

Review

Cadherins: an integral role in inflammatory bowel disease and mucosal restitution

ANDREW P. ZBAR¹, CONSTANTINOS SIMOPOULOS², and ANASTASIOS J. KARAYIANNAKIS²

¹Department of Surgery, University of the West Indies, School of Clinical Medicine and Research, Bridgetown, Barbados

²Second Department of Surgery, Democritus University of Thrace, Medical School, 6 I. Kaviri Street, 68 100 Alexandroupolis, Greece

The intestinal epithelium is characterized by rapid cellular turnover with continuous proliferation of multipotential stem cells within Lieberkuhn's crypts, cellular migration along the crypt–villus axis, cellular differentiation, polarization, apical apoptosis, and luminal shedding. These processes are critical for the development and maintenance of normal intestinal epithelial architecture and function and involve complex cell–cell and cell–substratum interactions, which are mediated by epithelial (E)-cadherin and the integrins, respectively. This review outlines the role of E-cadherin and its cytoplasmic binding proteins, the catenins, as well as the interplay with other mucosal adhesion and restitution molecules during physiological processes in the intestinal epithelium mediating embryogenesis, cellular differentiation, cellular migration, and mucosal repair, as well as what is known about the dysregulation of assembly of the E-cadherin–catenin adhesion complex in inflammatory bowel disease.

Key words: E-cadherin, catenins, cell adhesion, inflammatory bowel disease, mucosal restitution, trefoil peptides

Introduction

The regulated formation of cell–cell and cell–matrix junctions plays a crucial role in normal embryogenesis and in the induction and physiological maintenance of cellular architecture and integrity in adult tissues.^{1–3} Over the past decade, several related families of cellular adhesion molecules involved in these processes have been identified, including the cadherins (which control

cell–cell interactions), the integrins (which mediate cell–stromal cross talk), the selectins (which are coexpressed on leukocytes and endothelial cells), and members of the immunoglobulin and proteoglycan superfamilies.⁴

Specialized intestinal epithelium undergoes rapid cellular turnover with controlled proliferation of multipotential stem cells within the crypts of Lieberkuhn, ordered cellular migration along the crypt–villus axis, cellular differentiation, polarization, apical apoptosis, and luminal shedding. This coordinated process is central for the development, morphogenesis, and maintenance of normal intestinal epithelial architecture, providing a specific interaction between molecules responsible for cellular integrity and mucosal repair. Inflammatory bowel disease and its *in vitro* wounded colonic epithelial monolayer models have shown altered cadherin expression^{5,6} as well as alterations in the constitutively expressed trefoil peptides, which have been implicated in normal gastrointestinal epithelial restitution.^{7–9}

This review outlines the role of epithelial (E)-cadherin and its cytoplasmic binding proteins, the catenins, and the interplay with other mucosal adhesion and restitution molecules during physiological processes in the intestinal epithelium mediating embryogenesis, normal mucosal differentiation, cellular migration, and mucosal repair, as well as what is known about the dysregulation of assembly of the E-cadherin–catenin adhesion complex in inflammatory bowel disease.

The E-cadherin–catenin complex and integrins in normal intestinal mucosa: evidence for receptor cross talk

E-cadherin (also known as L-CAM, uvomorulin, ARC-1, or cell-CAM 120/80), is the principal mediator of cell–

cell adhesion between epithelial cells.¹ It is a 120-kDa transmembrane glycoprotein that is localized mainly to the zonula adherens junction, signaling extracellular domain cell–cell adhesion through calcium-dependent homotypic interactions. Its carboxy-cytoplasmic domain is associated with a group of undercoat cytoplasmic proteins, the α -, β -, and γ -catenins, which share common structural and functional features and which are necessary for direct and indirect E-cadherin binding.¹⁰ Each catenin contains the Arm domain (a 42-amino-acid motif originally described for the *Drosophila* segment polarity gene product armadillo),¹¹ with both β - and γ -catenins binding directly to E-cadherin and α -catenin linking bound β - and γ -catenin to the

actin microfilament network of the cellular cytoskeleton.^{12,13} Binding is essential for the establishment of stable cell–cell adhesion, and survival studies have shown that structural and functional integrity of the components of the complex is necessary for the basic adhesive function of E-cadherin.^{14–20} Recently, a novel member of the catenin family, the p120 protein, has been described.²¹ This molecule has been identified as a common substrate for several receptor tyrosine kinases, sharing some homology with both β - and γ -catenin and binding directly with the cytoplasmic domain of E-cadherin^{22,23} (Fig. 1).

