

The Apical Junctional Complex in Respiratory Diseases

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The epithelium, including the respiratory system, acts as a selective gate between the outside environment and underlying tissue. Epithelial cells are polarized due to the formation of the apical junctional complex, which includes adherent junctions and tight junctions. Endothelial cells are one of the most important cellular constituents of blood vessels. Endothelial junctional proteins play important roles in tissue integrity as well as in vascular permeability, leukocyte extravasation, and angiogenesis. This review focuses on the apical junctional complex in respiratory diseases.

Key Words: *Epithelium; Permeability; Tight junctions*

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INTRODUCTION

The epithelium, including the respiratory system and other organs, acts as a selective gate between the outside environment and underlying tissue.¹ Epithelial cells are polarized by the formation of specialized cell-cell junctions, which are referred to as the apical junctional complex (AJC) (Fig. 1). The AJC consists of adherent junctions (AJs) and tight junctions (TJs).¹

Endothelial cells are one of the important cellular con-

stituents of blood vessels. Endothelial junctional proteins play important roles in tissue integrity as well as in vascular permeability, leukocyte extravasation, and angiogenesis. Inter-endothelial junctions play pivotal roles in tissue integrity, barrier function, and cell-cell communication.² Cell-cell junctions are the sites of attachment between endothelial cells but also function as signaling structures that communicate cell position, limit growth and apoptosis, and regulate vascular homeostasis.²

Some proteins³ are particular to endothelial cells [e.g., VE-cadherin and claudin (CLDN)-5], whereas others are common to epithelial cells [e.g., occludin, junctional adhesion molecule (JAM)-A, nectins, claudins, and connexins], blood cells (e.g., PECAM/CD31, endothelial cell selective adhesion molecule, JAM-A, JAM-C, CD99), smooth muscle cells (e.g., S-endo-1/CD146), and mesangial/trophoblast cells (e.g., protocadherin12/VE-cadherin-2). These proteins may be components of organized junctional structures, such as VE-cadherin in AJs, claudins and occludins in TJs, or connexins in gap junctions. However, others are independent, such as PECAM, CD99, and S-endo-1.^{3,4}

Junctional complexes start off intracellular signaling in various ways. Signaling can be directly activated by engaging signaling proteins or growth-factor receptors or can be indirectly activated by tethering and retaining transcription factors at the cell membrane, thereby limiting their nuclear translocation.⁵⁻⁸

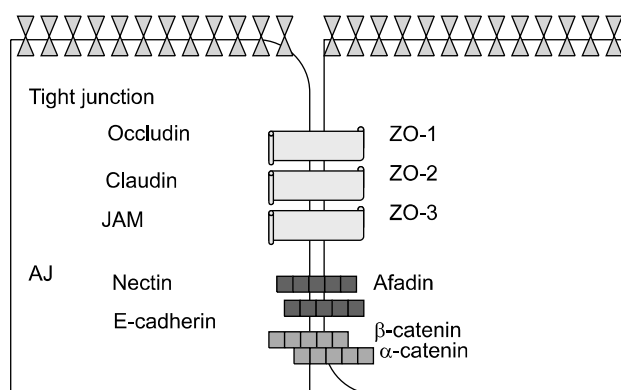


FIG. 1. Apical junctional complexes. The AJC includes adherent junctions (AJs) and tight junctions (TJs).

