

A new understanding of enteroaggregative *Escherichia coli* as an inflammatory pathogen

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Abbreviations: AAF, aggregative adherence fimbriae; SCID-HU-INT mice, human intestinal xenografts in severe-combined immunodeficient mice; EAEC, enteroaggregative *E. coli*; HXA₃, heparin-binding hemagglutinin A₃; 12-LOX, 12-lipoxygenase; MRP2, multidrug resistance-associated protein 2; PKC- δ , protein kinase C- δ ; PLA₂, phospholipase A₂; PMN, polymorphonuclear leukocyte

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Enteroaggregative *Escherichia coli* (EAEC) is an important cause of endemic and epidemic diarrheal disease worldwide. Although not classically considered an inflammatory pathogen in the style of *Shigella* and *Salmonella* species, clinical data from patients suggests that inflammatory responses may play an important role during EAEC disease. However, the specific role of inflammation during EAEC pathogenesis has not been investigated in detail. To better understand how EAEC may induce inflammation, we have focused our attention on the intimate interactions between EAEC and the host epithelium and the subsequent induction of host cell signaling events leading to innate immune responses. Here, we discuss our recent findings on the signaling pathway by which EAEC promotes transepithelial migration of polymorphonuclear leukocytes (PMNs), the role of aggregative adherence fimbriae in triggering this event and the implementation of human intestinal xenografts in immunodeficient mice for studying EAEC pathogenesis *in vivo*. Our findings suggest that EAEC shares conserved mechanisms of inducing PMN recruitment with other intestinal pathogens, providing new insight into the potential pathological consequences of EAEC-induced inflammation.

as well as worldwide foodborne outbreaks.¹ While EAEC may in fact be one of the most common bacterial causes of diarrhea, the lack of global routine surveillance systems for detecting EAEC has likely rendered it underreported.² In the past year, however, focus on EAEC has increased following a major outbreak in Germany in 2011.³

EAEC pathogenesis results from colonization of the intestinal mucosa via a stepwise process of adherence and subsequent biofilm formation, which is followed by toxin release leading to secretion of intestinal fluids.¹

Several enteric pathogens, including *E. coli* pathotypes, are agents of inflammatory diarrhea, the histopathologic hallmark of which is infiltration of polymorphonuclear leukocytes (PMNs).^{4–7} The role of inflammation during EAEC pathogenesis has only recently been considered, and increasing evidence suggests that inflammatory responses may play a substantial role in EAEC pathology. Clinical studies have documented elevated levels of pro-inflammatory markers, including interleukin (IL)-8, IL-1 β and fecal lactoferrin and leukocytes, in EAEC-infected individuals.^{8–10} Therefore, unraveling the mechanisms underlying EAEC-induced inflammation and dissecting the role of these events in disease are important steps toward advancing the understanding of this emerging pathogen.

Introduction

Enteroaggregative *Escherichia coli* (EAEC) is an enteric pathogen increasingly recognized for causing acute and persistent diarrheal illness in developing countries,

Induction of Host Cell Inflammatory Signals by EAEC

Accumulation of PMNs at inflamed sites is a common outcome of colonization of mucosal surfaces by pathogenic bacteria.

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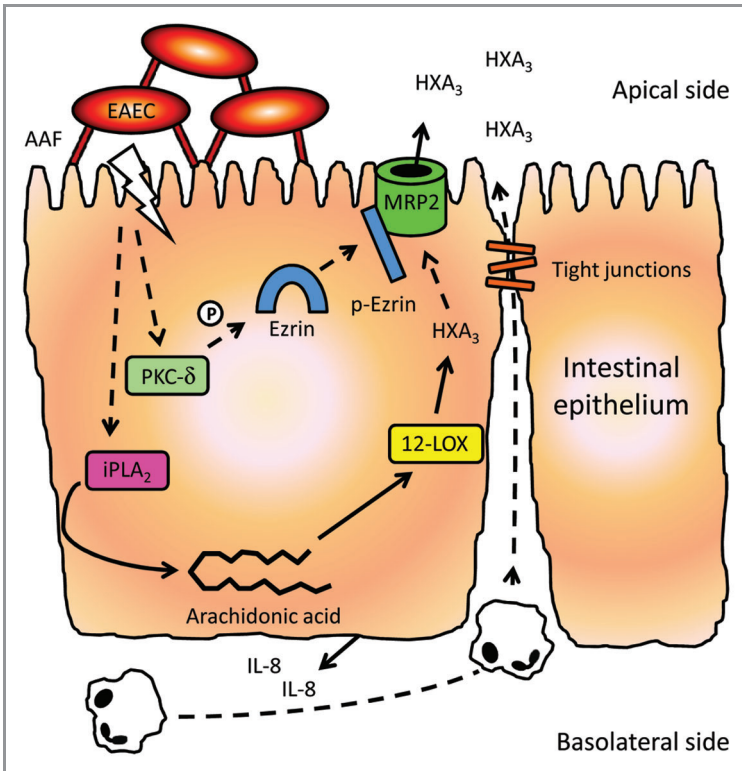


