

When microbiome meets one health principle: Leading to the holy grail of biology and contributing to overall well-being and social sustainability

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Introduction

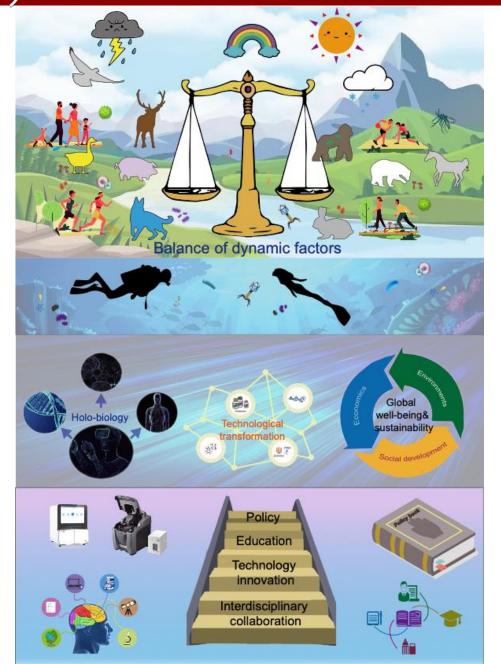
"Balance of Dynamic Factors" spotlights the equilibrium within microbial and human-animal-environment interactions, offering a revolutionary pathway to global health and social well-being. What does biology truly encompass?

How can we universally and systematically unveil the intricate biology woven within all life forms on our diminutive planet?



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Highlights





The microbiome is key for interdisciplinary solutions to global health issues.

A systems-thinking approach for disease surveillance and environmental care should



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be adopted.

Integrate the microbiome and One Health for understanding life complexity and shaping health policies.

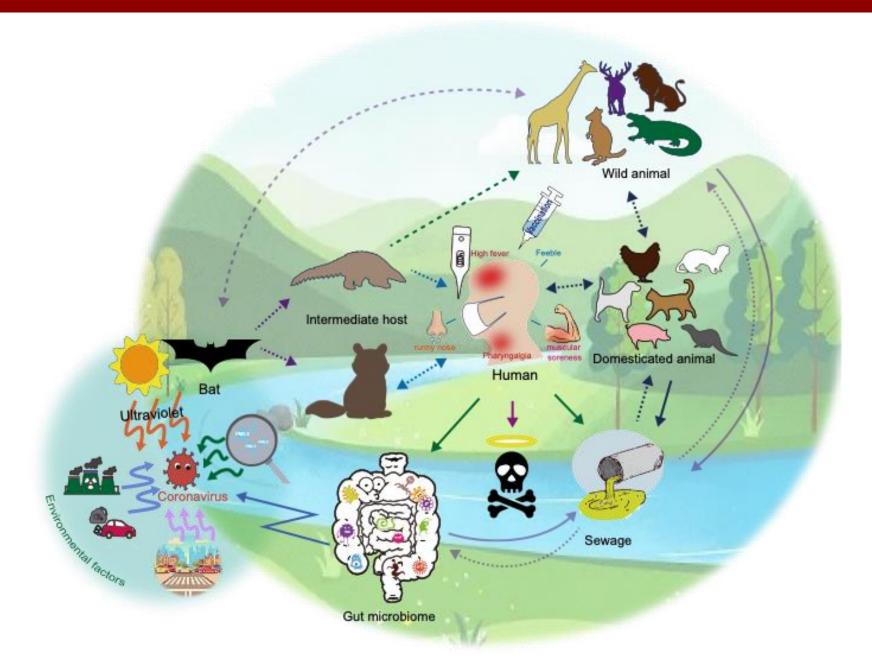
Microbiome: the Indispensable Genome for Creatures and Key Player for the Environment and Ecosystems





