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Established and Emerging Concepts in Hirschsprung's-Associated Enterocolitis

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Abstract

Hirschsprung's disease (HSCR) is a common cause of neonatal bowel obstruction and the approach to diagnosis and surgical treatment is well defined and accepted. Hirschsprung's-associated enterocolitis (HAEC) remains a frequent cause of pre-operative and post-operative morbidity and mortality, with unchanged treatment guidelines over multiple decades. Recent advances in our understanding of the genetics underlying HSCR have allowed the development of animal models, some of which recapitulate the HAEC phenotype. These animal models, along with recent translational studies, have implicated multiple facets of mucosal immunity and microbiome dysbiosis in the development of HAEC. Here, we will review the established epidemiology, modes of diagnosis and treatment of HAEC. Furthermore, we will explore emerging concepts in the pathogenesis of this disease; including animal models, alterations in mucosal immunity, dysbiosis of the intestinal microbiome, specific genetic susceptibility, and novel treatment modalities.

Keywords

Hirschsprung disease; Hirschsprung's disease; enterocolitis; mucosal immunity; microbiome; pathogenesis

Introduction

In 1886, Harald Hirschsprung presented the clinical course and pathology specimens of two unusual patients to the International Congress for Children's Disease in Berlin[1]. Both patients had profound constipation from birth, demonstrated megacolon without dilation of the rectum, and ultimately expired from a sudden onset of acute abdominal distention, profound diarrhea, and weight loss/dehydration. The presentation of these cases by a world-renowned pediatrician prompted an outpouring of other, similar cases, being presented worldwide. The underlying disease ultimately came to be referred to as "Hirschsprung's disease" (HSCR, Online Mendelian Inheritance in Man #142623), although we now

recognize that the two children Professor Hirschsprung described were, in fact, suffering from Hirschsprung's-Associated Enterocolitis (HAEC).

Since the time of Harald Hirschsprung, some of the greatest minds in all of medicine engaged in the quest to understand the etiology and pathophysiology of HSCR and to develop treatment strategies[2]. This has led to an understanding of the underlying genetics of HSCR, methods to establish the diagnosis, and a variety of surgical options to resect the aganglionic portion of bowel. However, despite technically adequate surgery and histologic confirmation of removal of the pathologic, aganglionic segment, there is increasing recognition that the majority of Hirschsprung's patients continue to suffer from bowel dysfunction, soiling or incontinence, HAEC, and decreased quality of life[3]. Of these complications, HAEC has come under increasing scrutiny over the last 10–15 years (Figure 1), likely due to an increasing recognition of its associated morbidity and mortality. Here, we will review aspects of HAEC that are well established (Epidemiology, Diagnosis, Treatment) as well as those that, based on recent translational research, are emerging (Animal Models, Mucosal Immunity, Microbiome, Genetics, Treatment).

Established Concepts

Epidemiology

HSCR is a common cause of intestinal obstruction in the newborn, with an incidence that ranges worldwide from 1:2000 to 1:12000 live births, but is most commonly reported as 1:4000[4]. While previously thought to be rare in premature infants, recent analyses indicate that the incidence of HSCR is similar for pre-term and term births[5–7]. The incidence of HAEC reported in the literature is highly variable, with reports as high as 60%, although an incidence of 25–35% is generally quoted in current series[8, 9]. HAEC can occur pre-operatively or post-operatively, with a similar incidence in both settings. It is the presenting symptom of HSCR in 25% of infants and it is these infants that are most likely to experience mortality from the disease[10–13]. Mortality from HAEC ranges from 1–10%, with the majority of deaths occurring in newborns prior to definitive operation[13]. It is unclear if higher mortality in newborns is due to delay in diagnosis, immature immune mechanisms in the newborn, or disease-intrinsic differences from the pre-operative to post-operative period.