Figure 1. Model of EAEC-induced PMN transepithelial migration. AAF-mediated binding of EAEC to the surface of the intestinal epithelium triggers basolateral release of pro-inflammatory cytokines, e.g., IL-8, thus recruiting PMNs to the subepithelial space. Moreover, AAF binding also induces iPLA₂-mediated release of arachidonic acid from the host cell membranes. Through 12-LOX activity, arachidonic acid is then metabolized into HXA₃, which is then transported across the apical membrane by MRP2, thus generating a chemotactic gradient of the lipid across the tight junctional complex driving transepithelial migration of PMNs to the apical surface. PKC-δ also plays a key role in these inflammatory events, presumably by activating ezrin which in turn aids in facilitating transport of MRP2 to the apical membrane. Solid lines represent events supported by published results. Hashed lines represent speculations from our group. Modified from Boll et al.¹³

Intestinal epithelial cells respond by releasing cytokines and distinctive PMN-specific chemoattractants that—in combination with numerous adhesion molecules—recruit PMNs from the blood stream and direct their movement through endothelial and epithelial barriers to the luminal surface. Recruitment of PMNs is the first line of response of the host immune system to bacterial infection, geared toward destruction of invading pathogens. However, the nonspecific neutrophil effectors can cause collateral damage, thus potentially contributing to disease pathology. Moreover, several pathogens have evolved strategies to resist neutrophil killing or even benefit from eliciting inflammation.

In line with other enteric pathogens, previous studies have shown that EAEC infection of polarized intestinal epithelial

cells triggers mitogen-activated protein kinase signaling cascades that lead to nuclear factor kappa-B (NFκB) activation, which in turn stimulates the release of an array of pro-inflammatory cytokines, including the potent PMN chemokine interleukin (IL)-8.^{11,12} Thus, basolaterally released IL-8 likely plays a major role in recruiting PMNs to the subepithelial space in response to EAEC infection.

A recent study from our group shows that EAEC-induced migration of PMNs across the epithelium requires apical secretion of a second, lipid-based PMN chemoattractant (Fig. 1). Specifically, EAEC infection of polarized T84 colonic epithelial cells triggers calcium-independent phospholipase A₂ (iPLA₂)-mediated release of arachidonic acid from the cell membrane. Through enzymatic action of 12-lipoxygenase (12-LOX), arachidonic

acid is then metabolized into heptoxilin A₃ (HXA₃), a member of the eicosanoid class of lipids with potent PMN chemoattractant properties. EAEC infection also triggers an increase in expression of the apically located membrane ATP-binding cassette (ABC) transporter multidrug resistance-associated protein 2 (MRP2), which subsequently functions as an efflux pump for the vectorial release of HXA₃ to the apical surface. Secreted HXA₃ then forms a chemotactic gradient through the tight junctional complex, thus directing paracellular transit of PMNs across the epithelial monolayer to the luminal surface¹³ (Fig. 1).