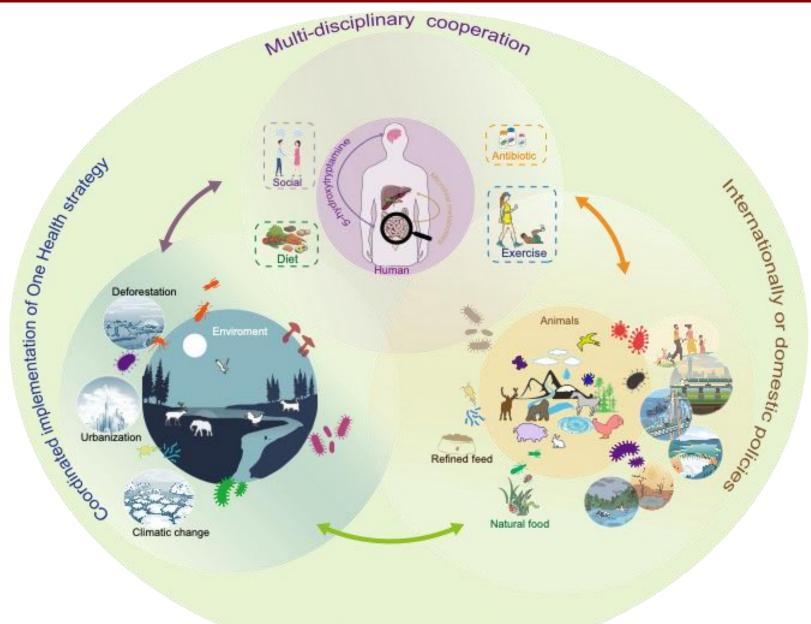


One Health: A Concept to Unite All Well-Beings in Our Globe



Microbiome Meets One-Health Principles: A Universal Balance Theory for Overall Health and Social Sustainability





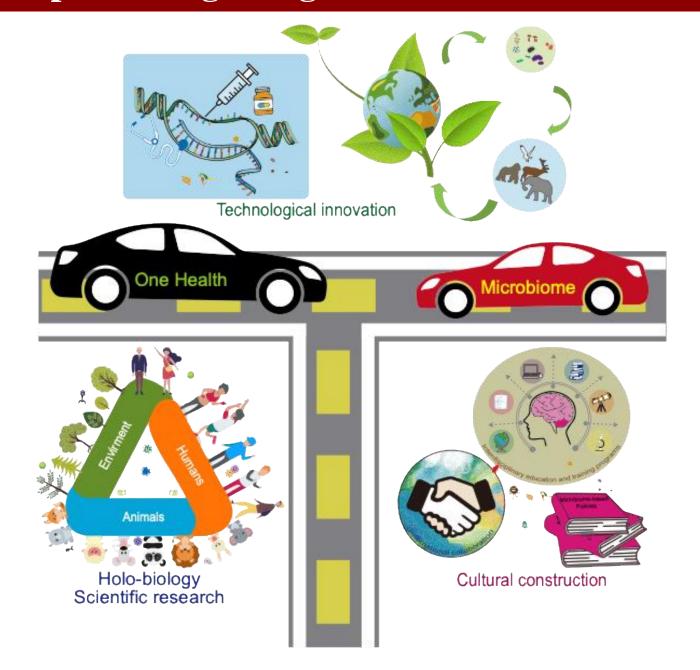


How Microbiome and One Health Principles Shape Biology in the Future





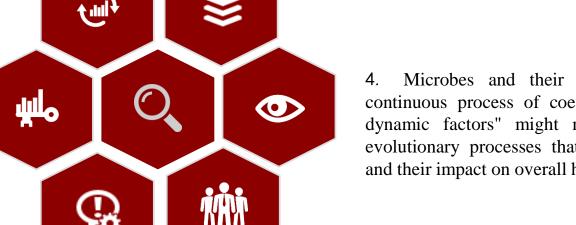
Next Step for Integrating Microbiome and One Health



Limitations of Theory "Balance of Dynamic Factors"

- 1. The complexity and heterogeneity of microbiomes make their combination with the One Health principle incredibly diverse and dynamic, varying across different hosts and environments. The "balance of dynamic factors" may not fully capture the complexity of these interactions because of the vast diversity of microbes and their functions.
- 3. Human activities significantly alter can microbiomes, sometimes leading to imbalances. The theory needs to consider the impact of these changes on the balance it seeks to maintain, especially with increasing recognition of the role of microbiomes in planetary health.
 - 5. Advances in microbiome research are often driven by new technologies, which can introduce biases or require new definitions and standards. This can make it difficult to apply the theory consistently across different studies and environments.