ADHERENT JUNCTIONS

The important AJ transmembrane proteins belong to the cadherin family.⁹ E-cadherin is expressed primarily in epithelium. The cytoplasmic tail of E-cadherin is linked to the actin cytoskeleton and other signaling elements through many peripheral membrane proteins, including catenins, vinculin, and α -actinin.¹⁰ Through their cytoplasmic tail, junctional adhesion proteins bind to cytoskeletal and signaling proteins, which permits anchoring of the adhesion proteins to actin microfilaments and the transfer of intracellular signals inside the cell.¹⁰ The association with actin is required for stabilization of the junctions as well as for dynamic regulation of junction opening and closure. In addition, the interaction of junctional adhesion proteins with the actin cytoskeleton might be relevant for maintaining cell shape and polarity.¹¹ In addition to acting as adaptors that mediate the binding of adhesion proteins to actin, some intracellular junctional proteins, upon release from junctions, translocate to the nucleus and modulate transcription.¹² Another characteristic of some junctional proteins is their function as scaffolds and binding of several effector proteins, which makes their reciprocal interaction easy. A typical example is the TJ component zona occludens-1 (ZO1), which can associate with many transmembrane proteins, including claudins, occludin, or JAMs, as well as with cytoskeletal binding proteins, such as cortactin, cingulin, α -CATENIN, and indirectly with vinculin and α -actinin.^{13,14} ZO1 also associates with other PDZ-DOMAIN-containing proteins such as ZO2 and with signaling mediators such as ZO1-associated nucleic-acid binding (ZONAB).^{13,14} VE-cadherin is the transmembrane component of endothelial AJs.¹⁵ The VE-cadherin complex also transiently or permanently associates with signaling partners, as well as with a specific apical-basal polarity complex, through direct interaction with PAR-3 and PAR-6.¹⁵ VE-cadherin is located at the junctions of all endothelial subtypes. β -Catenin is often noticed in the nucleus and cytoplasm of endothelial cells during pathological angiogenesis or vascular remodeling.¹⁵ Nectins are Ca^{2+} -independent Ig-like CAMs.¹⁵ Nectin-based cell-cell adhesions set up and maintain AJCs, both independently and by working together with the cadherin-based AJs.¹⁵ Afadin binds to both nectins and α -catenin and recruits the cadherin- β -catenin complex to the nectin-based cell-cell adhesion site to form AJs.^{14,15}

TIGHT JUNCTIONS

TJs are the apical-most constituents of the AJC in vertebrate epithelial cell sheets.¹ They are also observed in vascular endothelial cells and mesothelial cells. TJs function as a semipermeable gate for the paracellular transport of ions, solutes, water, and cells and are considered to function as a fence that divides the apical and basolateral domains of plasma membranes.¹ TJs coordinate a variety of signaling and trafficking molecules that regulate cell dif-

ferentiation, proliferation, and polarity.¹ These functions enable epithelial and endothelial cell sheets to establish distinct tissue compartments within the body and to maintain homeostasis.^{16,17} Moreover, disturbance of TJ function likely causes or contributes to a variety of pathological conditions, such as inflammatory bowel disease, infection, cancer, vasogenic edema, and blood-borne metastasis.^{16,17} The barrier function of the TJ also restricts drug delivery to underlying tissues.^{16,17} Therefore, determining how to overcome paracellular barriers, such as the blood-brain barrier, is critical for the treatment of human disease.^{16,17} Occludin is an approximately 60-kDa tetraspan membrane protein with two extracellular loops, a short intracellular turn, and N- and C-terminal cytoplasmic domains.¹⁸ Claudins¹⁹ are 18- to 27-kDa tetraspan proteins with short cytoplasmic N-termini, two extracellular loops, and a C-terminal cytoplasmic domain. They do not show any sequence similarity to occludin. Claudins are capable of forming TJ strands and thus are the backbone of the TJ.¹⁹ The claudin family consists of 24 members in mice and humans and exhibits distinct tissue- and cell-type-specific expression patterns.¹⁹ Many claudin species are generally expressed in most epithelial cell types. Cldn2, Cldn3, Cldn4, Cldn7, Cldn8, Cldn12, and Cldn15 are abundantly expressed in the duodenum, jejunum, ileum, and colon.¹⁹ The segment- or axis-specific expression of different claudins has also been reported in several other epithelial tissues, including the liver lobule, nephron, and inner ear.

