The influence of nutrition on the development of the child's intestinal microflora

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Abstract: The intestinal microbiota, also known as the intestinal microflora, or the intestinal flora, is a group of microorganisms (microbiome), mainly bacteria, which form a complex ecosystem in the digestive system. In humans, it is one of the elements of his physiological biota [1].

Childbirth is an important stage in the colonization of the human digestive system by the microbiota. The form of feeding also plays an important role in the colonization of the intestines of a newborn and infant [2]. Breast milk has many benefits. Breastfeeding plays an important role in shaping baby's gut microbiota. Mother's milk contains health-promoting bacteria that support the primary colonization of the newborn's intestines. Diet has a significant influence on the formation of the intestinal microbiota. Therefore, it is important that a woman not only during the entire pregnancy, but also before pregnancy, takes care of a varied, balanced and rich in prebiotics diet [3].

Keywords: microbiota of a pregnant woman, diet, breastfeeding, childbirth.
1. Introduction

The gut microbiota is a group of microorganisms (microbiome), mainly bacteria, that form a complex ecosystem in the digestive system [4]. The intestinal biota bacteria are found mainly in the large intestine and are the basic mass of the stool [5]. The intestines contain 1,000 species of bacteria and 100 times more genes than the human genome [6]. The intestinal microbiota also includes fungi and protozoa [7].

In newborns, the digestive system is sterile, but it is later colonized by bacterial biota [8]. With aging, the total number of microorganisms does not change significantly, but there is usually a significant change in the share of particular groups. You can observe an increase in the number of enterobacteria, Clostridium and enterococci and a simultaneous decrease in the number of Lactobacillus and Bifidobacterium, which are considered beneficial to health [9].

The composition of the gut microbiota changes under the influence of a number of different factors [11], e.g.

1. food
2. hormones
3. the environment,
4. the type of delivery - whether by natural means or by caesarean section,
5. taking antibiotics and medications,
6. century
7. stress
8. diseases.

In recent years, there has been a significant increase in interest in the role of the intestinal microbiota in shaping human health and in the occurrence of intestinal dysbiosis in various disease entities, including Parkinson's disease, breast cancer, acute myeloid leukemia [12–14]. The intestinal microbiota is not only involved in the processes related to digestion and absorption of nutrients [15]. The bacteria that are part of the intestinal microbiota perform various important functions, and their correct quantitative and qualitative structure, the so-called eubiosis state, supports the homeostasis of the whole organism, thus shaping immunity, metabolism and the
synthesis of many chemical compounds, for example serotonin and neurotransmitter 
precursors [16–18].

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intestinal flora, is a group of microorganisms (microbiome), mainly bacteria, which 
form a complex ecosystem in the digestive system. In humans, it is one of the elements 
of his physiological biota [19,20].

The relationship between the host and the bacterial biota on commensalism is a 
kind of favorable symbiosis [21]. Microorganisms perform many useful functions, such 
as:

1. fermentation of certain nutrients,
2. stimulation of the immune system in the fight against pathogenic 
   microorganisms,
3. regulation of intestinal development,
4. production of vitamins (biotin and vitamin K),
5. hormone production.

Under certain conditions, certain species of the gut microbiota can cause disease 
states (opportunistic infections) or contribute to carcinogenesis. The gastrointestinal 
microbiome, which is regulated by the composition of the diet and nutritional status, 
also significantly affects the maintenance of the proper functioning of the brain [17,22].

Scientific research shows that the gut bacteria of a pregnant woman, her diet and 
health status have a significant and direct impact on the future profile of the baby's gut 
bacteria [3,23]. During pregnancy, the female body undergoes hormonal, metabolic, and 
immunological changes to preserve the health of both the mother and the offspring [24]. 
These changes alter the mother microbiota at different sites such as the gut, the vagina, 
and the oral cavity [25]. A number of factors might influence the microbiota profile 
such as the diet, antibiotic, or other supplement intakes, as well as the methodology of 
research [4,24,26].
2. Materials and Methods

Systematic searches were conducted in August 2022 using electronic databases such as Pubmed, Science Direct and Google Scholar in accordance with PRISMA guidelines (Moher, Liberati, Tetzlaff, Altman and Group, 2009). The databases were checked by two independent authors. The following items were searched: (microbiota or microbiome) AND pregnancy; intestinal microflora and child development; microbiota and child nutrition. The articles you downloaded were first selected based on title and abstract. As a result, we identified a total of 235 articles related to the topic of interest to us. After considering the inclusion / exclusion criteria and eliminating duplicates, 32 studies were selected for analysis.

