

Guardians of Intestinal Homeostasis: Focus on Intestinal Epithelial Cells

Maha M. Elbrashy^{1,2}, Hozaiifa Metwally^{1,*}, Tadimitsu Kishimoto^{1,*}

¹Laboratory of Immune Regulation, The World Premier International Research Center Initiative (WPI) Immunology Frontier Research Center, Osaka University, 565-0871 Osaka, Japan

²Biochemistry Department, Biotechnology Research Institute, National Research Center, P.O. 12622, Giza, Egypt

*Correspondence should be addressed to Hozaiifa Metwally, hozaiifa1@ifrec.osaka-u.ac.jp; Tadimitsu Kishimoto, kishimoto@ifrec.osaka-u.ac.jp

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Abstract

The intestinal epithelium not only facilitates the absorption of nutrients, but also plays a pivotal role in guarding intestinal homeostasis and preventing opportunistic gut microbiome invasions. The intestinal epithelial cells have diverse and coordinated regulatory networks that provide intricate lines of defense, in order to maintain the integrity of the intestinal barrier. The epithelial defense comprises the anatomical structure of intestinal epithelial cells as a stout physical barrier and its covering mucus layer containing diverse antimicrobial peptides. In addition, epithelial cells are well-equipped with diverse microbial sensors, which upon activation induce the expression of downstream immunomodulatory networks of cytokines and chemokines. In this review, we summarize the intestinal epithelial cells' defense mechanisms and their role in maintaining intestinal homeostasis.

Keywords: Cytokine biology, Epithelial cells, Immunomodulation, Inflammation

Introduction

The intestinal epithelial cells (IECs) form a semi-permeable barrier that allows the absorption of nutrients and electrolytes, meanwhile preventing harmful external environmental antigens from entering the host's internal environment, in order to maintain the host's homeostasis [1]. The mammalian intestine accommodates a dynamic community of trillions of microorganisms allowing the adaptation of diverse saccharolytic enzymes that complement the limited saccharolytic diversity encoded in the mammalian genome [2]. Although the host-microorganisms relationship is symbiotic in nature, such a dense bacterial community poses a serious threat to the host, and opportunistic invasion of the host's internal environment by gut resident bacteria can lead to serious pathologies such as chronic inflammation or,

in extreme cases, bacteremia. Taking into consideration the huge numbers and the high diversity of gut microbiome, and the large surface area of the intestine, the best defense strategy for the host is to prevent microorganisms from breaching the epithelial barrier. IECs play a central role in maintaining the epithelial barrier through shaping physical and chemical layers of defense to prevent the invasion of gut microorganisms into the internal environment of the host [3,4]. However, gut bacteria have evolved evasion strategies to escape from these layers of defense and breach epithelial barrier. Remarkably, epithelial cells utilize a next layer of defense against the invaded pathogens through cell-intrinsic microbial sensors and the activation of immune responses to eradicate invading pathogens [5,6]. In this review, we discuss the adaptations of IECs to limit opportunistic invasions of resident microorganisms, and how the intestinal epithelium

minimizes the contact between luminal microbiota and the immune system, in order to prevent destructive immune responses and maintain intestinal homeostasis.

First Line of Defense: The Mucus Layer

Goblet cells, which are specialized epithelial cells, secrete mucin glycoproteins. These mucins are arranged into a viscous gel-like layer covering the epithelial surface, forming inner and outer mucus layers [7]. Whereas the small intestine has only one loose mucus layer, the large intestine displays the two mucus layers to defend against trillions of inhabiting gut bacteria [8]. Mice deficient in Muc2, a key mucin glycoprotein, show bacterial translocation across the mucosal barrier and develop spontaneous colitis [9]. In addition, probiotic microbiota, such as lactobacilli, enhance the intestinal epithelium barrier function through the stimulation of mucin production [10]. On the other hand, colitogenic microbiota, such as *Entamoeba histolytica*, precipitate intestinal inflammation by degrading the C-terminal region of mucin [11].

Second Line of Defense: The Apical Junctional Complex (APC)

Epithelial cells exhibit an anatomical structure that separates the internal host environment from the external environmental stresses [12]. This separation is partly achieved by apical junctional complex (APC), which ensures the impermeability of both commensal and pathogenic bacteria inhabiting the gut. The APC consists of three types of junctional proteins. First, tight junction proteins such as claudins, occludin, junctional adhesion molecules, and zonula occludens (ZO). Second, adherens junction proteins such as E-cadherin. Lastly, the desmosomes [13]. Mice deficient in occludin expressed morphologically intact tight junction structures but exhibited elevated inflammation and a defective gut barrier [14]. Bacteria such as *Clostridium difficile* and *Listeria monocytogenes* target occludin and claudin, respectively, thus weakening the intestinal epithelial barrier and promoting their invasion leading to an increase in intestine permeability [15]. Moreover, epithelial cells display a constant turnover cycle every 3-5 and 5-7 days in the small and large intestines, respectively, which renews the epithelial barrier, and thus maintains intestinal homeostasis [16,17].

