



OPEN ACCESS

EDITED AND REVIEWED BY
Sinead Waters,
University of Galway, Ireland

*CORRESPONDENCE
Francesca Lembo
✉ frlembo@unina.it

RECEIVED 04 November 2025
REVISED 17 November 2025
ACCEPTED 18 November 2025
PUBLISHED 11 December 2025

CITATION
Turrone S, Coretti L, Magzal F, Barone M,
O'Riordan KJ and Lembo F (2025) Editorial:
Innovative therapeutic strategies targeting
early-life gut microbiota: pathways to
long-term health benefits.
Front. Microbiol. 16:1739538.
doi: 10.3389/fmicb.2025.1739538

COPYRIGHT
© 2025 Turrone, Coretti, Magzal, Barone,
O'Riordan and Lembo. This is an open-access
article distributed under the terms of the
[Creative Commons Attribution License \(CC BY\)](#). The use, distribution or reproduction in
other forums is permitted, provided the
original author(s) and the copyright owner(s)
are credited and that the original publication
in this journal is cited, in accordance with
accepted academic practice. No use,
distribution or reproduction is permitted
which does not comply with these terms.

Editorial: Innovative therapeutic strategies targeting early-life gut microbiota: pathways to long-term health benefits

Silvia Turrone¹, Lorena Coretti², Faiga Magzal³, Monica Barone¹,
Kenneth J. O'Riordan⁴ and Francesca Lembo^{2*}

¹Department of Pharmacy and Biotechnology, University of Bologna, Bologna, Italy, ²Department of Pharmacy, University of Naples Federico II, Naples, Italy, ³Department of Nutritional Sciences, Tel-Hai College, Kiryat Shmona, Israel, ⁴College of Medicine and Health, University College Cork, Cork, Ireland

KEYWORDS

early life, therapeutic tools, host health, gut microbiota perturbations, long term effects

Editorial on the Research Topic

[Innovative therapeutic strategies targeting early-life gut microbiota: pathways to long-term health benefits](#)

The gut microbiota is increasingly recognized as a cornerstone of human development. During early life, this ecosystem of microorganisms plays critical roles in immune programming, neurodevelopment, and systemic homeostasis. It is considered a key intermediary between the “exposome,” which encompasses the full spectrum of environmental exposures across the lifespan, and the host’s genetic background, influencing the organism’s ability to either resist or be predisposed to pathological events. Indeed, perturbations of gut microbiota during sensitive developmental windows have been linked to a wide range of adverse outcomes (Lynch et al., 2023; Stokholm et al., 2018; Hyvönen et al., 2025; Beck et al., 2022; Zhang et al., 2025).

Despite substantial progress in this field, critical knowledge gaps persist regarding the role of early gut microbiota colonization in regulating host developmental processes and maintaining physiological homeostasis. Activities of early resident microbial communities are increasingly recognized as pivotal contributors to long-term health outcomes. Furthermore, advances in knowledge of the dynamic interactions between the microbiota and the host hold significant potential for the development of targeted therapeutic strategies.

This Research Topic brings together 15 contributions that span basic science, clinical research, meta-analysis, and methodological advances. This Research Topic highlights the principal topics of microbiota-early life area of study as emerged by the analysis of Yang et al. and contributes to delineating an increasingly clear picture of host response trajectories to the microbiota asset in early life. There are two broad themes covered.

The first theme, explored through a variety of methodological approaches including bibliometric analysis, Mendelian randomization, and basic science studies, focuses on the characterization of the microbiota in early childhood, animal models, and pathophysiological conditions.

In a prospective cohort study, very low birth weight newborns exhibited profound gut dysbiosis, with a disrupted microbiota trajectory characterized by a lack of obligate anaerobes and a fecal metabolomic shift during the first 28 days after birth. The substantial number of pediatric participants in the study supports the recommendation to monitor the fragility of the microbial ecosystem in high-risk neonates (Liu, Chen et al.). Similarly, a cohort study has shown that pregnancy-related conditions, such as hypertensive disorders, can have intergenerational effects (Yang et al.). Indeed, these conditions have been associated with concordant microbial alterations in both mothers and newborns. Several studies comparing *Helicobacter pylori*-infected and uninfected infants have identified an altered gastrointestinal microbiota profile in infected individuals (Chen et al., 2021). A meta-analysis by Wang, Tan et al. highlighted the pediatric implications of *H. pylori* infection, particularly its association with an increased risk of iron deficiency and anemia. This work lays the groundwork for future studies on infection-driven dysbiosis and its extra-gastrointestinal manifestations, such as iron deficiency anemia.