Diagnosis

The classic manifestations of HAEC are similar to those originally described by Professor Hirschsprung; abdominal distention, fever, and diarrhea. However, the spectrum of symptoms that children may present with additionally includes vomiting, rectal bleeding, lethargy, loose stools, and obstipation. These symptoms are non-specific, and may lead to alternative diagnoses (e.g. infectious enteritis) or delays in diagnosis. This is reflected the wide range of incidence of HAEC reported in the literature. Ultimately, this has resulted in difficulties in comparing outcomes based on treatment strategies, operative approaches amongst different centers. In an attempt to address the difficulty in establishing the diagnosis of HAEC, a group of 27 gastroenterologists and surgeons participated in a Delphi process[14], starting with 38 features (history, patient characteristics, physical exam signs, laboratory findings, radiology findings, pathology findings) and narrowing this list down to

16 items to develop a HAEC score[15]. Unfortunately, this tool has proven difficulty in routine bedside use, and has not been widely adopted. Currently, overall clinical gestalt based on observation of the classic manifestations remains the mainstay of diagnosis.

Treatment

Current therapy for HAEC is relatively non-specific and consists of systemic antibiotics, rectal irrigations and bowel rest. These measures are directed towards managing the factors that are thought to contribute to pathogenesis and treating the acute symptoms. Systemic antibiotics are used empirically in HAEC, with metronidazole typically chosen to treat anaerobes. This agent has activity against *Clostridium difficile*, an organism that has been associated with HAEC[16]. Fecal stasis and bacterial overgrowth are treated by rectal irrigations, which are distinct from enemas in which large volumes of fluid may be retained in the already-distended bowel. Additionally, rare patients presenting with severe sepsis or with perforation will benefit from proximal enteric diversion, although cases of persistent and recurrent enterocolitis following diversion have been described[17, 18].

Some authors have advocated for the use of preventive measures in selected patient populations[9]. These measures include routine use of rectal irrigations in the post-operative period, long-term administration of oral metronidazole, use of pro-biotic therapy, and diverting enterostomy. Unfortunately, no definitive evidence to support routine use of these “preventive” measures exists[19].

Development of a targeted therapy for HAEC has the potential to decrease mortality and increase quality of life.

Emerging Concepts

Animal models

Because of familial occurrence in 5–20% of patients, association with syndromes, and the existence of spontaneous mutant animals (piebald lethal and lethal spotted mice), a genetic basis for HSCR had long been suspected. The identification of gene defects contributing to HSCR has advanced our ability to study the disease in the laboratory. The *Rearranged during transfection (Ret)* gene, on the short arm of chromosome 10, was identified in 15 Hirschsprung’s disease pedigrees in the early 1990s[20, 21]. Subsequent to this, multiple genes contributing to Hirschsprung’s disease have been identified (*GDNF*, *NRTN*, *EDNRB*, *EDN3*, *ECE1*, *ZFHX1B*, *SOX10*, etc.)[22]. Although these gene defects account for less than 30% of sporadic mutations found in humans, animal models in which these genes are mutated have provided valuable insight into the pathogenesis of HSCR and HAEC.

Mutations of *RET* and of *Endothelin receptor B (EdnrB)*, both of which are involved in ENS formation, are the two most commonly identified HSCR gene defects in humans[22]. Recently, *Ret* has been shown to be necessary for the development of Peyer’s patches (PP), the primary inductive site for gastrointestinal host defense, which suggests a potential developmental link between the enteric nervous system (ENS) and gastrointestinal mucosal immunity[23]. Unfortunately, *Ret*^{-/-} animals exhibit a severe phenotype with an ENS absent distal to the stomach and renal agenesis, resulting in death shortly after birth and limiting

their utility in studying HAEC[24]. The second most common gene defect identified in patients with HSCR are mutations of *EdnrB*. *EdnrB* and its ligand, endothelin 3 (*ET-3* or *EDN3*), regulate enteric neural crest cell proliferation, migration and differentiation[25]. Murine models of *EdnrB* mutation display aganglionosis of the distal hindgut, mimicking the most common clinical finding in HSCR patients[26, 27]. For this reason, mutation of *EdnrB* in mice, both in the whole animal (*EdnrB*^{-/-})[28–34] and specifically in neural crest cells only (*EdnrB*^{NCC-/-})[35, 36], has been used to study the pathogenesis of HAEC, with 9 publications using these models over the last five years.