The integrins are transmembrane glycoproteins with overlapping receptor specificity expressed along the

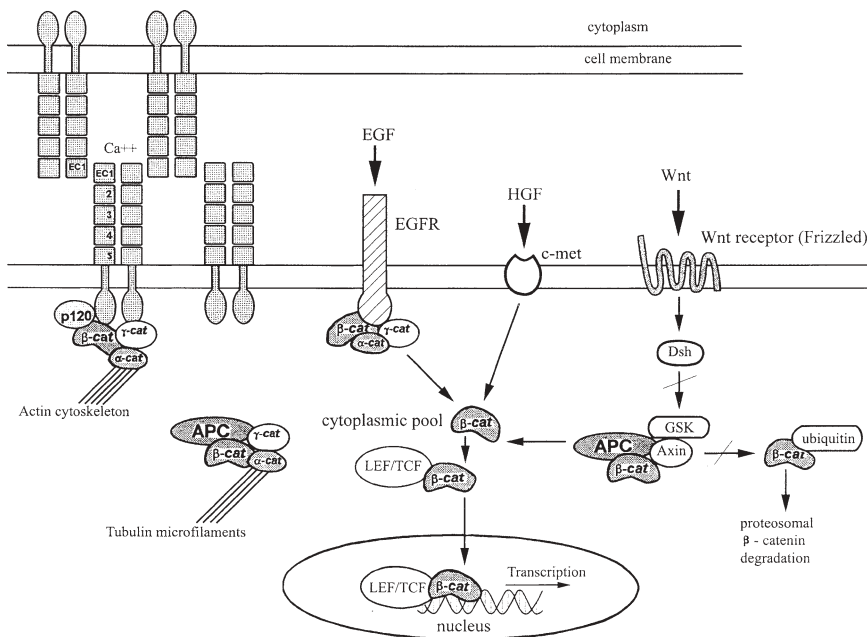


Fig. 1. Schematic presentation of E-cadherin and the catenins and their cellular interactions with each other and with other proteins. The extracellular domains of the E-cadherin homodimers bind in the presence of calcium ions in a “zipper” fashion to establish tight physical cell–cell adhesion. Their cytoplasmic domains bind to β - or γ -catenin in a mutually exclusive pattern whereas α -catenin links bound β - or γ -catenin to the actin cytoskeleton. p120 also binds to the cytoplasmic domain of E-cadherin in a distinct and separate site to that of other catenins. Catenins also form E-cadherin-independent complexes with other growth regulatory and signaling proteins such as the epidermal growth factor receptor (*EGFR*) and the adenomatous polyposis coli (*APC*) gene product and through this gene product to the tubulin microfilament network, the Wnt-signaling pathway, T-cell factor 4 (*TCF*), and lymphoid enhancer factor 1 (*LEF*) transcription factors (see text). The presence of a free cytoplasmic β -catenin pool is postulated to play an important role in controlling the stoichiometry of cadherin–catenin interactions and to influence the titration of catenins between cadherin and APC complexes. In the absence of Wnt ligand–receptor interaction, glycogen synthetase kinase 3- β (*GSK*) is activated and, with wild-type APC and axin, binds to free cytoplasmic β -catenin, resulting in its degradation through an ubiquitination and proteasome pathway. In the presence of a Wnt signal, *GSK* activity is inhibited in response to activation of the phosphoprotein dishevelled (*Dsh*), resulting in decreased degradation of β -catenin and elevated cytoplasmic levels of β -catenin, permitting interaction with transcription factors (*TCF/LEF*). This complex translocates to the nucleus and modulates the transcription of target genes including negative regulation of E-cadherin gene expression itself. Other abnormalities in the Wnt signaling pathway, such as mutated APC gene product, mutated β -catenin, or tyrosine phosphorylation of β -catenin on epidermal growth factor receptor (*EGFR*) and hepatocyte growth factor receptor (*c-met*) activation, can also result in an increased cytoplasmic pool of β -catenin, with modulation of the stoichiometry of cadherin–catenin interactions and subsequent perturbation of E-cadherin-mediated cell adhesion resulting in enhanced cell motility and cell migration. α -cat, α -catenin; β -cat, β -catenin; γ -cat, γ -catenin; *APC*, adenomatous polyposis coli; *c-met*, hepatocyte growth factor receptor; *Dsh*, dishevelled; *EC1*, extracellular domain; *EGF*, epidermal growth factor; *EGFR*, epidermal growth factor receptor; *GSK*, glycogen synthetase kinase 3- β ; *HGF*, hepatocyte growth factor; *LEF*, lymphoid enhancer factor 1; *TCF*, T-cell factor 4

