Necropsy showed that these mice suffered from intestinal obstruction [40]. This was not due to fibrosis or dysmotility. We therefore considered the possibility that insufficient luminal hydration could result in fecalith formation and obstruction. In contrast to humans, this led to obstruciton in mice, where the luminal space is dramatically narrowed by mucosal inflammatory expansion. To prevent obstruction, polyethylene glycol was added to the drinking water of claudin-2 knockout and wild-type mice. This did not affect disease severity but improved survival of claudin-2 knockout mice [40]. Thus, increasing luminal water was sufficient to prevent obstruction and death in claudin-2 knockout mice. Notably, claudin-2 knockout, but not wild-type, mice upregulated intestinal epithelial claudin-15 expression during colitis. This can be taken as an additional piece of evidence that, despite our inability to detect functional differences at present [52], claudin-2 and claudin-15 are not functionally interchangeable, as such claudin-15 upregulation did not disrupt the effects of claudin-2 knockout.

Occludin S408 Phosphorylation Regulates Claudin-2 Channel Function

The occludin C-terminal tail can be separated into membrane-proximal and distal portions. The distal half (residues 373 to 522) is most highly conserved [66] and can be broken into an unstructured region (373–412) and a coil-coil domain (413–522) composed of three alpha helices [67]. Biochemical studies found that the



Fig. 4 Intestinal epithelial claudin-2 overexpression exacerbates, and claudin-2 knockout limits immune-mediated colitis severity. (**a**) Weight loss and disease activity induced by T-cell transfer in $Cldn2^{+/+}Rag1^{-/-}$ mice (blue circles), $Cldn2^{-/-}Rag1^{-/-}$ mice (red circles), and $Cldn2^{Tg}Rag1^{-/-}$ mice (green circles). (**b**) Histopathology of mice sacrificed at day 56 after T-cell transfer shows more severe disease in $Cldn2^{Tg}Rag1^{-/-}$ mice (and milder disease in $Cldn2^{-r}Rag1^{-/-}$ mice, relative to $Cldn2^{+/+}Rag1^{-/-}$ mice. (**c**) T-cell recruitment (CD3, green) parallels disease severity. ZO-1 or E-cadherin (red) and nuclei (blue) are shown for reference. Bars, 50 µm (Data from Raju et al. J Clin Invest, 2020, with permission)

unstructured region within the distal occludin tail contains multiple sites that can be phosphorylated by casein kinase 2 (CK2) [68–70]. The development of highly specific CK2 inhibitors as pharmacological therapies for neoplastic disease created an opportunity to assess the impact of CK2-mediated occludin phosphorylation on a barrier function. In Caco-2 intestinal epithelial cell monolayers, enzymatic CK2 inhibition reduced paracellular permeability of small cations, consistent with inhibition of the claudin-2 channel [71]. CK2 inhibitors had no effect in the absence of CK2, occludin, or claudin-2 expression [71]. Mutagenesis studies showed that serine-408 dephosphorylation was required for the effect of CK2 inhibition on barrier function. Fluorescence recovery after photobleaching (FRAP) studies showed that CK2 inhibition reduced the occludin mobile fraction and that this behavior also mapped to serine-408 [71]. Further analyses using pull-down assays showed that the non-phosphorylatable tail of occludin^{S408A} captured greater amounts of endogenous occludin, claudin-1, and claudin-2 from Caco-2 cell lysates than the phosphomimetic occludin^{S408D} [71]. Although no direct binding sites between claudins and occludin are known, ZO-1 includes PDZ1 and U5-GuK domains that bind to each of these, respectively [72, 73]. Consistent with an essential role for ZO-1-mediated interactions between occludin and claudins, neither S408A nor S408D occludin tails effectively recovered claudin-2 from ZO-1-deficient Caco-2 cell lysates [71]. Moreover, CK2 inhibitors had no effect on TER of ZO-1-deficient Caco-2 monolayers. The TER response to CK2 inhibition could be restored by expression of fulllength ZO-1, but not ZO-1^{Δ PDZ1} or ZO-1^{Δ U5GuK} [71]. Together, these data demonstrate that dephosphorylation of occludin^{S408} triggers assembly of an occludin-ZO-1claudin-2 complex that interferes with claudin-2 channel function (Fig. 5a) [73]. As a final test of this hypothesis, T84 cell monolayers, which do not ordinarily express significant levels of claudin-2, were treated with IL-13 to induce claudin-2 expression. This reduced TER and increased cation selectivity, and both of these changes could be reversed by CK2 inhibition. Thus, occludin phosphorylation at S408 is a molecular switch that, indirectly, regulates claudin-2 channels [71].