Mucosal Immunity

The gut-associated lymphoid tissue (GALT) is the largest lymphoid organ in the body and is charged with providing protection against a variety of antigens that may gain access to the host, including food particles, commensal and pathogenic bacteria and their toxins[37]. The GALT can be organized into inductive and effector sites for the ease of discussion and investigation. Peyer's patches (PP) are the primary inductive site for gut mucosal immunity[38]. PP are distinct collections of immune cells that display follicular architecture similar to lymph nodes and are located along the anti-mesenteric surface of the bowel. Circulating naïve T-lymphocytes and B-lymphocytes, which display $\alpha 4\beta 7$ integrin and L-selectin, migrate from circulation to the PP via binding of MADCAM-1 (mucosal addressin cell adhesion molecule 1), which is expressed in the high endothelial venules associated with PP[39, 40]. The mucosa overlying PP contains specialized epithelial M cells that sample and transport antigen from the lumen of the bowel to the underlying PP, where it is presented to naïve lymphocytes by antigen presenting cells, such as dendritic cells (DC). Activated lymphocytes then enter efferent lymphatic channels to travel to mesenteric lymph nodes (MLN). From the MLN, activated lymphocytes travel through the thoracic duct into the systemic circulation. Homing of activated lymphocytes to the lamina propria (LP), the primary effector site of mucosal immunity, occurs via the coordinated action of a number of cellular adhesion molecules, cytokines and chemokines[41]. Once in the LP, T-lymphocytes produce the pro-inflammatory Th-2 cytokines interleukin-4 (IL-4) and IL-10 to stimulate immunoglobulin A (IgA) production by terminally differentiated B-lymphocytes, termed plasma cells. IgA, the most abundantly produced immunoglobulin isotype in the body, is then actively transported to the mucosal surface in dimeric form by polymeric immunoglobulin receptor (pIgR)[42]. During transport, five of the seven pIgR domains, termed the secretory component (SC), remain attached to the IgA dimer. The secreted complex, referred to as secretory IgA (SIgA), is the principle effector of antigen-specific immune defense along the mucosal surface.

There is abundant evidence in the literature to suggest that multiple aspects of the mucosal immune system detailed above are altered in the setting of HSCR/HAEC. Cheng, etl.al., using the *EdnrB*^{-/-} mouse, developed a histopathology grading system for quantitative assessment of HAEC[28]. In this system, severity and depth of inflammation are graded separately, and the score summed to arrive at the final number. Similar histologic changes were observed in *EdnrB*^{NCC-/-} mice during the development of HAEC[35, 36], with mortality in the fourth week of life. Additionally, there was a linear correlation between histology score and bacterial colonization of intraperitoneal organs[28].