basolateral surface of enterocytes and on the cell surface of all cells derived from the three primary germ cell layers, being composed of α - and β -chains associating noncovalently as heterodimers. Their extracellular domains contain ligand-binding regions whereas the cytoplasmic domains comprise regions capable of binding to particular cytoskeletal elements including talin, vinculin, and α -actinin.² These molecules modulate cellular interactions with components of the extracellular matrix including collagen, laminin, and fibronectin as well as with other integral cell-surface plasmalemmal proteins, with at least 15 different α - and 8 different β -chains for combination as more than 20 heterodimeric combinations.² They represent a vital heterophilic interaction between integrins and cadherins because specialized intraepithelial lymphocytes with the cytotoxic (CD8+) surface receptor phenotype expressing the α E β 7 (CD 103) integrin also lie along the basolateral region of enterocytes, where they are believed to regulate tissue-specific lymphocyte tracking and adherence during lymphocyte movement from the circulation to gastrointestinal-associated lymphatic tissue of the Peyer's patches.^{24,25} Here, the inhibition of cadherin-mediated cellular adhesion has been shown to down-regulate constitutive integrin expression in differentiating cells.^{26,27} Further evidence of cadherin-integrin cross talk in intestinal epithelium is provided by studies showing that the *c-erbB-2* oncogene coinhibits E-cadherin and $\alpha_2\beta_1$ integrin (an integrin acting as a principal collagen receptor in epithelial cells) gene transcription in human mammary epithelial cell culture.²⁸

Although integrins were originally identified as mediators of cell-matrix adhesion, they have recently been shown to participate in the intrinsic organization of the actin cellular cytoskeleton and in bidirectional transmembrane signal transduction through tyrosine phosphorylation of focal adhesion kinases, increases in intracellular calcium levels, inositol lipid and cyclin synthesis, activation of mitogen-stimulated kinases, and changes in crucial cellular growth genes.^{3,29} Catenins, apart from linking E-cadherin to the actin cytoskeleton, also link the cell membrane to downstream intracellular signaling pathways, providing a further interaction between integrins and cadherins as well as contributing to E-cadherin-independent binding to growth regulatory and signaling proteins such as epidermal growth factor (EGF),³⁰ the *c-erbB-2* oncogene product,^{31,32} and the adenomatous polyposis coli (APC) gene product.^{33,34} Here, the APC gene product, which is an integral mutated protein in the colorectal adenoma-carcinoma sequence as well as in colitis-associated carcinoma,³⁵ has been shown to control intracellular β -catenin levels and APC-catenin and E-cadherin-catenin complex assembly^{36,37} by competing directly with E-cadherin for β -catenin binding and forming separable α -, β -, and γ -

catenin complexes that are independent of cadherin.³⁷⁻³⁹ In this respect, it has recently been shown that APC mutations (similar to β -catenin mutations) appear to enhance the association of β -catenin with transcription factors such as T-cell factor 4 and lymphoid enhancer factor 1, both of which have been implicated in the transcriptional regulation of E-cadherin⁴⁰⁻⁴² (see Fig. 1).