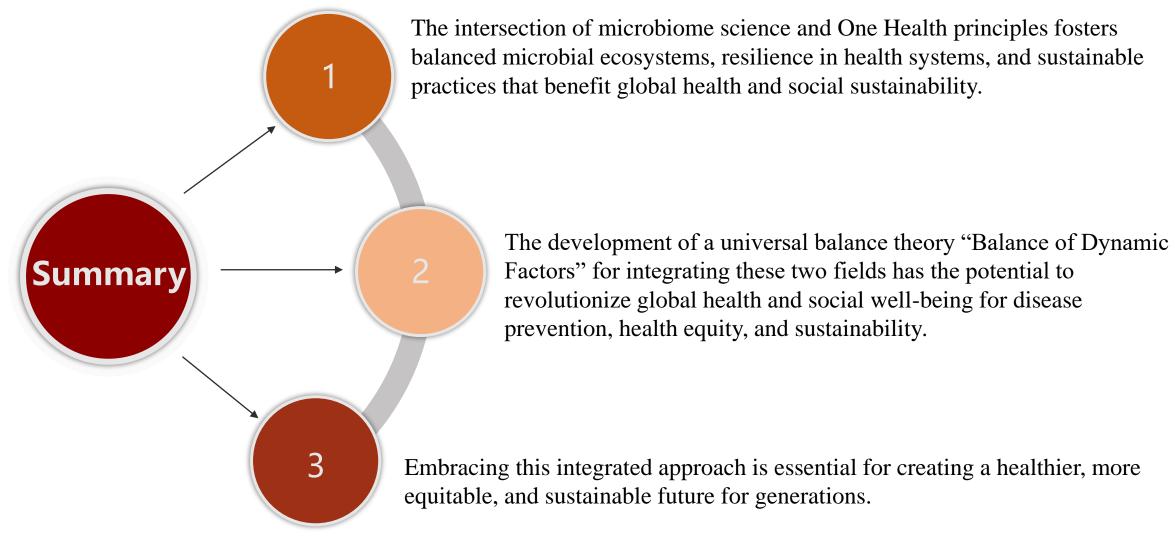
2. Lack of a unified definition of "balance of dynamic factors" causes issues in establishing a universally accepted framework for the theory.



- Microbes and their hosts are engaged in a continuous process of coevolution. The "balance of dynamic factors" might not fully encapsulate the evolutionary processes that drive these relationships and their impact on overall health and sustainability.
- 6. While this theory provides a conceptual framework, its practical application in managing microbiomes for health and sustainability is still in its infancy. There is a need for more research on how to manipulate microbiomes effectively to achieve the desired balance.
- 7. The "balance of dynamic factors" cuts across various disciplines, requiring a high level of interdisciplinary collaboration. This can be challenging owing to differences in terminology, methodologies, and research objectives.



Conclusion and Future Prospectives



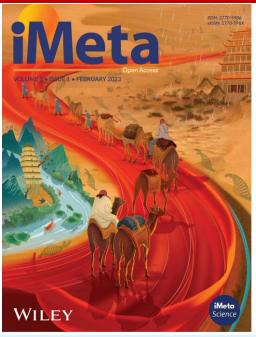
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iMeta: Integrated meta-omics to change the understanding of the biology and environment













"iMeta" is a Wiley partner journal launched by iMeta Science Society in 2022, receiving its first impact factor (IF) of 23.7 in 2024, ranking 2/165 in the microbiology field. It aims to publish innovative and high-quality papers with broad and diverse audiences. Its scope is similar to Nature Biotechnology, Nature Microbiology, and Cell Host & Microbe. Its unique features include video abstract, bilingual publication, and social media dissemination, with more than 500,000 followers. It has published 200+ papers and been cited for 4000+ times, and has been indexed by **ESCI/WOS/JCR**, **PubMed**, **Google Scholar**, and **Scopus**.

"iMetaOmics" is a sister journal of "iMeta" launched in 2024, with a target IF>10, and its scope is similar to Microbiome, ISME J, Nucleic Acids Research, Briefings in Bioinformatics, Bioinformatics, etc. All contributes are welcome!

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