A role for IgA in HAEC pathogenesis was demonstrated by Imamura, et.al, who found an increase in IgA containing plasma cells along the length of resected aganglionic bowel from patients with HAEC as compared to HSCR patients without HAEC[43]. They also found decreased luminal IgA (SIgA) in the same patients, suggesting decreased production or impaired transport to the lumen. Recently, a similar observation, decreased luminal SIgA, has been made in *EdnrB^{NCC-/-}* mice[44]. Interestingly, this finding was specific to the gut, with normal levels of bronchial and nasal IgA seen in these animals. In the mouse, a specific subset of B-lymphocytes (IgM⁺IgD^{high}), termed mature B-lymphocytes, from the PP are responsible for the majority of intestinal IgA production[45]. Unlike the *Ret^{-/-}* animals, which lack PP, the *EdnrB^{NCC-/-}* animals display a normal number of PP, but demonstrate decreased B220⁺IgM⁺IgD^{high} (mature) B-lymphocytes in PP[44]. In the mouse, mature B-lymphocytes in PP are primarily of splenic origin[45]. Small splenic size and splenic lymphopenia was first noted in the *EdnrB^{-/-}* animal[30]. They noted a relative reduction of B-versus T-lymphocytes as well as a negative correlation between splenic lymphocyte counts and intestinal inflammation on histologic analysis. These findings were extended in the *EdnrB^{NCC-/-}*, in which B-cell lymphopenia was also observed[36]. Furthermore, they observed a decrease in marginal zone B-lymphocytes, suggesting impaired B-lymphocyte development or trafficking. Both groups have noted that similar lymphocyte population alterations may be observed in the setting of stress (e.g. HAEC). In an attempt to delineate the contribution of the *EdnrB^{-/-}* genotype to the HAEC phenotype, Frykman, et.al., performed bone marrow transplants from *EdnrB^{-/-}* animals to *Rag2^{-/-}* recipients and, separately, induced bowel obstruction in WT animals[33]. They concluded that stress from obstruction resulted in lymphocyte alterations similar to those seen in the HAEC models. However, they have additionally shown that, in the *EdnrB^{-/-}* model, animals that undergo surgical relief of obstruction continue to carry a 40% risk of developing HAEC[46]. Additionally, they noted small splenic size in the *EdnrB^{-/-}* bone marrow transplant recipients, further underscoring a role for *EdnrB* in lymphocyte development and/or trafficking.

Another critical facet of mucosal immunity is barrier defense. In the steady state, bacterial invasion across the colonic epithelium is prevented mucus, composed of glycosylated proteins of the MUC family and secreted by goblet cells[47]. This prevents the adherence of pathologic organisms to enterocytes, reducing susceptibility to infection. Multiple studies have demonstrated impaired intestinal barrier function in HAEC patients, specifically decreased secretion of mucin and increased adherence of bacteria to enterocytes[48–50]. Recently, Thiagarajah, etl.al., used human HSCR colonic biopsies and the *EdnrB^{-/-}* mouse to investigate goblet cell alterations in the setting of HSCR and HAEC[32]. They found increased goblet cell numbers in HSCR patients and distal *EdnrB^{-/-}* colons, along with up-regulation of goblet cell differentiation genes. Finally, they noted decreased amounts of membrane-bound Mucin 4 (Muc4), but not secreted Muc2, in the distal colons of *EdnrB^{-/-}* animals. Further, functional studies have been performed in the same animal model[34]. In these studies, reduced rates of transport of microparticles and bacteria, both active and passive, were observed in the aganglionic and ganglionated bowel. Of note, increased goblet cell size was also noted in the *EdnrB^{NCC-/-}* model, further suggesting alterations in goblet cell structure and function in the setting of HAEC[35]. In this study, the authors focused on

the proximal, ganglionated bowel, and investigated Paneth cells, which are the primary source of antimicrobial compounds for epithelial protection[51]. They found reduced Paneth cell sPLA₂ expression in ileal tissue and decreased sPLA₂ activity in the lumen, along with a blunted sPLA₂ response to bacterial challenge[35].

Together, these studies have delineated a series of mucosal immune defects in HSCR/HAEC that involve both the aganglionic, as well as ganglionated bowel, and extend beyond the GI tract. Further work to determine the mechanistic underpinnings of these defects, and potential therapeutic targets, is warranted.

Microbiome

Over the last decade, many lines of investigation have revealed the influence of the microbiome on human health and disease[52]. These studies have demonstrated both spatial and temporal variability of the microbiome within and amongst individuals, along with demonstrating dysbioses that may contribute directly to disease pathogenesis. Historically, investigations into an infectious cause for HAEC have failed to identify a single, responsible organism, although viral and bacterial etiologies have been suggested. However, recent studies in mouse models of HSCR/HAEC have identified broader changes in the intestinal microbiome that may contribute to HAEC. These advances have been made possible by next-generation sequencing technologies, which are culture-independent.