In effect, integrins, catenins, and cadherins play complex integrative roles in signaling for the cellular motility and migration, proliferation, differentiation, and apoptosis, with stimulatory and inhibitory interactions that are affected by constitutive mutations and mutations in central genes like the APC gene. This complicated association in normal intestinal homeostasis and its interaction with growth factors, cytokines, and hormone-induced signaling in health and colitic disease remain to be defined.^{3,43}

The E-cadherin-catenin complex in embryogenesis, morphogenesis, and differentiation

The expression of cadherins is regulated during development. In embryogenesis, a complex variety of morphological events denote cellular aggregation and epithelial/mesenchymal transition, partly as a result of differential expression and spatiotemporal changes in the surface cellular expression of the different cadherins.⁴⁴ Experimental evidence in developmental models has shown a crucial role for E-cadherin in early morphogenesis where expression of a dominant negative cadherin mutant in *Xenopus laevis* embryos results in perturbation of normal cell adhesion and normal tissue polarity.^{45,46} Similarly, mutations of the β -catenin homologue in *Drosophila* (armadillo) create defective segment polarity⁴⁷ whereas ectopic overexpression of either β - or γ -catenin induces body axis duplication in the *Xenopus* embryo.^{48,49} Here, the timed administration of antisense β -catenin oligonucleotides or β -catenin-specific monoclonal antibodies leads to defective dorsal development.^{50,51}

The role of the cadherins and catenins in embryonic development and morphogenesis of the intestinal epithelium has been evaluated in vivo by gene targeting experiments in mouse embryonic stem cells and by generating transgenic and chimeric animals. Studies on E-cadherin gene knockout mice have revealed that E-cadherin-mediated cell adhesion is essential for the compaction of mesenchymal cells and their transition into a polarized epithelium. Homologous deletion of E-cadherin is also associated with lack of trophoblast epithelium or blastocyst cavity formation, resulting in early embryonic lethality.^{52,53} Similarly, β -catenin deletion results in failed mesoderm formation with a disturbed ectoderm development and early

death.⁵⁴ Further evidence for the crucial role of cadherin-mediated cell adhesion within the intestinal epithelium comes from studies using chimeric-transgenic mice in which islets of enterocytes transfected with truncated N-cadherin lacking the extracellular domain were introduced into C57 BL/6 mouse blastocysts, where isolated expression of this construct in the villus enterocytes results in disruption of normal cell–cell and cell–matrix adhesion.⁵⁵ Perturbation in cell–cell adhesion in this model was associated with increased cell migration along the crypt–villus axis, loss of cellular differentiation and polarization, and increased apoptosis.⁵⁶ When the dominant negative N-cadherin mutant was expressed along the entire crypt–villus axis, disrupted cell–cell adhesion resulted in increased cell proliferation and apoptosis in the crypts and increased cell migration in the villi. Alterations of the cell cycle in the crypts caused adenoma formation, impairing mucosal barrier function and inducing a progressive transmural inflammation resembling Crohn’s disease. In this model, histopathological changes observed in the adult chimeric mice included lymphoid aggregates, cryptitis, and crypt abscesses with goblet cell depletion, Paneth cell hyperplasia, and aphthoid or linear mucosal ulcers.⁵⁵ In contrast, forced expression of E-cadherin in the same model resulted in decreased cell proliferation and migration with increased apoptosis of cells in the upper part of the intestinal crypt.⁵⁷

Cell adhesion and epithelial cell migration

Disturbance or modulation of cadherin–catenin and APC–catenin cell–matrix adhesion and phenotypic expression may have an important place in the cellular migration process, which is believed to be necessary for the initial stages of tumor metastasis. Cellular migration is a fundamental process involved in many physiological and pathological processes. It is now apparent that cell motility requires dynamic and coordinated interactions between cells, their extracellular substratum, and the cytoskeletal network.⁵⁸ The transition from a stationary to a motile cellular phenotype is associated with spatial and temporal changes in cell–cell and cell–matrix interactions, which involve perturbation of E-cadherin-mediated cell adhesion and the coordinated modulation of the expression and function of integrins.^{59,60}

