REVIEW ARTICLE -

The Developing Intestinal Ecosystem: Implications for the Neonate

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ABSTRACT

Interactions of resident intestinal microbes with the luminal contents and the mucosal surface play important roles in normal intestinal development, nutrition, and innate and adaptive immunity. The neonate, especially the premature, who possesses a highly immunoreactive intestinal submucosa underlying a single layer of epithelial cells that are continuously exposed to the luminal environment, is highly susceptible to perturbations of the luminal environment. Understanding the interactions of the intestinal ecosystem with the host and luminal nutritional environment, especially in regard to human milk and pre- and probiotics,

A better understanding of the developing intestinal ecosystem in the neonate may hold the key to prevention of several important diseases. This intestinal ecosystem is composed of three closely interacting components: host cells, nutrients, and microflora. Changes in intestinal microflora and host cells during maturation affect the developing intestine, and these interact with the changing nutrient environment. Reduction of normal commensal bacteria in the context of infection or after antibiotic treatment may interfere with availability of critical nutrients and impair beneficial stimulation of GI mucosal development and the innate and adaptive immune responses. Here we will briefly summarize some of our current knowledge about the intestinal microflora, its interaction with the maturing intestine, and the nutritional environment ("cross-talk"). We also discuss consequences of disrupted interaction between these components, especially pertaining to the intestinal mucosal inflammatory and immune capability, and why human milk is the appropriate substrate in this environment.

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has major implications for the pathogenesis of diseases that affect not only the intestine but distal organs such as the lung and brain. (*Pediatr Res* 58: 625–628, 2005)

Abbreviations

GI, gastrointestinal LPS, lipopolysaccharide NEC, necrotizing enterocolitis TLR, toll-like receptor

NORMAL MICROFLORA

The microflora of the adult human is found primarily in the colon and distal small intestine and consists of more than 10¹³ microorganisms (the "microbiota"), comprising nearly 500 species and nearly 2 million genes (the "microbiome") (1). This is mostly a mutually beneficial relationship, as evidenced by the important role commensal bacteria play in nutrition (2), angiogenesis (3), and mucosal immunity (4,5). The role of bacteria in providing nutrients to the host is exemplified by the fermentative salvage of lactose that remains unabsorbed in the distal intestine because of low lactase activities in the immature intestine of premature infants (6). The resulting short-chain fatty acids such as acetate, propionate, and butyrate are used for energy or synthetic processes (6), stimulate intestinal blood flow, affect intestinal proliferation and differentiation, and alter intestinal interepithelial tight junctions (7). Other beneficial effects of commensal bacteria include lipid hydrolysis, protein breakdown into peptides and amino acids, and vitamin production.

DEVELOPMENT OF THE INTESTINAL MICROFLORA

Marked changes occur in the intestinal microflora from birth to adulthood. At birth, the intestine is sterile but becomes rapidly colonized thereafter (8). Neonatal human intestine contains large numbers of facultative anaerobes such as *Streptococcus* and coliforms. These decline in number during weaning as obligate anaerobes such as *Bacteroides* and *Clostridium* establish a foothold and eventually become the predominant community residing in the gut. These changes are accompanied by transition from a high-fat, milk-based diet to a diet rich in carbohydrates and maternal immunoglobulins (8). In the preterm infant, the methods of neonatal intensive care, such as treatment with antibiotics, total parenteral nutrition, or nursing in incubators, may delay or impair the intestinal colonization process. They are thus slower to acquire commensals such as bifidobacteria and more susceptible to pathogenic colonization (9). The intestine of the premature infant also has poor motility and is predisposed to bacterial overgrowth (10).

Profound changes occur in the intestinal ecosystem when young mammals are weaned from their mother's milk. An example of such a change that normally occurs is in the angiogenins, potent antimicrobial peptides released from the Paneth cells of the crypt (11). In mice raised in a conventional environment, angiogenin mRNA expression is markedly increased at the time of weaning from mother's milk to an adult diet. This change occurs to a much lesser degree in germ-free mice (11,12). It is also thought that commensal bacteria, through "cross-talk" and the production of angiogenins, are critical for normal villus capillary development. One dramatic example of this is seen in the lag in development of the villus capillaries in the mouse small intestine grown in a germ-free environment versus normal growth in an environment containing normal mixed flora or even only one species of commensal bacterium (Bacteroides thetaiotaomicron) (3).