Two studies have used mouse models to examine the role of the microbiome in HAEC. Using the *EdnrB*^{-/-} model, Ward, et.al., demonstrated increasing diversity of the microbiome over time (from post-natal day 7–24), with a greater increase in the *EdnrB*^{-/-} versus control animals[31]. Using principal coordinate analyses, which allow visualization of how the microbiome “clusters”, the HSCR animals clearly demonstrated distinct microbiota from the controls at all ages. When specific bacterial species were examined, the HSCR animals had higher levels of *Bacteroidetes* and *Firmicutes* than the control animals. They also noted decreased *Lactobacillus* species over time in both HSCR and control animals. In another study, using the *EdnrB*^{-/-} model, survival could be extended to 36 days by changing to a liquid diet and the addition of oral antibiotics[28], further supporting a role for the microbiome in the development of HAEC. Another group has employed the *EdnrB*^{NCC-/-} model in a similar fashion[35]. They noted similar compositions of the microbiota between HSCR animals and controls at an early time point, with a shift towards “dysbiosis” occurring prior to the onset of HAEC. Additionally, unlike the findings in the *EdnrB*^{-/-} model, they noted a decrease in *Lactobacillus* in HSCR animals preceding HAEC. They also observed increases in *Bacteroidetes* and *Clostridium* species, and noted that only HSCR animals demonstrated *Escherichia coli* in their microbiome samples. Finally, unlike models of inflammatory bowel disease, they noted increased species diversity at the time of enterocolitis, suggesting distinct mechanisms of intestinal inflammation between HAEC and IBD.

Culture-independent techniques to study the microbiome have recently been applied to human HSCR/HAEC patients as well. DeFilippo, et.al., followed a single child with HSCR over a five month time period, obtaining 15 fecal samples[53]. While the majority of clustering they observed was explained by the passage of time, they did note progression of

microbiota changes from “pre-enterocolitis”, through acute HAEC, and on to remission phases. Yan, et.al., applied a similar approach to four patients with HSCR, two with HAEC and two without[54]. They found that the patients with HAEC clustered separately from those without HAEC. Additionally, similar to the *EdnrB^{NCC-/-}* finding[35], they noted increased species diversity in the HAEC patients. Finally, another group has examined both the bacterial and fungal microbiome in a group of 19 HSCR patients, half with a history of HAEC[55]. They noted HAEC patients had increased fungal diversity, with increased *Candida species* and reduced *Malassezia* and *Saccharomyces species*, suggesting the microbiota changes contributing to HAEC may extend beyond bacteria.

Genetics

From the accumulating basic science evidence, it is tempting to postulate that a specific genotypic mutation confers a higher risk for development of HAEC.

Nucleotide oligomerisation domain 2 (NOD2), which is a pattern recognition receptor, has been associated with Crohn’s disease and graft versus host disease[56]. In the intestine, NOD2 is involved in post-natal PP development and its deficiency results in overgrowth of lymphoid tissues in the gut[57]. However, a study of human patients with HSCR showed that while 19.2% of patients were carriers of NOD2 mutations, none of the patients that developed HAEC carried NOD2 variants[58].

Integrin beta-2 (ITGB2) is involved in cell-surface mediated signaling and is required for leukocyte extravasation from the blood into tissues. ITGB2 has been associated with chronic colitis conditions, and participates in T-cell development. In a study of blood and colonic tissue samples from 33 patients with HSCR, 66% of patients demonstrated ITGB2 variants[59]. Of these, 13 patients had experience episodes of HAEC, suggesting a molecular target for further investigation.