The extending lamellipodia at the leading edge of the migrating cell bind to the substratum through integrin receptors to establish stable cell–substratum adhesion and to generate adequate traction force for forward movement, followed by release of adhesion contacts at the rear of the cell.⁵⁹ Both β - and α -catenin have been shown to bind to the actin bundling proteins fascin and α -actinin, respectively,^{61,62} with these proteins being cru-

cial for the dynamic assembly and organization of actin bundles and networks during the extension of the cellular lamellipodia. The interaction between catenins and proteins involved in cell motility requires an organized regulation of E-cadherin/catenin-mediated cell–cell adhesion and cell migration where disruption of E-cadherin-mediated intercellular adhesion can occur through downregulation of E-cadherin expression and/or function, altered E-cadherin binding to β -catenin through tyrosine phosphorylation of β -catenin by type I tyrosine kinase growth factor receptors (such as the EGF receptor), or stoichiometric changes in the cytoplasmic pool of catenins and their complexes (see Fig. 1). This latter mechanism influences the titration of catenins between E-cadherin and other protein complexes impairing the adhesive function of E-cadherin. Here, the interactions between β -catenin and APC gene products appear to play a central role in cell migration, with APC localizing to the tip of the microtubule bundles⁶³ as well as to the end of cell processes.⁶⁴ In this respect, wild-type APC has been shown to induce disordered cell migration in the intestinal epithelium⁶⁵ and to downregulate the cytoplasmic levels of β -catenin,³⁷ where it is possible that a shift toward the “motility complexes” of β -catenin–APC or β -catenin–fascin could result in decreased β -catenin binding to E-cadherin, loss of E-cadherin-mediated intercellular adhesion, increased cell motility, and an induction of cell migration. An improved understanding of these interactions may provide future avenues for potential gene therapies in disordered cell migration associated with micrometastatic disease.^{36,66}

Cellular adhesion and inflammatory bowel disease

Ulcerative colitis and Crohn’s disease represent gastrointestinal diseases which, although clinicopathologically distinct, share common characteristic features of chronic inflammation, epithelial damage, mucosal ulceration, and epithelial restitution. It is now apparent that the reparative mechanism of epithelial restitution is induced immediately after superficial ulceration of the gastrointestinal mucosa⁶⁷ consisting of a rapid response mechanism of migrating surviving epithelial cells from the ulcer margins over the denuded area to cover the defect with reestablishment of epithelial continuity to restore mucosal barrier function.^{68–70} During this phase, there is no cell proliferation as such with cell migration acting as the main factor contributing to mucosal repair. Later, both cell proliferation and differentiation occur to establish normal villus architecture and absorptive/secretory function.

There is considerable evidence that cell migration occurring during epithelial restitution is associated with

alterations in E-cadherin and catenin expression and E-cadherin-mediated cell adhesion. This change has been noted in *in vitro* wounded monolayer models using colorectal cancer cells⁶ where there is reduced membranous expression and a cytoplasmic localization of E-cadherin, α -catenin, and p120, an effect that has also been demonstrated *in vivo* in peptic ulcer disease, Crohn's disease, and ulcerative colitis tissue.^{5,6,71} The changes observed in these *in vivo* models occur in the regenerating and migrating epithelium adjacent to ulcers and appear to correlate directly with disease activity. Differential changes in the expression of the E-cadherin–catenin complex in inflammatory bowel disease result in focal upregulation of these adhesion molecules in areas of active inflammation with downregulation in the reparative epithelium of healing ulcers, absent expression of the E-cadherin–catenin complex in the ulcer-associated cell lineage, and dysregulation of epithelial desmoid and tight junction molecules and mRNA expression.^{72,73} All these changes correlate with the recognizable severity of intestinal inflammation.