The link between the microbiota and allergic diseases is becoming increasingly clear, and meta-analyses are needed to aggregate findings and build platforms for causal effects studies. By subsequently applying Mendelian Randomization studies (either forward or reverse), Zheng, Chen, Zhuang, Xu et al. and Zheng, Chen, Zhuang, Zhao et al. established causal associations between specific microbial taxa and genetic predisposition to asthma and allergic rhinitis. Complementing this, an updated bibliometric analysis mapped the evolution of microbiota research trends toward atopic dermatitis (AD) in human infants, showing enrichment in publications linking microbiota to immune mechanisms and prebiotic intervention in AD (Wang, Wang et al.). This Research Topic also includes an observational study underscoring the composition and dynamics of the intestinal microbiota in infants with congenital heart disease (CHD), as well as potential influencing factors. It has been reported that the observed microbial patterns are linked to oxygenation and perfusion in CHD patients shortly after birth (Renk et al.).

A focus on host-microbe immune interactions, employing human-like advanced research models suitable to investigate the impact of commensal microbiota on immune function, has been developed by Liu, Zhang et al. In germ-free and specific-pathogen-free piglets, they observed that commensal microbiota impacted the abundance of various immune cell types and the proliferation and differentiation of lymphocytes.

In a different research context focused on neurobehavioral functions during early development, Nankova et al. employed a microbiota perturbation model to investigate the underlying mechanisms. Using a mouse model of postnatal antibiotic-induced dysbiosis, they demonstrated that altering the normal seeding and maturation of the postnatal microbiome affected neuroendocrine signaling, stress responses, and behavior in a sex-specific manner, providing mechanistic evidence that microbial composition can shape neurodevelopment.

The second theme covers approaches to delve into the effects of microbiota remodeling through therapeutic tools to impact early life and long-term health benefits. Included studies pinpoint the

use of complementary and alternative medical techniques and microbiota-based therapies in human and animal models.

Generally, both ozone rectal insufflation and electroacupuncture are being investigated for their modulatory effects on the gut microbiota. Through mechanisms involving the autonomic nervous system, anti-inflammatory pathways, and changes in the intestinal environment, these non-pharmacological interventions may influence microbial composition and host-microbe interactions, especially during early development or in dysbiotic conditions. In an experimental model of atherosclerosis (ApoE^{-/-} mice on a high-fat diet), ozone rectal insufflation was found to reduce plaque formation and serum low-density lipoprotein cholesterol levels, likely through an improvement in gut microbial health and regulation of microbial metabolites (Li et al.). In a rat model of perinatal nicotine exposure (PNE), electroacupuncture in the dams reduced the PNE-induced impairment of lung development in the offspring (Xie et al.). This effect was shown to be mediated by microbiota functions, since maternal antibiotic treatment blocked the beneficial effects of electroacupuncture on pulmonary function and lung morphology in offspring. Treatments able to support an eubiotic trajectory of microbiota colonization in neonates and preterm infants, and manage various dysbiosis-related diseases, are under study. Among these, probiotic supplementation is gathering attention as an adjunct therapy to manage neonatal gut health and recovery from diverse neonatal diseases. Probiotic-based complementary therapies, namely supplementation with *Clostridium butyricum* and *Bifidobacterium infantis*, were shown to ameliorate antibiotic-mediated dysbiosis in preterm infants with neonatal respiratory distress syndrome, contributing to disease recovery (Fu et al.).