ROLE OF MICROFLORA IN INFLAMMATION

The commensal microflora may represent a key regulatory checkpoint for the intestinal inflammatory response. The intestinal epithelium partially relies on TLR to act as an interface between the luminal microflora and signal transduction pathways. TLR are cell-surface receptors that recognize specific microbial ligands, from both pathogens and commensals, enabling the innate immune system to recognize nonself, and activating both innate and adaptive immune responses (13). Studies of the interaction between resident microflora and TLR are beginning to shed light on how the healthy intestinal surface defuses the threat of commensal bacteria to the lumen, and how this interaction is actually required to maintain the architectural integrity of the epithelium (14,15). These studies suggest that the epithelium and resident immune cells do not simply tolerate commensal microorganisms but are dependent on them. Commensal bacteria secrete molecules such as LPS and lipoteichoic acid, which in turn interact in the normal intestine with a population of surface TLR. The resultant ongoing signaling enhances the ability of the epithelial surface to withstand injury while also priming the surface for enhanced repair responses. Therefore, either the disruption of TLR signaling or the removal of TLR ligands compromises the ability of the intestinal surface to protect and repair itself in the face of inflammatory or infectious insult (5). Another study reported nonvirulent *Salmonella* strains whose direct interaction with model human epithelia attenuate IL-8 production elicited by various proinflammatory stimuli (16). This interaction abrogates the polyubiquitin-mediated degradation of the inhibitor of nuclear factor kappa-B (alpha subunit; $I\kappa$ -B α) (16). These studies raise intriguing questions: Is the TLR signaling pathway fully functional in the premature infant? Does the common practice of treating premature infants with broad-spectrum antibiotics shortly after birth adversely affect the commensal population? Does the lack of enteral nutrition or the provision of commercial formula rather than mother's milk result in undergrowth of commensal bacteria?

The microbial population of the developing GI tract of the premature infant affects maturation and optimal function of the intestinal innate and adaptive immune system, suggesting a pathogenic role in several diseases related to systemic inflammation such as NEC, chronic lung diseases, intraventricular hemorrhage, periventricular leukomalacia, and hematopoietic abnormalities. Over the past few years, a body of literature has emerged supporting that overproduction of IL-8 is a critical, and arguably the most important, component of the systemic inflammatory response in neonates. The role of IL-8 in chorioamnionitis, intracerebral hemorrhage, and lung disease is compelling (17-21). High concentrations of IL-8 are frequently noted in the plasma and affected tissues of patients who exhibit these problems. This leukocyte chemoattractant chemokine, which can be produced in large quantities by the intestine, has thus been causally linked to some of these problems, implying that modulation of intestinal production of IL-8 may be a key to prevention. It is notable that infant intestine appears to be much more sensitive to stimuli, such as LPS and flagellin that induce IL-8, than is adult intestine (22,23). Figure 1 represents a hypothetical scheme as to how commensal microorganisms or probiotic bacteria might interact with the intestine to prevent these complications.

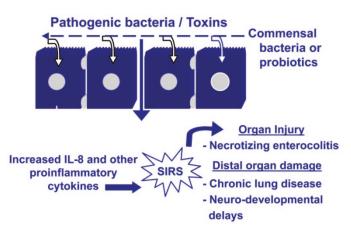


Figure 1. Hypothetical scheme demonstrating the effect of commensal microorganisms on the intestinal production of proinflammatory mediators such as IL-8 and the subsequent prevention of propagation of this inflammatory mediator to proximal (intestine) as well as distal organs (lung and brain).

ENTERAL NUTRIENTS, HUMAN MILK, AND THE INTESTINAL MICROFLORA

Nutrients, as well as probiotic bacteria and/or prebiotics (nutritional substrates that promote the growth of probiotic bacteria), through their stimulation of a normal commensal microflora may also play an important role in the regulation of IL-8 (24–27) and neutrophil infiltration into the intestine. The reliance on parenteral nutrition while providing few or no enteral nutrients may be highly significant in the promotion of intestinal inflammation because the presence of enteral nutrients can prevent gut-derived inflammation (28). Whether this is partially due to stimulation of commensal bacterial growth is speculative. Several in vitro studies have shown that probiotics (live, heat-killed, and DNA from probiotic bacteria) can downregulate intestinal IL-8 production (26,27,29-31) induced by proinflammatory stimuli such as LPS, but studies in neonatal animals (in vivo) showing regulation of intestinal inflammation by pre- or probiotics have not yet been published.