Fig. 5 (continued) channels. (b) IL-13 administration increased claudin-2 (green) expression in proximal colonic crypt epithelium. This was not affected by CK2 inhibition. Nuclei (blue) are shown for reference. (c) Ussing chamber analyses of proximal colonic mucosae from IL-13-treated mice (red squares) shows specific increases in permeability of small cations relative to vehicle-treated mice (blue circles). Treatment with CK2 inhibitor (green-outlined symbols) blocked IL-13-induced permeability increases but had no effect in the absence of IL-13. (d) Weight loss following T-cell transfer was attenuated in $Rag1^{-/-}$ mice treated with a CK2 inhibitor (green diamonds) relative to controls receiving vehicle (blue circles). (e) Histopathology was reduced by CK2 inhibitor treatment. (f) T-cell (CD3, green) infiltration was reduced by CK2 inhibition. E-cadherin (red) and nuclei (blue) are shown for reference. Bars, 50 µm (Data from Raju et al. J Clin Invest, 2020, with permission)



Fig. 5 CK2 inhibition limits immune-mediated colitis severity by inactivating claudin-2 channels. (a) Casein kinase-2 (CK2) inhibition results in occludin S408 dephosphorylation and triggers assembly of a trimolecular complex composed of occludin, ZO-1, and claudin-2 to inactivate claudin-2

Inhibition of Occludin Phosphorylation Is Therapeutic in Experimental, Immune-Mediated Colitis

In order to determine whether CK2 inhibition could also block claudin-2 channel function in vivo, mice were treated with saline or IL-13, to induce claudin-2 expression, in combination with a CK2 inhibitor or vehicle [40]. Bi-ionic potential measurements were then used to characterize excised proximal colonic mucosae [40, 53, 74, 75]. As expected, IL-13 specifically increased claudin-2 expression (Fig. 5b) as well as paracellular permeability to Na⁺, methylamine, and, to a lesser degree, ethylamine (Fig. 5c) [40]. CK2 inhibition failed to prevent increased claudin-2 expression but reversed changes in paracellular permeability (Fig. 5b, c). In contrast, CK2 inhibition had no effect on permeability in the absence of IL-13 treatment. CK2 inhibition is, therefore, able to block claudin-2 channel function in vivo.

Daily treatment with an orally bioavailable CK2 inhibitor, beginning 10 days after T-cell transfer, dramatically reduced immune-related colitis severity by all measures in claudin-2 wild-type, $Rag1^{-/-}$ mice [40]. In contrast, CK2 inhibition afforded no benefit to claudin-2 knockout $Rag1^{-/-}$ mice [40]. Thus, although the nearly ubiquitous expression and substrate promiscuity of CK2 make it impossible to fully exclude other targets, the requirement for claudin-2 expression suggests that the effect of CK2 inhibition reflects claudin-2 channel inactivation [40]. Thus, in vivo inhibition of CK2-mediated occludin phosphorylation prevents progression and attenuates severity of immune-mediated colitis [40]. CK2 inhibition can affect many processes and, therefore, is not an ideal therapeutic approach. However, future definition of how S408 phosphorylation regulates occludin tail interactions may lead to more specific means of inhibiting claudin-2 function.