Increasing evidence suggests that 12-LOX-mediated apical release of HXA₃ to promote PMN transepithelial migration is a conserved mechanism by which the intestinal epithelium responds to intruding inflammatory pathogens, including *Salmonella enterica* serovar Typhimurium (*S. Typhimurium*), *Shigella flexneri*, *Campylobacter* species and EAEC.¹³⁻¹⁵ Yet, the upstream events by which these pathogens elicit inflammation are very much distinct, reflecting their discrete strategies for promoting infection. As an example, *S. Typhimurium* and *S. flexneri* rely on effector proteins, translocated into the host cells by type III secretion systems (T3SS), to interact with host cells, leading to invasion of the epithelium and triggering and manipulation of innate immune responses.^{16,17} While *E. coli* pathotypes such as enteropathogenic *E. coli* and enterohemorrhagic *E. coli* also employ T3SS-dependent infection strategies, this does not appear to be the case for EAEC. Moreover, unlike *S. flexneri* and *S. Typhimurium*, EAEC strains generally do not invade the epithelium, and instead remain anchored in the intestinal mucosa.¹⁸ Our recent work shows that EAEC-induced PMN transmigration requires only binding of the bacteria to the apical epithelial surface, an event facilitated by aggregative adherence fimbriae (AAF), the principal adhesins of EAEC.¹⁹

The pro-inflammatory properties appear to be conserved among different variants of these adhesins as all four AAF subtypes identified thus far promote PMN transepithelial migration.¹⁹ The AAF subunits

are phylogenetically related to those of the Afa/Dr family of *E. coli* adhesins, all of which employ the chaperone-usher pathway for fimbrial assembly.²⁰ Notably, other members of this family, such as the F1845 adhesin and Dr hemagglutinin of diffusely adhering *E. coli* (DAEC), have also been shown to promote PMN transepithelial migration,⁶ thus inferring a common strategy of F1845/Dr/AAF-mediated inflammatory responses among these two *E. coli* pathotypes.

How Does EAEC Activate the 12-LOX Pathway to Trigger Inflammation?

How AAF-facilitated adherence of EAEC to the epithelium is linked to activation of the 12-LOX pathway is yet to be determined. However, binding of AAF to the extracellular matrix (ECM) proteins fibronectin, laminin and type IV collagen has been demonstrated. It is possible that AAF trigger host signal transduction indirectly by binding to ECM proteins that then interact with host cell receptors such as integrin $\alpha_5\beta_1$.²¹

ECM protein-mediated integrin signal transduction in epithelial cells has been shown to involve phosphorylation of protein kinase C delta (PKC- δ).²² Notably, EAEC infection of T84 cell monolayers triggers phosphorylation and translocation of PKC- δ to the cell membrane, and blocking of PKC- δ activity strongly attenuates EAEC-induced PMN transmigration.¹³ The exact role of PKC- δ in EAEC-induced inflammation warrants further investigation. However, a different PKC isoform, PKC- α , has been shown to play a central role in *S. Typhimurium*-induced PMN transmigration by phosphorylating ezrin which then associates with MRP2 and mediates localization of the membrane transporter to the apical surface.²³ We speculate that PKC- δ plays a similar role in the event of EAEC-induced HXA₃ secretion, as PKC- δ has been shown to be involved in facilitating translocation of MRP2 to the plasma membrane in rat hepatocytes.²⁴ In addition, PKC- δ activity has been shown to cause disruption of tight junctions of intestinal Caco-2 cells through oxidative injury,²⁵ whereas inhibition of PKC- δ

appears to confer enhanced barrier function to the cells by promoting expression and assembly of the tight junction proteins occludin and claudin-1.²⁶ Conversely, EAEC has been shown to cause epithelial barrier disruption in T84 cell monolayers through delocalization of occludin and claudin-1.²⁷ Based on these findings, it is evident that PKC- δ activity may play a role in several aspects of EAEC-induced inflammation.

The findings described above point to a role for ECM proteins and integrins in EAEC adherence and possibly in the elicitation of inflammatory responses. However, as ECM proteins are generally localized to the epithelial basement membrane, other receptors are likely to be involved in at least the early stages of infection, at which point the bacteria are restricted from gaining access to the ECM proteins. Additional unknown AAF receptors may therefore likely play a role in activating the 12-LOX pathway and triggering inflammation.