JAMs contain two extracellular Ig-like domains, a single transmembrane region, and a C-terminal cytoplasmic domain.²⁰ The JAM family is divided into two subgroups based on their sequence similarities.²⁰ The first subgroup, JAM-A, JAM-B, and JAM-C, has a class II PDZ domain-binding motif at the C-terminal end and directly interacts with ZO-1 and PAR-3.²⁰ In contrast, members of the second subgroup, coxsackie and adenovirus receptor, endothelial cell-selective adhesion molecule (ESAM), and JAM4, contain a class I PDZ domain-binding motif at their C-terminus.²⁰ CAR and JAM4 associate with Ligand-of-Numb protein X1, and JAM4 and ESAM bind the membrane-associated guanylate kinase protein.²¹

RESPIRATORY DISEASES

Epithelial cells form a tight barrier against environmental stimuli via TJs and AJs. Defects in TJ and AJ proteins may cause changes in epithelial morphology and integrity and potentially result in more rapid trafficking of inflammatory cells through the epithelium. House dust mite (HDM) allergens are important factors in the increasing prevalence of asthma. The lung epithelium forms a barrier that allergens must cross before they can cause sensitization. The cysteine proteinase allergen Der p 1 from fecal pellets of the HDM *Dermatophagoides pteronyssinus* disrupts intercellular TJs, which are the principal components of the epithelial paracellular permeability barrier. These data suggest that opening of TJs by environ-

mental proteinases may be the initial step in the development of asthma in response to a variety of allergens.²²

Lipoxin A4 (LXA4) is a biologically active eicosanoid produced in human airways that displays anti-inflammatory properties. LXA4 plays an important role in the regulation of TJ formation, and stimulation of ZO-1 localization and expression at the plasma membrane occurs through a mechanism involving the LXA4 receptor.²³

HDM peptidase is capable of increasing epithelial permeability and thereby creating conditions that favor trans-epithelial allergen delivery.^{24,25} Lower epithelial α -catenin, E-cadherin, and/or ZO-1 expression in patients with atopic asthma contributes to defective airway epithelial barriers and higher influx of eosinophils into the epithelium. Der p 1 could contribute to sensitization and allergic responses by degrading airway epithelial barrier function.²⁴ Pollen peptidases directly or indirectly disrupt epithelial tight junctions, and this activity should be considered a possible mechanism for facilitating allergen delivery across epithelia.²⁶

In the normal lung, bronchiolar epithelial cells predominantly express CLDN 1, 3, 4, 7, and 18, whereas alveolar type II epithelial cells predominantly express CLDN 3, 4, 7, and 18.²⁷ In epithelial cells, transgenic expression of

CLDN1 with CLDN3 increases transepithelial resistance and decreases paracellular permeability, whereas CLDN4 confers a selective ion transport function without affecting paracellular solute permeability.²⁸ CLDN4 inhibition decreases transepithelial resistance without altering paracellular permeability in primary rat and human epithelial cells.²⁹ CLDN5, although expressed weakly in the epithelium, is strongly expressed in normal lung endothelium and is very highly expressed in interstitial pneumonia endothelium.³⁰ Newborn gene-targeted *Cldn5*^{-/-} mice die within 10 hours of birth, possibly as the result of altered permeability of the blood-brain barrier.³¹ When CLDN5 is transfected into airway epithelial cells, paracellular permeability increases even in the presence of excessive CLDN1 and CLDN3.³² Moreover, inducing CLDN5 expression in leaky rat lung endothelial cells can enhance paracellular barrier function against large, but not small, molecules.³³

An integral membrane protein, CLDN5 (Fig. 2), is a critical component of the endothelial TJs that control pericellular permeability. The breach of endothelial barriers is a key event in the development of pulmonary edema during acute lung injury. A major irritant in smoke, acrolein, can induce acute lung injury, perhaps by altering CLDN5