Decreased E-cadherin and catenin expression and subsequent disruption of E-cadherin-mediated cell adhesion (probably in response to the stimulatory effect of various motogenic factors) will enhance cell motility and promote cell migration in the regenerating mucosa. Importantly, a decrease in α E β 7-integrin-positive intestinal intraepithelial lymphocytes has also been demonstrated in the intestinal mucosa from patients with Crohn's disease⁷⁴ along with a focal upregulation of the E-cadherin–catenin complex in the subclinically inflamed bowel mucosa derived from spondyloarthropathy patients,⁷⁵ indicating an inverse relationship between α E β 7-integrin-positive lymphocyte infiltration and E-cadherin expression. This associated downregulation of epithelial tight junction molecules with cellular permeability and lymphocyte transmigration occurs in accordance with colitis activity and the activity of extracolonic disease.⁷⁶ Further evidence suggests that there is a differential expression of both E- and P-cadherins in inflammatory bowel disease, where left-sided ulcerative colitis results in either the downregulation of E-cadherin or specific single-base-pair mutations resulting in upregulation of P-cadherin specifically in dysplastic ulcerative colitis tissue.^{66,71}

Many recent studies have shown that aberrant 5'-CpG DNA methylation plays a vital role in the inactivation of tumor suppressor and DNA repair genes or as an early *de novo* phenomenon in colorectal carcinogenesis.⁷⁷ This finding has been shown with gene methylation-specific polymerase chain reaction techniques⁷⁸ in aberrant crypt foci,⁷⁹ adenomas,⁸⁰ and in dysplasia-associated ulcerative colitis.⁸¹ These CpG islands are short stretches of CpG-rich regions frequently associ-

ated with the promoter regions of specific genes that may undergo age-related methylation (so-called type A methylation), affecting the growth and differentiation of cells and cancer-specific methylation (so-called type C methylation), which produce a tumor-specific CpG island methylator phenotype.^{82,83} The manner in which these changes in DNA methylation of critically important cell regulators such as E-cadherin cooperate in aberrant mucosal healing and dysplasia is at present relatively poorly understood, although as a feature of dysplastic samples in ulcerative colitis mucosa, it suggests that it is subject to epigenetic regulation during colonic ulceration and may be potentially useful as a marker in those patients with long-standing colitis at risk for malignant degeneration.⁸⁴ An improved understanding concerning the small subsets of colorectal cells that express such mutant cadherins associated with areas of active ulceration and epithelial resistance to mucosal damage and denudation, or which are more likely to display an inherently dysplastic phenotype, will assist in the development of new therapies and in defining those patients at risk for the colitis–carcinoma sequence.

The trefoil peptide family of molecules represent a group of highly protease-resistant peptides actively secreted onto the intestinal mucosal surface that promote mucosal repair through epithelial restitution following injurious insult. These factors are allied to a number of agents such as EGF, the transforming growth factors alpha (TGF- α) and beta (TGF- β), and the hepatocyte growth factor/scatter factor (HGF/SF), which with the trefoil peptides has a specific motogenic effect promoting cell migration during epithelial restitution and mucosal repair through modulation of cell–cell and cell–matrix adhesion. Both EGF and TGF- α have been shown to promote cell migration of colonic epithelial cells *in vitro* and to upregulate the functional activity of $\alpha_2\beta_1$ and $\alpha_3\beta_1$ integrin receptors for laminin and collagen.^{85,86} Conversely, blocking of α_2 and β_1 integrin chains by specific monoclonal antibodies inhibits cell migration, whereas TGF- α stimulation results in an increased binding of $\alpha_2\beta_1$ integrin to the E-cadherin–catenin complex through β -catenin,⁸⁷ suggesting that the interactions between integrins and E-cadherin affect the stability of cell–matrix binding that is normally necessary during cell migration.

The trefoil peptides are a highly conserved family of small, secretory-stable molecules bearing one or more trefoil structural motifs of three interchain loops secured by disulfide bonds.⁷ To date, three trefoil peptides have been identified in humans; namely, pS2 (trefoil factor 1, TFF1), human spasmolytic polypeptide (trefoil factor 2, TFF2), and intestinal trefoil factor (trefoil factor 3, TFF3). These peptides are constitutively expressed in the gastrointestinal tract in a region-