Research studies by Hudcovic et al. and Yoon et al. to shape host diseases outcomes demonstrated the importance of probiotic-based therapies in early life to establish the correct microbiota signature and promote long-term health benefits, including protection from later pathological events. In animal models, the researchers specifically focused on a probiotic intervention aimed at protecting against intestinal inflammation (inflammatory colitis and enterotoxigenic *Escherichia coli*, ETEC, infection, respectively). Experiments in gnotobiotic mice showed that susceptibility to colitis was determined by the order of colonization by probiotic vs. pathogenic *E. coli* strains, emphasizing the importance of microbial priority effects to shape host outcomes (Hudcovic et al.). Also, Yoon et al. identified *Lactiplantibacillus argentoratensis* as a novel probiotic candidate that protects against ETEC infection, enhances barrier integrity, and promotes short-chain fatty acid production.

In conclusion, the articles included in this Research Topic help to shed light on the role of the gut microbiota in early development, its link to pathophysiological states and predisposition to disease, and the therapeutic opportunities of modifying aberrant microbiota structures to impact long-term health outcomes. As gut microbiota not only determines gut health, but also regulates systemic development affecting the nervous, respiratory, cardiovascular, metabolic, and immune systems, it is essential to strengthen our understanding of its role in shaping future health. This is particularly important from the earliest stages of development, given its early integration as a key modulator of the host's response capacity.

We thank all contributing authors for their valuable efforts, and we hope this Research Topic inspires further exploration and innovation at the intersection of microbiota, development, and health.

Author contributions

ST: Writing – original draft, Writing – review & editing. LC: Writing – review & editing, Writing – original draft, Conceptualization. FM: Writing – original draft, Writing – review & editing. MB: Writing – original draft, Writing – review & editing. KO'R: Writing – review & editing, Supervision, Writing – original draft. FL: Supervision, Conceptualization, Writing – review & editing, Writing – original draft.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

References

- Beck, L. C., Masi, A. C., Young, G. R., Vatanen, T., Lamb, C. A., Smith, R., et al. (2022). Strain-specific impacts of probiotics are a significant driver of gut microbiome development in very preterm infants. *Nat. Microbiol.* 7, 1525–1535. doi: 10.1038/s41564-022-01213-w
- Chen, C. C., Liou, J. M., Lee, Y. C., Hong, T. C., El-Omar, E. M., Wu, M. S., et al. (2021). The interplay between *Helicobacter pylori* and gastrointestinal microbiota. *Gut Microbes* 13, 1–22. doi: 10.1080/19490976.2021.1909459
- Hyvönen, S., Saarikivi, A., Mälkönen, J., Solasaari, T., Korpela, K., de Vos, W. M., et al. (2025). The association of maternal and infant early gut microbiota with respiratory infections in infants. *Pediatr. Res.* doi: 10.1038/s41390-025-04326-0
- Lynch, C. M. K., Cowan, C. S. M., Bastiaanssen, T. F. S., Moloney, G. M., Theune, N., van de Wouw, M., et al. (2023). Critical windows of early-life microbiota disruption on behaviour, neuroimmune function, and neurodevelopment. *Brain Behav. Immun.* 108, 309–327. doi: 10.1016/j.bbi.2022.12.008
- Stokholm, J., Blaser, M. J., Thorsen, J., Rasmussen, M. A., Waage, J., Vinding, R. K., et al. (2018). Maturation of the gut microbiome and risk of asthma in childhood. *Nat. Commun.* 9:141. doi: 10.1038/s41467-017-02573-2
- Zhang, M., Ding, L., Strodl, E., Yin, X., Wen, G., Sun, D., et al. (2025). Early supplement of probiotics reduces the risk of obesity among preschool children: a real-world observational study. *Front. Nutr.* 12:1597894. doi: 10.3389/fnut.2025.1597894

The author(s) declared that they were an editorial board member of Frontiers, at the time of submission. This had no impact on the peer review process and the final decision.

Generative AI statement

The author(s) declare that no Gen AI was used in the creation of this manuscript.

Any alternative text (alt text) provided alongside figures in this article has been generated by Frontiers with the support of artificial intelligence and reasonable efforts have been made to ensure accuracy, including review by the authors wherever possible. If you identify any issues, please contact us.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.