Eliciting Inflammation as a Pathogenic Strategy to Circumvent and Exploit the Host Immune Response

While host inflammatory responses are intended as a first line of defense, several pathogens have evolved sophisticated ways of subverting these events to promote infection and cause disease. For instance, *S. Typhimurium* utilizes an electron acceptor generated by the host respiratory burst during inflammation to gain a growth advantage over the intestinal microbiota.²⁸ In another example, *S. flexneri* benefits from the opening of tight junctions—a direct effect of PMN transmigration—to invade the intestinal epithelium from the basolateral surface.²⁹ Moreover, antimicrobial proteins released from neutrophil granules have been shown to enhance adherence of *S. flexneri* to epithelial cells during the initial steps of infection.³⁰ In a third example, PMN transmigration induced by Afa/Dr-expressing DAEC has been shown to trigger synthesis of tumor necrosis factor α and IL-1 β . This, in turn, upregulates apical surface expression of decay-accelerating factor, the receptor for Afa/Dr adhesins,

thus promoting enhanced bacterial colonization.³¹

Similar to DAEC, our studies show that PMN transepithelial migration enhances subsequent adherence of EAEC to T84 cell monolayers, suggesting that EAEC may also benefit from eliciting inflammation. This enhanced adherence is not due to loss of barrier integrity but rather a direct consequence of post-transmigratory PMN-mediated events that alter host cell signaling.¹³ Distinguishing them from Afa/Dr adhesins, AAF do not appear to bind to decay-accelerating factor.²¹ However, adenosine derived from neutrophils that infiltrate the lumen during active intestinal inflammation has been shown to trigger apical secretion of fibronectin from T84 cells.³² Moreover, adenosine facilitates enhanced adherence of EAEC to T84 cell monolayers, likely through AAF-fibronectin binding.²¹ Thus, enhanced EAEC adherence following PMN transmigration could be mediated by increased availability of ECM proteins on the apical surface. In addition, adenosine secretion stimulates fluid secretion from the epithelium.³³ Triggering of inflammation may therefore both enhance colonization of EAEC on mucosal surfaces as well as aid in the diarrheal spreading of the bacteria.

Apart from AAF, at least one other EAEC virulence factor has been identified as a contributing factor in causing or modulating host immune responses. Pic, a serine protease autotransporter of Enterobacteriaceae (SPATE) found in EAEC, *S. flexneri* and uropathogenic *E. coli*, cleaves mucin, induces mucus release and confers a growth advantage in a mouse model of mucosal colonization.^{34,35} More intriguingly, Pic was recently shown to target several Sialyl Lewis-X-modified glycoproteins on the immune cell surface, including CD43 and CD45. The effects of Pic activity on these leukocytes included impaired chemotaxis and migration of PMNs, activation of the PMN oxidative burst, and activation and apoptosis in T-cells. Moreover, a Pic mutant *S. flexneri* strain was found to induce a greater inflammatory response than the wild-type strain in a guinea pig keratoconjunctivitis model, implying that the overall effects of Pic are predominantly anti-inflammatory.³⁶ Thus, it is tempting to speculate a dual role

for Pic in EAEC pathogenesis in which Pic mediates penetration of the mucosal layers by the bacteria as well as counteracts the inflammatory response induced by AAF-mediated adherence. However, Pic may also act to enhance inflammatory responses by facilitating premature activation of PMNs and subsequent tissue damage or by causing mislocalization of adherent PMNs.³⁶ Thus, whether Pic-mediated immune modulation by EAEC contributes to pro- and/or anti-inflammatory effects in the human intestine remains to be determined.

Human Intestinal Xenografts as a Model for Studying EAEC Pathogenesis

Animal models provide useful means to study aspects of bacterial pathogenesis that cannot be addressed using cell culture-based studies such as the roles of mucus or the commensal flora in intestinal colonization, or how the complex interplay between different host cell types affect immune responses. However, implementation of suitable small animal models for studying EAEC disease has long been hampered by the fact that EAEC appears to be pathogenic only in the human intestinal tract.^{34,37} This may result from the inability of AAF adhesins to bind to different versions of receptors present on the mucosal surfaces of tested animal species as compared with humans.