Third Line of Defense: Antimicrobial Peptides (AMPs) and Immunoglobulin A (IgA)

AMPs are secreted by gut epithelial cells and diffused through the mucus layer to prevent unwanted colonization of microbes. AMPs include defensins, cathelicidins, and C-type lectins, which have a wide spectrum of antibiotic activity [18,19]. Besides the microbicidal activity of AMPs, they have several other actions, like stimulating mucus secretion [20], expression of tight junction proteins [21], chemotaxis, cell

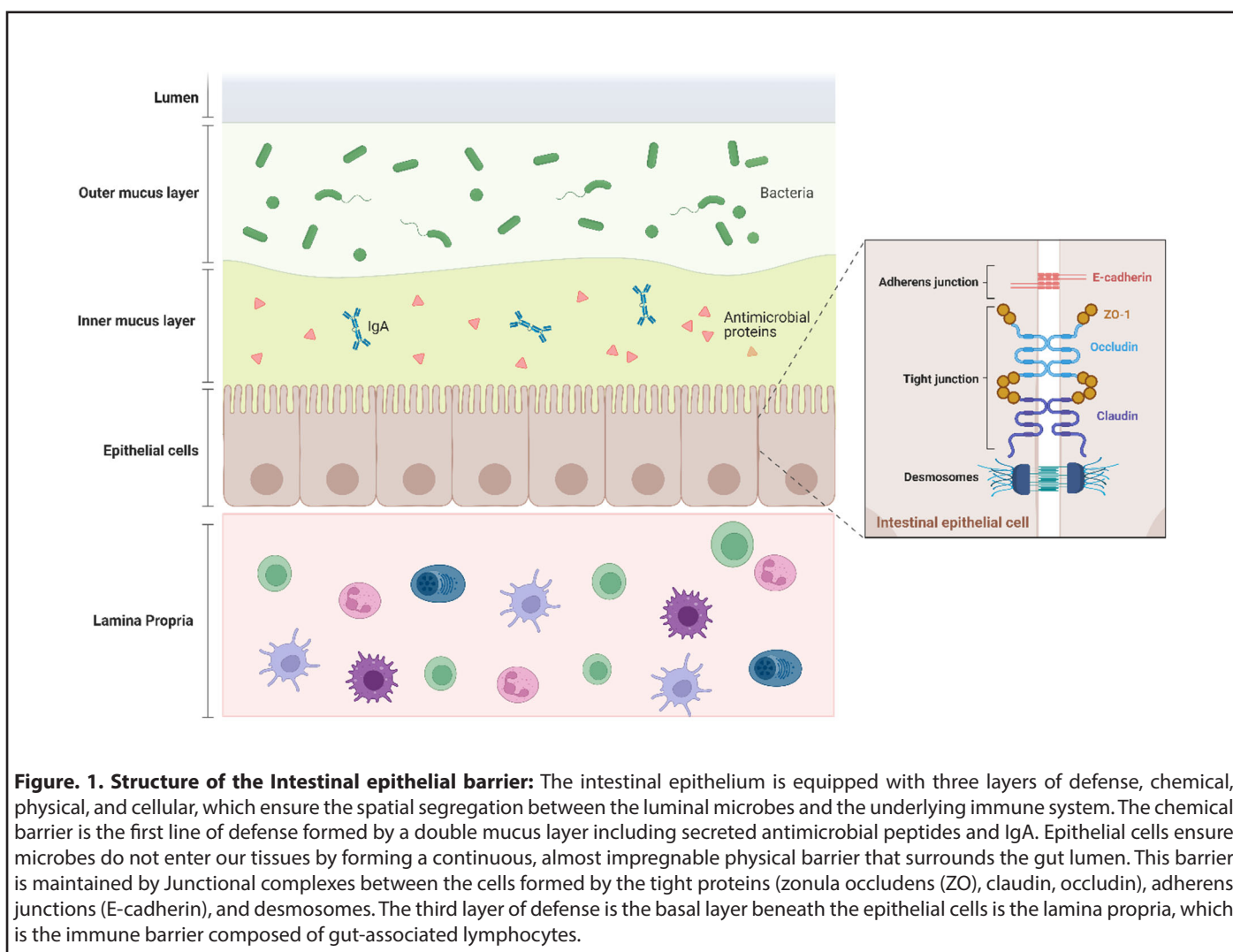
proliferation, and enhancing the production of extracellular matrix proteins, confirming their roles in wound healing [22]. Defects in endogenous AMP expression and function have been linked with intestinal inflammation in mice. For example, Mice with a deficiency in Regenerating islet-derived protein 3 gamma (Reg IIIγ), a C-type lectin, exhibited increased mucosal bacterial burden and impaired spatial relationships between bacteria and their host tissues [23]. In addition, Ly6/Plaur domain-containing 8 (Lypd8) has been recently identified as an antimicrobial peptide, which contributes to the segregation of intestinal bacteria and intestinal epithelia in the large intestine [24]. Mice lacking Lypd8 demonstrate the disappearance of the bacteria-free space just above the epithelial layer of the colon [25].