In an effort to further our understanding of how genotype is linked to phenotype, the Hirschsprung Disease Research Collaborative has gathered geneticists, pediatric surgeons, pediatricians, pathologists, and gastroenterologists together. This group is enrolling children and families worldwide in order to perform a global genetic analysis with deeply phenotyped clinical information. The goal of this research is to provide improved genetic counseling for patients and families and to discover possibilities for improved and individualized treatment.

Treatment

Based on the findings in basic science and pilot human studies, a few recent efforts have been made to advance our approach to HAEC treatment. El-Sawaf, et.al., randomized children undergoing surgery for HSCR to receive either probiotic or placebo postoperatively[60]. While they did not observe a difference in the incidence of HAEC (28.3% overall) or recurrent HAEC (19–21%), this may be a limitation of the choice of probiotic formulation, rather than the overall treatment strategy. In contrast, Wang, et.al., similarly randomized patients to receive probiotic for a duration of 4 weeks[61]. Over the subsequent three months, the probiotic group demonstrated decreased incidence and severity of HAEC, along with decreased circulating pro-inflammatory cytokines. These studies

suggest that strategies aimed at altering susceptibility factors for HAEC may hold therapeutic potential.

Conclusions

Since its original description by Harald Hirschsprung 130 years ago, HAEC remains a common complication of HSCR, and carries significant morbidity and potential mortality. Recent advances in our mechanistic understanding of HAEC have focused on mucosal immunity, the role of the intestinal microbiome, and the role of genetics in determining disease phenotype (Figure 2). Initial efforts have been made to translate these discoveries into therapeutic measures, with mixed success. Moving forward, further basic science efforts combined with collaborative translational efforts will be required to advance our understanding and treatment of HAEC.

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PubMed HAEC Publications Over Time