Inclusion criteria for selecting the study:

• the samples have been taken from the stool or placenta,
• randomized clinical trials, systemic reviews, meta-analyzes,
• Human Research.

Exclusion criteria:

• case reports, conference summaries, comments,
• insufficient amount of data,
• full-text article not available for review,
• a language other than English,
• non-human studies.

Key review references were hand searched to identify any relevant references that were omitted.

2.1. Feeding

The form of feeding also plays an important role in the colonization of the intestines of a newborn and infant [8]. Mother's milk is recommended for most newborns. Breastfeeding plays an important role in shaping your baby's gut microbiota [27]. Mother's milk contains health-promoting bacteria that support the primary
colonization of the newborn's intestines. Bifidobacterium, Lactobacillus and representatives of the Enterobacteriaceae family, as well as substances stimulating their growth (prebiotics), including the most important human milk oligosaccharides (HMO), enter the digestive tract of the newborn along with the mother's food. The mixture of protective bacteria, prebiotics with immunoactive proteins in breast milk is responsible for [28]:

1. proper digestion, development of intestinal villi,
2. intestinal peristalsis,
3. shaping the immune system of a newborn.

It has been shown that in breastfed infants, the protective bacteria Bifidobacterium with an admixture of Lactobacillus dominate, while bacteria from the genera Clostridium and Escherichia are much smaller [29–31]. Bacterial diversity grows into an adult microbiota-like syndrome in the first two years of a child's life as a result of exposure to new foods and the microbiota of the child's environment, including siblings, pets and immediate surroundings [32,33]. It should be remembered that an excessive increase in the hygienization of life, resulting in a reduced contact of children with microbes of human and animal origin, and the abuse of antibiotics during pregnancy and the first two years of life, may significantly disturb the development of microbiota [34,35].

2.2. Diet

Diet has a significant influence on the formation of the intestinal microbiota [36]. Therefore, it is important that a woman not only during the entire pregnancy, but also before pregnancy, takes care of a varied, balanced and rich in prebiotics diet [37]. As a result, immune processes are improperly shaped, which can lead to an excessive immune reaction to pollen, dust mites (allergies) or own tissues (autoimmune diseases) [38,39,39]. Additionally, it should be borne in mind that the addition of antibiotics perinatal to the mother or baby and preterm labor strongly influences the microbiota and may have significant consequences for later health [40,41]. Preterm newborns have, inter alia, reduced size of the Bacteroidaceae family in the first months of life and a higher initial percentage of Lactobacillaceae, compared to full-term infants [42].
Antibiotics that are taken perinatally also significantly affect the quantitative and qualitative structure of the intestinal microbiota [43,44].

The intestinal microbiota forms a kind of ecosystem that is comprehensive and metabolically active. However, he is also very sensitive [45,46]. The factors most strongly disrupting the microbiota include e.g.:

1. incorrect diet low in fiber,
2. antibiotics,
3. proton pump inhibitors,
4. non-steroidal anti-inflammatory drugs,
5. some stress,
6. alcohol,
7. environmental and food pollution.

Studies also show a relationship between intestinal microbiota disorders and civilization diseases, such as type 2 diabetes, obesity and hypertension, the risk of cardiovascular disorders, autoimmune diseases and inflammatory bowel diseases (IBD) [34,47–49].

A very important test is the assessment of the intestinal microbiota, which includes detailed microbiological cultures and genetic analyzes for non-breeding bacteria [50]. Such studies include a detailed qualitative and quantitative analysis of indigenous microorganisms:

1. protective (anaerobic bacteria of the genus Bacteroides and Bifidobacterium, lactobacilli of the genus Lactobacillus)
2. immunostimulatory bacteria (Enterococcus and E. coli), nourishing the intestinal epithelium (Faecalibacterium prausnitzii and Akkermansia muciniphila),
3. proteolytic bacteria (Clostridium, Enterobacteriaceae family including Klebsiella spp. Enterobacter spp., Citrobacter spp., Proteus spp.) And Pseudomonas and the number of bacteria [51,52].