Occludin Endocytosis Is Required for TNF-Induced, MLCK-Dependent Leak Pathway Permeability Increases

Remarkably, beyond the perijunctional actomyosin condensation seen by transmission electron microscopy, occludin internalization was the primary morphologic change associated with T-cell activation-induced, TNF-dependent barrier loss [33]. Further study showed that occludin was internalized into caveolin-1-positive endosomes. Although TNF-induced MLCK activation and MLC phosphorylation were increased, caveolin-1 knockout mice failed to internalize occludin and were protected from TNF-induced barrier loss [76]. Caveolar endocytosis is, therefore, required for TNF-induced, MLCK-mediated leak pathway permeability increases. Occludin was the only tight junction protein consistently identified within these endosomes, but the data do not exclude the possibility that occludin is merely a marker of some other critical change in tight junction structure. To assess this, transgenic mice that overexpress occludin within intestinal epithelial cells were treated with TNF. Although endocytosis did occur, tight junction-associated occludin pools were preserved in transgenic, relative to wild-type, mice [76]. Moreover, the magnitude of barrier loss was attenuated, and TNF-induced diarrhea was prevented by transgenic occludin expression [76]. Occludin is, therefore, an essential regulator of leak pathway permeability. Finally, this result is consistent with leak pathway barrier loss and reduced occludin expression in human inflammatory bowel disease [36, 77, 78].

Occludin Regulates Epithelial Survival

The initial reports of occludin knockout mice failed to identify intestinal or renal barrier defects or spontaneous disease involving these organs [79]. Nevertheless, occludin knockout was associated with male infertility, inability of females to effectively suckle pups, gastric epithelial hyperplasia, brain calcifications, testicular atrophy, and osteoporosis, and subsequent studies showed that occludin knockout mice became deaf over time [80]. Finally, our unpublished experience that the frequency of knockout pups when occludin heterozygotes were bred was far less than the expected 25% predicted by Mendelian genetics suggests embryonic loss; crosses of occludin floxed and occludin floxed; villin-cre transgenic mice resulted in equal numbers of cre-positive and cre-negative pups. Embryonic loss of occludin-deficient embryos is, therefore, not due to *Ocln* deletion within the intestinal epithelium.

The absence of intestinal disease in occludin-deficient mice could be due to compensatory mechanisms that overcome occludin loss. Universal and intestinal epithelial-specific occludin knockout mice were, therefore, stressed in order to unmask functional deficits. Remarkably, dextran sulfate sodium (DSS) failed to induce weight loss in occludin knockout mice (Fig. 6a) [81]. Thus, intestinal epithelial occludin expression sensitizes mice to chemically induced colitis.

Further analyses showed that histopathology, inflammatory cytokine production, and epithelial apoptosis induced by DSS or trinitrobenzenesulfonic acid (TNBS) were suppressed in intestinal epithelial occludin knockout mice (Fig. 6b). To better define the mechanism by which occludin loss reduced epithelial apoptosis, mice were treated with 5-fluorouracil, which triggers intrinsic pathway apoptosis, systemic T-cell activation using anti-CD3, or TNF treatment to activate the extrinsic apoptotic pathway [81]. Occludin knockout protected intestinal epithelial cells from all three stimuli (Fig. 6c). Biochemical analysis of apoptotic signaling induced by TNF showed that ERK, p38 MAPK, IkB, and caspase-9 were all activated similarly in occludin-sufficient and knockout intestinal epithelia (Fig. 6d). However, TNFinduced caspase-3 cleavage was only detected in occludin-expressing epithelia (Fig. 6d). Further characterization showed that this was due to a ~ 50% reduction in overall caspase-3 mRNA and protein expression in the absence of occludin. Although it was surprising that an only 50% decrease in expression would be sufficient to prevent apoptosis, mice heterozygous for Casp3 knockout (Casp3^{+/-}), which express caspase-3 at ~50% of normal levels, were also protected from intestinal epithelial apoptosis induced by DSS or TNF [81]. Thus, occludin promotes



Fig. 6 Occludin deletion reduces caspase-3 expression, limits epithelial apoptosis, and attenuates experimental disease in mice; both occludin and caspase-3 expression are reduced in Crohn's disease. (a) DSS-induced colitis severity and weight loss were reduced in $Ocln^{-/-}$ (red circles) relative to wild-type (WT, blue squares) mice. (b) Numbers of apoptotic (ISOL-positive, red) epithelial cells were increased in DSS-treated wild-type, but not intestinal epithelial-specific occludin knock

caspase-3 transcription, and the reduced caspase-3 expression that follows occludin loss is sufficient to prevent intrinsic and extrinsic pathway apoptosis.