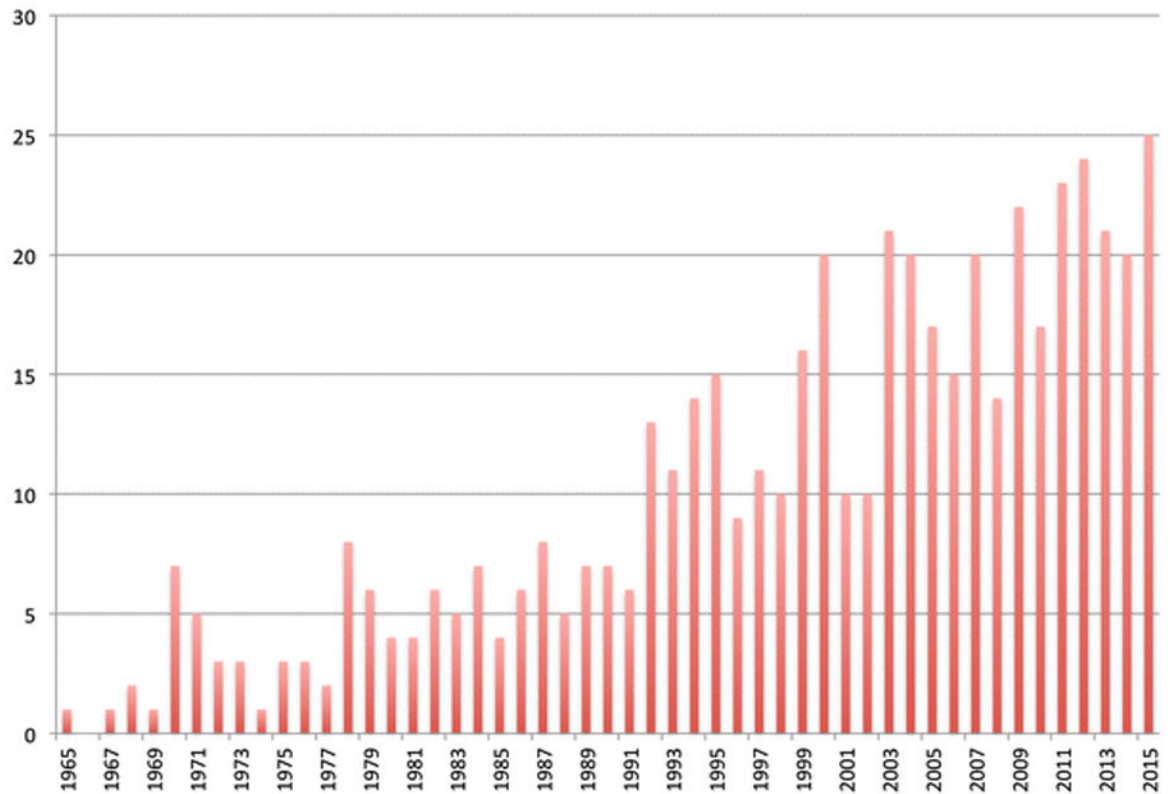


Figure 1.

Number of HAEC Publications in PubMed over time. A PubMed search for ((Hirschsprung[Title/Abstract] OR Hirschsprung's[Title/Abstract]) AND Enterocolitis) was conducted on 31 December 2015 to determine the number of publications related to HAEC over time. The number of publications related to HAEC is increasing, with 25 in 2015.

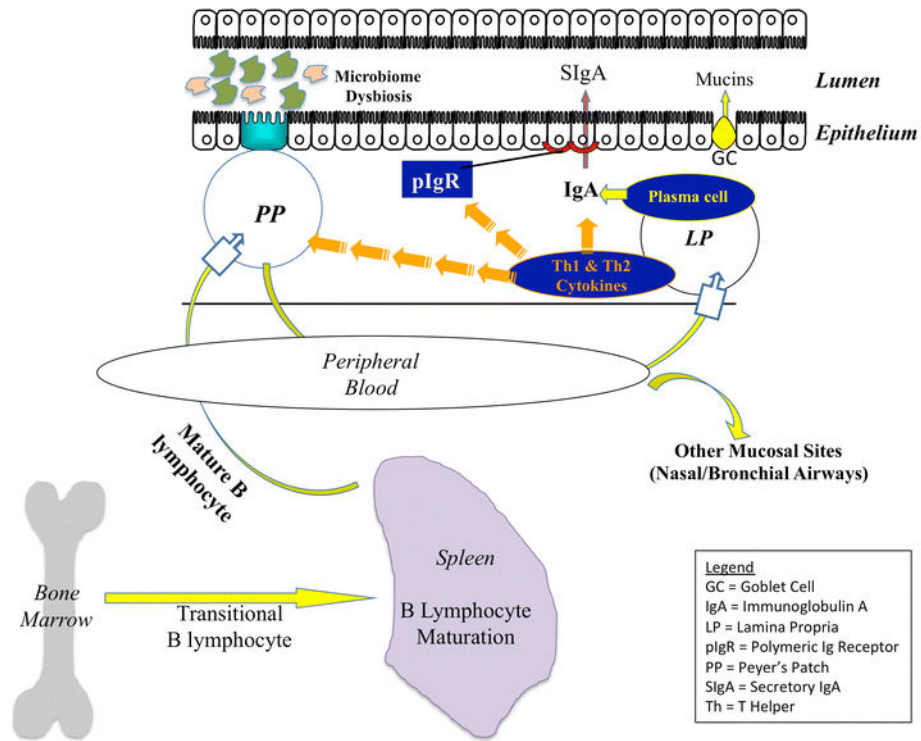


Figure 2. Emerging Concepts in HAEC Pathogenesis. Recent studies in the *EdnrB*^{-/-} and *EdnrB*^{NCC-/-} mouse, along with human studies, have implicated facets of mucosal immunity (encompassing innate and adaptive immunity, as well as epithelial barrier defense mechanisms) and microbiome dysbiosis in the development of HAEC.