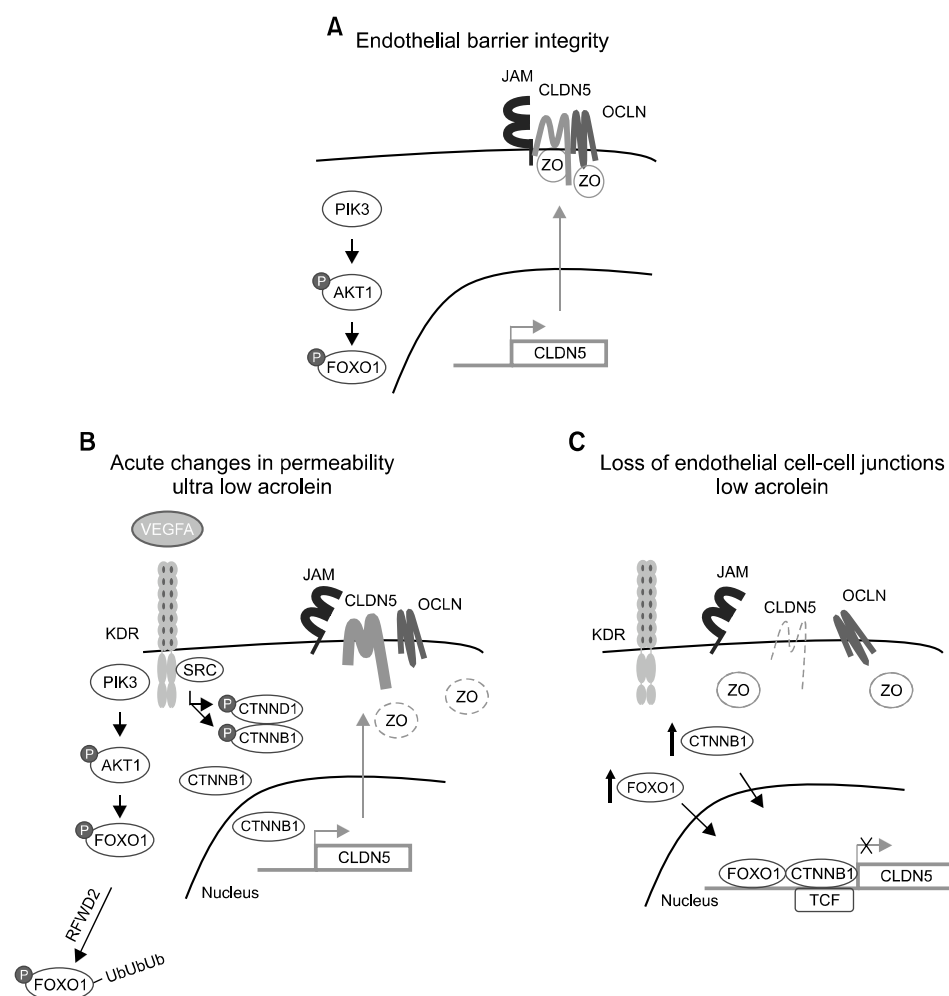


FIG. 2. Acrolein can induce acute lung injury through changes in claudin 5 expression. (A) Endothelial barrier integrity. (B) Acute changes in permeability. (C) Loss of endothelial cell-cell junctions.

expression. The phosphorylation status of the FOXO1 and CTNNB1 transcription factors is consistent with the observed alteration in CLDN5 expression.³⁴ Thus, preservation of endothelial CLDN5 may be a novel clinical approach to treatment of acute lung injury.³⁴

AJC MODULATORS

Safe and effective routes for drug delivery continue to be widely studied. Tissue barriers associated with such non-invasive delivery pose a considerable challenge, particularly for macromolecules.³⁵ This challenge can be met by employing permeation enhancers to facilitate paracellular transport.³⁶ Permeation enhancers can be selected empirically from among currently acceptable excipients for pharmaceutical use, or novel TJM excipients can be developed rationally.³⁶ The growing knowledge of TJ biology, the development of high-throughput cell/tissue-based assay systems, and the use of diverse molecular libraries have led to the identification of promising TJ-modulating compounds that safely and reversibly open TJs to enhance tissue permeability and drug transport.³⁷ New-generation peptide and lipid compounds, identified as TJ modulators, are amenable to optimization by structure-activity analyses and detailed mechanistic studies. This will continue to improve bioavailability and extend the molecular weight range of drugs that can be delivered effectively for chronic applications. The dynamic nature of TJs and the involvement of specific endocytic pathways and intracellular signaling mechanisms that regulate their function offer new possibilities for the development of drugs with more specific mechanisms of action, including the ability to restore normal barrier properties to TJs that are dysregulated during respiratory disease.

In conclusion, the AJC proteins, such as occludin, claudins, and JAMs, have substantially contributed to our understanding of the molecular mechanism of the AJC. Although knowledge regarding AJC membrane proteins has accumulated, several unsolved questions remain. In particular, the significance of AJC proteins in respiratory diseases should be examined further.

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CONFLICT OF INTEREST STATEMENT

None declared.

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