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Yan Y. Sweat, Jerrold R. Turner

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## Commentary

It is clear that excessive mucosal immune activation and intestinal barrier dysfunction both contribute to inflammatory bowel disease (IBD) pathogenesis. T cell protein tyrosine phosphatase (TCPTP), which extinguishes signaling in immune cells, is linked to IBD and other immune-mediated diseases. In this issue of the *JCI*, Marchelletta and Krishnan et al. demonstrate that, in intestinal epithelial cells, TCPTP regulates tight junction permeability in vivo. Intestinal epithelial TCPTP loss potentiated cytokine-induced barrier loss, and this synergized with effects of TCPTP loss in immune cells. This work implicates a single mutation as the cause of distinct functional aberrations in diverse cell types and demonstrates how one genetic defect can drive multihit disease pathogenesis.



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## *PTPN2* mutations cause epithelium-intrinsic barrier loss that synergizes with mucosal immune hyperactivation

#### Yan Y. Sweat and Jerrold R. Turner

Laboratory of Mucosal Barrier Pathobiology, Department of Pathology, Brigham and Women's Hospital and Harvard Medical School, Boston, Massachusetts, USA.

It is clear that excessive mucosal immune activation and intestinal barrier dysfunction both contribute to inflammatory bowel disease (IBD) pathogenesis. T cell protein tyrosine phosphatase (TCPTP), which extinguishes signaling in immune cells, is linked to IBD and other immunemediated diseases. In this issue of the *JCI*, Marchelletta and Krishnan et al. demonstrate that, in intestinal epithelial cells, TCPTP regulates tight junction permeability in vivo. Intestinal epithelial TCPTP loss potentiated cytokine-induced barrier loss, and this synergized with effects of TCPTP loss in immune cells. This work implicates a single mutation as the cause of distinct functional aberrations in diverse cell types and demonstrates how one genetic defect can drive multihit disease pathogenesis.

## TCPTP expression may regulate immune cell activation

Inflammatory bowel disease (IBD) has been associated with over 240 genetic polymorphisms. In most cases, these gene variants are insufficient to cause disease, and their relative contributions to pathogenesis are difficult to assess. Large genomewide association studies have linked many SNPs and associated genes to multiple immune-mediated diseases. The protein tyrosine phosphatase non-receptor type 2 (PTPN2) gene has been linked to Crohn's disease, ulcerative colitis, type I diabetes, and rheumatoid arthritis (1). In adults, PTPN2 polymorphisms increase the risk of IBD only modestly (odds ratio, approximately 1.35), but a recent report implicating a PTPN2 mutation in a patient with very early onset IBD suggests that the effects may be much greater in some contexts (2).

PTPN2 encodes T cell protein tyrosine phosphatase (TCPTP), which dampens

JAK/STAT pathway activation (3). Because JAK/STAT pathways intersect with many other signaling events, TCPTP regulates diverse processes in many cell types. Consequences of TCPTP loss include unrestrained T cell receptor signaling, exaggerated cytokine responses, increased MAPK signaling, excessive growth factor receptor activation, and defective autophagosome formation (4, 5). The observation that TCPTP expression is upregulated in response to IFN-y but also attenuates IFN- $\gamma$  signaling (3, 6) suggests that increased TCPTP expression in inflammatory diseases, including IBD, may be a critical mechanism of regulating immune cell activation. Consistently, TCPTP deletion in T cells causes severe, systemic autoimmune disease in mice (5).