To overcome this hurdle, we have recently employed human intestinal xenografts in severe-combined immunodeficient (SCID-HU-INT) mice as a novel model for studying EAEC disease and innate immune responses in vivo. These transplanted xenografts become extensively vascularized, secrete mucus and develop into morphologically normal human intestine.³⁸ The use of SCID-HU-INT mice as an infection model has previously been demonstrated for other enteric pathogens adapted to humans such as enterohemorrhagic *E. coli* and *Shigella* species.^{39,40} We have shown that EAEC induces extensive tissue damage and inflammation in the human intestinal tissues in this model as marked by PMN infiltration, goblet cell depletion and edema formation. Moreover, these

pathological markers—particularly inflammatory infiltrates—were strongly correlated with expression of AAF.¹⁹ The SCID-HU-INT mouse model offers an exciting opportunity to study other aspects of EAEC pathogenesis as well.

EAEC as an Emerging and Adaptable Pathogen: The 2011 German Outbreak

The need for an increased understanding of EAEC pathogenesis is emphasized by the major recent outbreak that took place in Germany in May–June 2011, which was caused by a highly virulent Shiga-toxin (Stx)-producing EAEC O104:H4 strain. Over 4,000 cases of diarrhea were reported during this outbreak, of which 22% of patients developed hemolytic-uremic syndrome (HUS), and 54 patients succumbed to the infection.³ This strain exhibited unusually high proportions of adults affected and ratio of HUS cases, as compared with previous outbreaks of Stx-producing enterohemorrhagic *E. coli* that typically caused more severe disease in children and the elderly and with an average rate of HUS of about 4%.⁴¹ In contrast to enterohemorrhagic *E. coli*, the 2011 German outbreak strain possesses EAEC-specific virulence factors, including AAF, as well as the three SPATE proteases Pic, SepA and SigA. Thus, the severity of clinical outcomes following infection during this outbreak suggests that the EAEC background conferred enhanced virulence to this strain.^{3,42}

Stx-induced systemic complications require transit of the toxin across the intestinal epithelium upon release from its bacterial host. Epithelial barrier disruption—caused either by the pathogen itself or by infiltrating PMNs responding to infection—is a potential route of paracellular Stx uptake.⁴³ Indeed, PMN migration induced by Stx-producing *E. coli* has been shown to enhance apical-to-basolateral translocation of Stx across polarized T84 monolayers.⁴⁴ Adding to the potential impact of inflammation on HUS development, H₂O₂ production by recruited PMNs has been shown to activate stress responses leading to induction of Stx prophages and thus Stx production.⁴⁵ Moreover, based on findings from clinical

studies, Exeni et al. have suggested that the intensity of PMN activation during infection with Stx-producing *E. coli* and the speed of onset of PMN impairment is proportionate to the severity of systemic disease.⁴⁶

Given the ability of EAEC prototype strains to induce both epithelial barrier disruption and PMN transmigration, host inflammation mediated by EAEC virulence factors is likely to play a key role in conferring enhanced virulence to the Stx-producing O104:H4 outbreak strain.

Concluding Remarks

Many enteric pathogens have evolved the ability to engage host cells in complex interactions that trigger inflammatory responses. Our recent work shows that intestinal cells respond to EAEC infection by releasing an arachidonic acid-derived eicosanoid generated through 12-LOX activity that causes PMN transepithelial migration. Distinguishing it from other inflammatory pathogens, EAEC-induced 12-LOX activation requires only binding of the EAEC-defining AAF adhesins. This emphasizes the concept of 12-LOX-mediated signaling as a conserved mechanism by which the intestinal epithelium instigates PMN recruitment to battle enteric pathogens. Reflecting co-evolution, several pathogens have in turn developed sophisticated ways to evade innate immune responses or even benefit from them. Recent studies by ours and other groups suggest that this is also the case for EAEC.

While our studies here provide important new insight into the role of inflammation in EAEC pathogenesis, there is still a lot to be learned. For example, we have yet to determine the “missing link” between AAF-mediated adherence to host cells and induction of the 12-LOX pathway. Moreover, the overall implications of inflammation in EAEC pathology requires further investigation.

Advances in understanding the interplay between EAEC and host cells contributes to our overall understanding of its pathogenesis and provides useful information that may ultimately help in preventing and/or treating disease caused by this emerging pathogen.

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