The feeding of human milk is an immediate way to promote the development of the intestinal ecosystem in the neonate, including the preterm. The beneficial effects generally relate to improvements in host defenses, digestion and absorption of nutrients, GI function, neurodevelopmental outcomes, and maternal psychological well-being and also may contribute to a reduction in the economic burden brought on by the long hospitalization of the premature infant. However, the special needs of the preterm infant that arise as a result of metabolic and GI immaturity, immunologic compromise, and associated medical conditions must be considered so that adequate nutrition can be provided to meet the needs for intrauterine rates of growth and nutrient accretion (32).

Clinical studies in nurseries throughout the world indicate that premature infants have fewer episodes of late-onset sepsis, NEC, diarrhea, and urinary tract infection, and need less antibiotic therapy when fed their own mothers' milk compared with similar infants fed formula (33–35). Although the greater the dose of human milk, the greater the protective effects, reductions in the incidence of late-onset sepsis and/or NEC occur in infants who only receive partial mother's milk feedings (>50 mL/kg) compared with similar infants fed either less mother's milk or preterm formula (36,37). A recent report indicated that 30% of mothers of extremely premature infants did not secrete the anti-inflammatory cytokine IL-10 in their milk, yet this cytokine was not found in 85% of the mothers whose milk was fed to similar infants who developed NEC (38). The presence of high concentrations of soluble CD14 (sCD14), a pattern recognition receptor molecule not found on the intestinal surface, in human milk is also of interest in this regard. sCD14 prevents inflammatory conditions of the gut and has been shown to induce B-cell growth and differentiation (39). This is especially intriguing because of previously reported studies that show that breast-feeding combined with providing probiotics to mothers and their infants is associated with a decreased development of atopic diseases as these children get older (40).

Human milk contains a wide array of biologically active components. The growth of pathogenic bacteria and viruses is

known to be inhibited by proteins like lactoferrin, secretory IgA, and peptide formed from human milk during digestion (41). Human milk is an important factor in the initiation, development, and composition of the neonatal gut microbiota. It has been found to be a significant source of lactic acid bacteria that appear to be of endogenous origin and not contaminants from the breast skin (42). Some of these bacteria belong to the species Lactobacillus, the probiotic potential of which has recently been evaluated (43). The lactobacilli isolated from milk of healthy mothers is at least similar to that of the strains commonly used in commercial probiotic products in terms of survival to GI tract conditions, production of antimicrobial compounds, adherence to intestinal cells, production of biogenic amines, and patterns of antibiotic resistance. Among the numerous substances present in human milk, oligosaccharides have a clear prebiotic effect, stimulating the development of bifidogenic flora in the colon (44). Breast-fed infants, unlike those who are formula- fed, have an intestinal ecosystem characterized by a strong prevalence of bifidobacteria and lactobacilli (8,9). Human milk can be considered a symbiotic (44), a substance that has the properties of both a pro- and prebiotic. Other anti-inflammatory components in human milk include antioxidants, epithelial growth factors, cellular protective agents, and enzymes that degrade mediators of inflammation (45). Although the above-mentioned microbes found in milk may be partially responsible in reducing infection and inflammation, the lower n-6/n-3 fatty acid ratio, through a higher relative concentration of the anti-inflammatory n-3 compared with proinflammatory n-6 fatty acids, may also play a role (46). Lastly, there is some indication that host-defense benefits persist even beyond hospitalization in former premature infants. Fewer upper respiratory signs and symptoms were reported when human milk was continued after discharge, even if the diet was only partial human milk, and the protective effects continued even beyond the time human milk feeding was discontinued (47).

The relationship between human milk and reduced incidence of sepsis and/or NEC reported in the above studies may have important economic implications. For example, a reduction in the incidence of NEC may have profound effects on the cost of health care. It has been reported that one case of medically treated NEC increases the cost of hospitalization by \$73,700 (48). Given conservative estimates of the incidence of NEC, a 10% versus 4% incidence of NEC in formula- versus human milk–fed premature infants means that 5600 versus 2240 infants might be affected in the United States (2002 statistics), with a cost differential of \$4.1 million versus \$1.6 million (49).

CONCLUSIONS

Establishment and maintenance of a "normal" intestinal microflora may help abrogate several neonatal inflammatory conditions that not only affect the GI tract but are initiated in the gut and cause damage to distal organs such as the lung and brain. Decreasing excessive use of antibiotics and increasing the use of pro- and prebiotics or symbiotic nutrients such as human milk that promote the establishment and growth of a beneficial microflora are means by which neonatologists can maintain their patients' intestinal ecosystem.

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