Non-invasive examination of feces allows for the assessment of the quantitative composition of the intestinal microbiota and enables the selection of the appropriate and most effective, targeted and individualized prebiotic therapy and probiotic therapy [53]. An individually selected diet and probiotics significantly support the intestinal
microbiota. It is also very important to properly select safe, tested preparations. Bacteria should reach the large intestine and stimulate the creation of favorable conditions for the development of normal microbiota [21,54,55].

Helen L. Barrett et al. conducted research that a vegetarian diet is the main determinant of the intestine. They also analyzed the composition of the microbiota in early pregnancy. This study explored the gut microbiota profile in women who were vegetarian or omnivorous in early gestation. Women were selected from participants in the Study of Probiotics in Gestational diabetes (SPRING) randomised controlled trial. Nine women identified as vegetarians were matched to omnivorous women in a 1:2 ratio. Microbiota analyses were performed using 16S rRNA gene amplicon sequencing and analysed using the Quantitative Insights Into Microbial Ecology (QIIME) and Calypso software tools. There was no difference in alpha diversity, but beta diversity was slightly reduced in vegetarians. There were differences seen in the relative abundance of several genera in those on a vegetarian diet, specifically a reduction in Collinsella, Holdemania, and increases in the relative abundances of Roseburia and Lachnospiraceae. In this sub-analysis of gut microbiota from women in early pregnancy, a vegetarian as compared to omnivorous diet, was associated with a different gut microbiome, with features suggesting alterations in fermentation end products from a mixed acid fermentation towards more acetate/butyrate [56]. This study explored the gut microbiota profile in women who were vegetarian or omnivorous during early gestation. There was no difference in alpha diversity, but beta diversity was reduced in vegetarians. There were differences seen in the relative abundance of several genera in those on a vegetarian diet, specifically a reduction in Collinsella, Holdemania, and an increase in Roseburia and Lachnospiraceae. Functional analyses suggested that women on a vegetarian diet had higher abundance of species involved in fatty acid and lipid synthesis. Collinsella is positively correlated with insulin and lipid levels in the SPRING cohort as well as outside pregnancy. In non-pregnant omnivorous people, Collinsella aerofaciens was also reported to be higher than in their vegetarian.

4. Discussion

Childbirth is an important stage in the colonization of the human digestive system by the microbiota. Natural childbirth is definitely the most beneficial for the
microbiota, during which the fetus, passing through the genital tract, is populated with the physiological microbiota of the mother's vagina. Numerous studies have shown that in the case of cesarean section, the newborn is colonized by skin bacteria from the mother and hospital staff, and in extreme cases by hospital-derived strains [10,53,57].

Research carried out by Dominguez-Bello found that the microbiota of newborns born physiologically is similar to the microbiota of the vagina with a high proportion of Lactobacillus bacteria, in contrast to those born by caesarean section, whose microbiota contained more skin bacteria, e.g. Streptococcus, Staphylococcus [58]. Unfortunately, the number of imperial burdens is now significantly increasing. Research clearly emphasizes the correlations between cesarean sections and intestinal dysbiosis, which increases the real risk of allergies, atopy, bronchial asthma, type 1 diabetes and obesity later in life [59]. The first 1000 days of a child's life, along with fetal life, programs the microbiota, having a key and long-lasting impact on human development and health [60,61].

5. Summary

In summary, proper nutrition has a major impact on health and well-being during pregnancy. The protection of the intestinal bacterial flora is important during pregnancy. A balanced diet very often covers the increased need for nutrients during pregnancy. Using the right dietary supplements with good bacteria can have significant health benefits for pregnant women and their babies. Probiotic bacteria are live microorganisms which, when administered in appropriate amounts, exert beneficial effects in the host organism. The intestinal microbiota of a pregnant woman has a significant influence on the child's development.

List of references