## TCPTP in intestinal barrier regulation

In this issue of the *JCI*, Marchelletta and Krishnan et al. focus on the role of TCPTP

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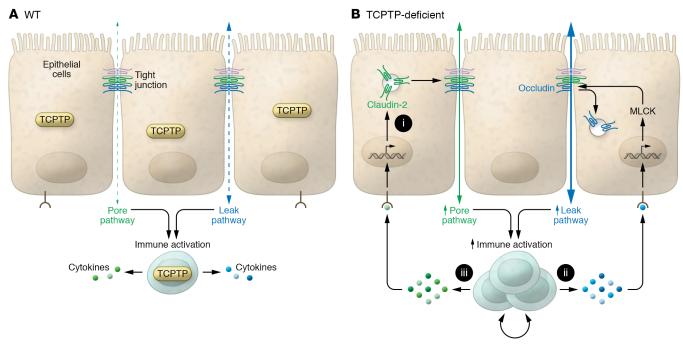
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in intestinal barrier regulation (7). Diseaseassociated human PTPN2 polymorphisms result in loss of TCPTP function, making Ptpn2 WT, knockout, and heterozygous mice a useful system. Both heterozygous and knockout mice showed markedly increased permeability to 4 kDa dextran, relative to WT mice. In contrast, the barrier to 70 kDa dextran was maintained (7). These results implicate increased flux across the low-capacity paracellular leak pathway, accommodating macromolecules up to approximately 125 Å in diameter, and exclude epithelial damage as the source of barrier loss (8). The leak pathway is regulated by myosin light chain kinase, which phosphorylates myosin II regulatory light chain to trigger endocytosis of the tight junction protein occludin (9, 10). Consistent with this process, Marchelletta and Krishnan et al. found that myosin II regulatory light chain phosphorylation was increased and occludin was internalized in epithelial cells of Ptpn2-knockout mice (also referred to as Tcptp-deficient mice). Although not explored here, it is likely that myosin light chain kinase expression was also increased (11, 12).

Marchelletta and Krishnan et al. also demonstrated increased expression of the pore-forming tight junction protein claudin-2 in Ptpn2-knockout mice. Claudin-2 may contribute to the increased transmucosal conductance, i.e., reduced resistance, measured in the distal ileum and cecum of Ptpn2-knockout mice, but cannot explain increased 4 kDa dextran flux, as claudin-2 channels exclude molecules larger than water and monovalent cations. The observed claudin-2 upregulation may be secondary to increased mucosal IL-13 and IL-22, each of which promotes claudin-2 transcription, were increased within intestinal mucosae of Ptpn2-knockout mice. In addition to these cytokine-mediated effects, Marchelletta and Krishnan et al. considered the possibility that TCPTP



**Figure 1. A cycle of barrier loss and mucosal immune hyperactivation. (A)** WT cells that express TCPTP are able to effectively dampen immune responses to limit changes in flux across the pore pathway (hatched green arrow), permeable to sodium and water, and the leak pathway (hatched blue arrow), permeable to larger molecules, including proteins and polysaccharides. **(B)** TCPTP loss in intestinal epithelial cells leads to epithelium-intrinsic signaling that upregulates claudin-2 expression (i). The resulting claudin-2-related increases in pore pathway (solid green arrow) permeability trigger excessive cytokine secretion in TCPTP-deficient immune cells. This cytokine release activates myosin light chain kinase (MLCK) to cause occludin internalization and leak pathway (solid blue arrow) permeability increases (ii). The exaggerated immune responses may also cause further increases in claudin-2 expression (iii). These cytokine-induced changes in permeabilities of and flux across pore and leak pathways trigger further immune activation, thereby creating a vicious cycle that leads to disease. In this manner, dysfunction of a single protein, TCPTP, results in excessive signaling in both epithelial and immune cells in development of a dysregulated cycle that leads to disease.

loss in epithelial cells could directly amplify STAT signaling and claudin-2 expression. To pursue this possibility, the authors developed intestinal epithelial-specific *Ptpn2*-knockout mice (also called TCPTP-deficient mice). In the absence of exogenous insults, intestinal epithelial STAT signaling and claudin-2 expression were upregulated in these mice, supporting the conclusion that epithelial TCPTP loss contributes to disease (7).

The complex in vivo milieu makes it difficult to definitively demonstrate epithelium-intrinsic effects of TCPTP mutation. Marchelletta and Krishnan et al. turned to a stably expressed dominant-negative TCPTP<sup>C1265</sup> mutant in human intestinal epithelial cells. This mutant was sufficient to increase both claudin-2 expression and pore pathway permeability. siRNA blockade of claudin-2 expression partially corrected this transepithelial electrical resistance (TER) loss (7). TCPTP dysfunction is therefore sufficient to activate claudin-2 expression and increase permeability of the high-capacity, charge- and size-selective pore pathway via an epithelium-intrinsic process (8).