ally specific pattern, particularly in association with mucins, implying a role in maintaining the mucosal barrier. They are also ectopically overexpressed within gastrointestinal mucosa affected by chronically ulcerating and inflammatory lesions such as peptic ulceration, Crohn's disease, and ulcerative colitis.^{8,88–90} In vitro studies using wounded monolayers as a model of mucosal injury have shown that at least two of these peptides (TFF1 and TFF2) stimulate cell migration and promote wound healing,^{91,92} whereas in vivo both TFF2 and TFF3 appear to provide sufficient mucosal protection to reduce the extent and severity of ethanol- and indomethacin-induced acute gastric injury in the rat.^{91,93,94} Following experimentally induced mucosal ulceration, there is an immediate increase in TFF2 expression in the ulcerated area, followed by an ectopic expression of TFF3.⁹⁵

Targeted deletion of the gene encoding TFF3 in particular produces a high sensitivity to colonic injury, in which activation of these pathways redistributes E-cadherin from the cell surface to an intracellular locale resulting in catenin complexing, kinase activation, and apoptosis. In this regard, the trefoil peptides appear to block p53-dependent and p53-independent apoptosis enabling cellular migration with or without the programmed cell death normally associated with cellular detachment. Other trefoil peptides, most notably TFF3, require disturbance of calcium-dependent E-cadherin function where they have been shown to stimulate the migration of HT29 colorectal cancer cells by phosphorylation of β -catenin and reduced E-cadherin expression.⁹⁶ Moreover, knockout TFF3 mice tend to show an expanded intestinal proliferative epithelial phenotype with impaired mucosal healing,⁹⁷ whereas transgenic mice overexpressing TFF1 have an enhanced resistance to mucosal intestinal injury.⁹⁸ In this way, local trefoil peptides modulate E-cadherin-mediated cellular adhesion and cadherin–catenin complex formation by their interaction with key growth factor receptors.

The evidence that the stimulatory effect of various growth and trefoil factors has on cell migration and epithelial restitution acts through direct modulation of the E-cadherin adhesion system comes mostly from in vitro work with tumor cell lines where frameshift deletions in E-cadherin gene repeat regions result in replication error-positive cell lines with altered cellular adhesion, differentiation, proliferation, and tumorigenicity as well as an altered sensitivity to motogenic trefoil factors; all these effects are restored to the cell lines by full-length E-cadherin cDNA transfection.⁹⁹

Because tyrosine phosphorylation of β -catenin appears to be a common mechanism by which the motogenic factors express their effects in migrating cells, it is conceivable that β -catenin phosphorylation results in dissociation of the E-cadherin–catenin com-

plex from the actin cytoskeleton with accumulation of β -catenin in the cytoplasmic pool. This effect may contribute to disassembly of adherens junctions, diminished cell–cell adhesion, and increased cell motility. The cytoplasmic accumulation of β -catenin may involve additional interactions with other proteins, most notably the APC protein as a central regulator of directed cell migration.³⁶ The synergistic effect of TFF3 and EGF on cell migration suggests a functional or physical interaction between these peptides and the relevant receptors, possibly through β -catenin.^{94,96}

In conclusion, the efficient and coordinated repair of the gastrointestinal mucosa safely prevents the entry of pathogens and antigens across the intestinal lumen. There is now considerable evidence from wounded monolayer and primary cell culture models showing that epithelial cell migration is vital in this epithelial restitution with a complex interplay between and tight spatiotemporal control of soluble extracellular molecules (growth factors, trefoil peptides, and cytokines), matrix components (collagen, laminin, and fibronectin), signaling surface receptor-specific kinases and principal regulators of cell–cell (cadherin) and cell–matrix (integrins, hyaluronate receptors) adhesion. The E-cadherin–catenin complex appears to function as a master molecule in the regulation of cell adhesion, polarity, differentiation, migration, proliferation, and survival of gastrointestinal epithelial cells. Its fundamental role in embryogenesis has been clearly defined by the lethality seen in E-cadherin–catenin knockout mice, and loss of E-cadherin–catenin function is an important step in the development of both inflammatory and neoplastic lesions of the gastrointestinal tract. Much needs to be elucidated in human colitis and colitis models to provide potential therapies that can primarily influence epithelial restitution through deliberate E-cadherin and catenin modulation.

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