IgA is secreted by plasma cells located in lamina propria. The secreted form of IgA (sIgA) is transcytosed across the epithelium to allow binding to luminal bacteria. The transcytosed IgA binds to bacteria on the luminal side of the epithelial barrier and prohibits their translocation [26]. The exact mechanisms by which IgA does these roles remain elusive but may include the ensnaring of bacteria in the mucus layer or enhancing fast phagocytic clearance of the pathogens that invade the epithelial cell barrier [27].

Fourth Line of Defense: Network of Microbial Sensors and Immunomodulatory Effectors

Despite the aforementioned defenses of IEC, pathogenic microorganisms have evolved escape strategies to escape. Remarkably, epithelial cells are well-equipped with intrinsic diverse and sophisticated regulatory networks to defend against the invaded pathogens. IECs utilize complex microbial sensors called pattern recognition receptors (PRRs) [28], which upon activation orchestrate the secretion of immunomodulatory molecules such as chemokines and cytokines to culminate in an appropriate immune response towards invading pathogens [29,30]

PRRs recognize conserved bacterial structures called pathogen-associated molecular patterns (PAMPs) that are not found in the host's cells. Additionally, PRR recognizes danger-associated molecular patterns (DAMPs) released from stressed or damaged cells [31]. PRRs comprise two main categories of receptors: membrane-bound toll-like receptors (TLRs), and intracellular nucleotide oligomerization domain (NOD)-like receptors (NLRs). Upon microbial recognition, activation of PRRs activates a signaling cascade culminating in the expression of pro-inflammatory cytokines and antimicrobial mediators, and the recruitment of immune cells to aid in the eradication of the bacterial threat and protect the epithelium from pathogenic invasion [32]. Further, it has been shown that TLR signaling is involved in epithelial cell proliferation [33,34], IgA production [35], maintenance of tight junctions [36], and antimicrobial peptide expression which are essential for



maintaining a healthy epithelial barrier [37]. TLR2 activation efficiently maintains the tight junction-associated barrier assembly in intestinal epithelial cells against stress-induced damage and inhibits mucosal inflammation [38]. High expression levels of TLR3 in intestinal epithelial cells correlate with resistance against rotavirus infection [39]. Mice that have deficiencies either in Tlr4 or Tlr5 exhibit impaired innate immune responses and are more vulnerable to dextran sodium sulfate (DSS)-induced colitis [40,41]. The best-characterized NLRs are NOD1 and NOD2, both of which identify PAMPs by leucine-rich repeats (LRRs) at the C-terminus like TLRs [42]. In accordance with their intracellular localization, NODs are specific for the detection of pathogens that invade the intestinal epithelial cells such as *Shigella* and *Salmonella* [43]. Consistently, Nod2-deficient mice exhibited severe colitis with increased bacterial invasion [44].

IECs present luminal antigens to intraepithelial lymphocytes (IELs) to modulate the adaptive immune system [45]. The bidirectional interactions between IELs and IECs are important to maintain immune homeostasis at the intestinal barrier [46].

Following bacterial invasion, IECs secrete chemokines such as CXCL8, CXCL1, CXCL3, and CXCL5 that recruit neutrophils. In turn, neutrophils secrete interleukin-22 (IL-22), which stimulates the expression of antimicrobial peptides by the colon epithelium and protects the epithelium from chemically induced damage [47,48]. Besides, IECs recruit dendritic cells (DCs) and T helper 17 (Th17) cells by secreting CCL20. In turn, Th17 secretes IL-17 which increases intestinal epithelial cell proliferation and reduces barrier permeability [49]. In addition, DCs cells secrete IL-28A, which induces intestinal epithelial proliferation [50].

Conclusion and Future Remarks

IECs are key players in shaping and maintaining intestinal homeostasis, which is critical for the host's metabolism and survival. IECs utilize diverse and complex networks to shape physical and chemical layers of defense against opportunistic gut microbiome invasions. Additionally, the cross-talk between IEC and the immune compartment strengthens the host defense against possible pathogenic invasions from gut

microbiome. It is thus of particular interest to understand how the IECs regulatory signaling networks operate, and how these sophisticated pathways shape cell-intrinsic and cell-cell responses. Harnessing such knowledge will eventually offer more opportunities for better and more specific therapies targeting IECs to strengthen the intestinal barrier and/or induce immune tolerance with minimal side effects. Besides, nutritional support to enrich certain microbiome and promote a gut anti-inflammatory environment may prevent provoking detrimental immune response. Also, stem cell therapies to regenerate damaged IECs may be potentially helpful for treating severe cases of intestinal inflammation. Of note, choosing the best therapeutic strategy will likely depend on the comprehensive assessment of the etiology and the severity of intestinal inflammation.

Author Contributions

M.M.E wrote the manuscript draft. H.M. and T.K edited the manuscript and supervised the work.

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