Analyses of tissues from intestinal epithelial-specific Ptpn2-knockout mice confirmed increased claudin-2 expression and reduced TER, but 4 kDa dextran permeability was unchanged. However, in vitro studies showed that both STAT1 activation and 4 kDa dextran permeability increases induced by IFN-y were potentiated in TCPTP<sup>C1268</sup>-expressing epithelial cells, consistent with the role of TCPTP in extinguishing cytokine signaling (7). Hyperactivation of STAT and MAPK pathways in TCPTP-deficient epithelial cells is therefore likely to explain increased pore and leak pathway permeabilities by promoting transcription of claudin-2 and myosin light chain kinase, respectively (13, 14).

### A vicious cycle that establishes disease

Why is permeability of the leak pathway increased in universal, but not intestinal epithelial-specific, *Ptpn2*-knockout mice? One possibility is that epithelial TCPTP

dysfunction leads to increased claudin-2 expression and pore pathway permeability that, secondarily, triggers mucosal immune cell activation (Figure 1). In TCPTPdeficient immune cells, cytokine release is excessive (15, 16). Cytokine signaling is exaggerated in TCPTP-deficient epithelial cells and leads to excessive leak pathway permeability increases that ultimately feed back onto a hyperactive mucosal immune system, cause further cytokine signaling and barrier loss, and create a vicious cycle that establishes disease. This model is supported by the enhanced mucosal immune activation observed in mice with intestinal epithelial claudin-2 overexpression and increased pore pathway permeability as well as the modest, microbiota-driven mucosal immune activation that occurs in mice with intestinal epithelial myosin light chain kinase hyperactivation and increased leak pathway permeability (16-18).

Neither pore nor leak pathway permeability increases alone are sufficient to induce disease (15–18). We can therefore infer that disease in universal *Ptpn2*-knock-

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out mice and humans with germline PTPN2 mutations requires events beyond those caused by epithelial TCPTP dysfunction. As described in the proposed model (Figure 1), these events likely include defective immunoregulation. Marchelletta and Krishnan et al. therefore provide insight into how modification of a single gene can trigger events in separate, but functionally synergistic, tissue types to generate phenotypes that could not be predicted by examining either tissue in isolation (7). This concept can be expanded to include synergistic effects of multiple mutations on the same or separate cell types, consistent with the polygenic nature of most immune-mediated disease.

#### Questions and insights

The work by Marchelletta and Krishnan et al. (7) is a substantial advance, but, as with any good study, the results also raise new questions. For example, if the deletion of *Ptpn2* acts primarily through the hyperactivation of the JAK/STAT signaling pathways, do JAK/STAT inhibitors reverse this phenotype? Do pore and leak pathway permeability increases make distinct or overlapping contributions to immune activation? This work also underscores the perennial question that arises when considering immune activation and mucosal barrier loss, which is the chicken and which is the egg?

Overall, Marchelletta and Krishnan et al. have deftly separated pore and leak pathways and discovered that, via epithelium-intrinsic processes, claudin-2mediated pore pathway permeability is increased when epithelial TCPTP function is perturbed in vivo (7). Leak pathway permeability increases require extrinsic stimuli, e.g., cytokines, whose effects are augmented in the absence of intact epithelial TCPTP. These findings shed light on the multiple mechanisms by which *PTPN2* polymorphisms contribute to disease pathogenesis (7). The data may also provide insight into some of the relationships between intestinal barrier loss and IBD risk (19, 20).

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Address correspondence to: Jerrold R. Turner, Brigham and Women's Hospital, Harvard Medical School, 77 Avenue Louis Pasteur, HNRB 730B, Boston, Massachusetts 02115, USA. Phone: 617.525.8165; Email: jrturner@bwh.harvard.edu.

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