Similar to mice, Caco-2 cells lacking occludin displayed reduced *CASP3* transcription and protein expression. Inducible occludin expression in these cells activated the *CASP3* promoter and restored transcription to normal levels [81]. Thus, although occludin has not been detected within nuclei, it promotes *CASP3* transcription, either directly or indirectly [81]. To assess the effect of inflammation-induced loss of occludin expression on epithelial survival, Caco-2 monolayers were treated with TNF at a low concentration that was insufficient to cause apoptosis. This was, however, sufficient to reduce both occludin and caspase-3 expression and protect cells from staurosporine-induced intrinsic pathway or high-dose TNF-induced extrinsic pathway apoptosis. In contrast, pretreatment of occludin-deficient Caco-2 monolayers with low-dose TNF had no effect on subsequent responses to staurosporine or high-dose TNF. Thus, low-grade inflammation, modeled in vitro by low-dose TNF, may lead to reduced occludin and caspase-3 expression that results in cytoprotection [81].

In order to determine whether occludin downregulation in human disease is associated with reduced caspase-3 expression, small intestinal and colonic biopsies from Crohn's disease and ulcerative colitis patients and age- and sex-matched healthy control subjects were analyzed by quantitative immunohistochemistry (Fig. 6e). Occludin expression was significantly reduced in both Crohn's disease and ulcerative colitis and was accompanied by caspase-3 downregulation (Fig. 6f). Moreover, reductions in caspase-3 expression correlated directly with decreased occludin expression. Thus, occludin downregulation in inflammatory disease may be an adaptive response that promotes epithelial survival. Further studies will be needed to understand how this cytoprotective function interfaces with the role of occludin in tight junction barrier regulation.

Fig. 6 (continued) out (KO^{IEC}), mice. Nuclei (blue) are shown for reference. Bar, 20 μ m. (c) Systemic T-cell activation induced by anti-CD3 treatment increased ISOL-positive (red) cell numbers in wild-type, but not KO^{IEC}, mice. Nuclei (blue) are shown for reference. Bar, 50 μ m. (d) Western blot of intestinal epithelia isolated from vehicle- and TNF-treated WT and *Ocln^{-/-}* mice. With the exception of cleaved caspase-3, signaling events downstream of TNF signaling were not affected by occludin deletion. (e) Ileal biopsies from healthy subjects and Crohn's disease patients show reduced occludin (green) and caspase-3 (green) expression in disease. E-cadherin (red) and nuclei (blue) are shown for reference. Bar, 50 μ m. (f) Quantitative morphometry confirms similar reductions in occludin and caspase-3 expression intensity in ileal epithelium from Crohn's disease patients (green diamonds) relative to healthy controls (yellow circles) (Data from Kuo et al. Gastroenterology, 2019, with permission)

Summary and Future Directions

The studies discussed here demonstrate the important contributions of pore and leak pathway permeability to intestinal disease. They also highlight the importance of considering these mechanisms as separate processes that are regulated by different signaling events and have distinct effects on barrier function and pathophysiology. The data also show that the impact of increased tight junction permeability may not be the same in infectious and immune-mediated disorders. Although available data cannot explain the reasons for these differences, one might hypothesize that increased claudin-2 expression in infectious disease both enhanced water efflux and augmented mucosal immune activation. Evolutionarily, where infectious diseases had much greater impact on survival than immune-mediated disorders, increased pathogen clearance and immune activation might be considered synergistic. The contradiction only occurs in the context of immune-mediated disease. Although the effects of occludin downregulation in infectious disease have not been defined, one could hypothesize that increased leak pathway permeability might promote pathogen clearance while simultaneously downregulating caspase-3 expression to limit tissue damage. Thus, regulatory mechanisms that evolved over billions of years to promote survival may now have unintended consequences in the context of modern maladies.

The growing synergy between in vitro and in vivo models makes this an exciting time for tight junction biology in which future studies will address the questions raised in this text and many others. Ultimately, we hope that both fundamental, foundational advances and translational studies will lead to clinical trials in which barrier and tight junction modulation are validated as therapeutic interventions for a broad range of intestinal and systemic diseases.

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Conflict of Interest Statement JRT is a founder and shareholder of Thelium Therapeutics and has served as a consultant for Entrinsic, Immunic, and Kallyope.

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