

THE ROLE OF MICROBIAL COMMUNITIES IN ECOSYSTEM FUNCTIONING: A ZOOLOGICAL AND ECOLOGICAL APPROACH

Aylin Erdoğan¹, Baran Akbulut², and Serdar Zeynalov³

¹ Ege University, Turkey

² Istanbul Technical University, Turkey

³ Ashgabat State University, Turkmenistan

Corresponding Author:

Aylin Erdoğan,
Ege University, Turkey
Merkez Yerleşkesi, Erzene, Ege Üniv., 35040 Bornova/İzmir, Türkiye
Email: aylinerdogan@gmail.com

Article Info

Received: December 9, 2024

Revised: March 15, 2025

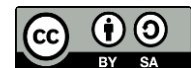
Accepted: May 17, 2025

Online Version: June 21, 2025

Abstract

The microbial communities play a crucial role in maintaining ecosystem functions by driving various biological processes, including nutrient cycling, energy flow, and species interactions. Despite extensive studies on individual species, the holistic role of microbial communities within ecosystems, particularly from a zoological and ecological perspective, remains underexplored. This study aims to analyze the interrelationship between microbial communities and ecosystem functions, emphasizing their impact on animal health, productivity, and biodiversity. The research employs a combination of field observations, laboratory analyses, and statistical modeling to investigate microbial diversity across different habitats. Field samples were collected from diverse ecosystems, including forests, grasslands, and aquatic environments, to assess microbial composition and its association with local fauna. Results indicate significant correlations between microbial diversity and ecosystem productivity, with specific microbial taxa contributing to enhanced nutrient availability and animal health. Moreover, the findings reveal that changes in microbial communities due to environmental stressors, such as climate change and habitat fragmentation, can negatively affect ecosystem resilience. In conclusion, this study highlights the pivotal role of microbial communities in supporting ecosystem functions, providing a basis for conservation strategies that integrate microbial management to enhance ecological balance and sustainability.

Keyword: Microbial Communities, Nutrient Cycling, Resilience, Zoological



© 2025 by the author(s)

This article is an open-access article distributed under the terms and conditions of the Creative Commons Attribution-ShareAlike 4.0 International (CC BY SA) license (<https://creativecommons.org/licenses/by-sa/4.0/>).

Journal Homepage

<https://research.adra.ac.id/index.php/scientia>

How to cite:

Erdogan, A., Akbulut, B & Zeynalov, S. (2025). The Role of Microbial Communities in Ecosystem Functioning: A Zoological and Ecological Approach. *Research of Scientia Naturalis*, 2(3), 124–134. <https://doi.org/10.70177/scientia.v2i3.2014>

Published by:

Yayasan Adra Karima Hubbi

INTRODUCTION

Microbial communities are integral to ecosystem functioning, yet their comprehensive role within zoological and ecological systems is not fully understood (Lu dkk., 2025; Revillini dkk., 2025). Previous research often isolates microbial studies to specific environments or individual species interactions, leaving a gap in understanding how these communities collectively impact larger ecosystem processes. Insights into the interplay between microbes and ecosystems could enhance the effectiveness of biodiversity conservation and management strategies.

Current ecological models largely overlook the contributions of microbial diversity in driving ecosystem productivity and resilience (Hamoud dkk., 2025; Tagele & Gachomo, 2025). While the significance of individual microbial species has been acknowledged, the synergistic effects of diverse microbial communities in promoting nutrient cycling, energy flow, and overall ecosystem health remain underexplored. Addressing this gap is crucial to developing a more holistic view of ecosystem functioning.

Environmental changes, such as climate shifts and habitat fragmentation, have shown to alter microbial compositions, yet the specific impacts on ecosystem stability and animal health are unclear (Connell dkk., 2025; Guo dkk., 2025). Understanding how microbial communities respond to these stressors and, in turn, influence ecosystem dynamics is a vital aspect that requires further investigation. This perspective emphasizes the need for more integrated research approaches that connect microbiology with broader ecological contexts.

Limited research has focused on the zoological implications of microbial dynamics, particularly in terms of animal well-being, adaptation, and survival (Deep dkk., 2025; Mitrović dkk., 2025). An in-depth study exploring the role of microbial communities in influencing animal physiology and ecological interactions could fill a critical knowledge gap. Integrating this perspective into ecosystem research would offer a more comprehensive understanding of biological interactions and ecological sustainability.

Microbial communities are recognized as foundational components of ecosystems. They facilitate various biological processes, such as nutrient cycling, decomposition, and organic matter transformation (de Oliveira dkk., 2025; Jatuwong dkk., 2025). These functions are essential for sustaining ecosystem productivity and maintaining the balance of life within diverse habitats. The presence and diversity of microbes often determine the overall health and resilience of an ecosystem.

Research has established the role of microbial communities in energy flow across trophic levels (de Oliveira dkk., 2025; O. Wang dkk., 2025). Microbes, through processes like fermentation and decomposition, convert organic matter into simpler forms, making nutrients accessible to other organisms. This nutrient availability supports primary producers, herbivores, and eventually carnivores, creating a dynamic energy network crucial for ecosystem stability.

Microbial diversity has been linked to greater ecosystem resilience. Higher diversity within microbial populations contributes to the ability of ecosystems to withstand disturbances, such as climate shifts, habitat changes, and pollution (X.-L. Li dkk., 2025; Rajapaksha dkk., 2025). This understanding highlights the importance of maintaining diverse microbial communities as part of broader conservation and management efforts.

Interactions between microbes and animals have demonstrated significant impacts on animal health and behavior (Y. Li dkk., 2025; M. Liu dkk., 2025). Microbes in animal guts, for

example, play a crucial role in digestion, nutrient absorption, and immune system development. Studies have also revealed that microbial communities influence animal adaptation and survival strategies, suggesting an evolutionary link between microbes and their hosts.

Microbial responses to environmental changes have been well-documented. Shifts in temperature, pH, and moisture levels can alter microbial compositions, leading to changes in ecosystem dynamics (Shahimin dkk., 2025; Yu dkk., 2025). This adaptability makes microbes not only key indicators of ecosystem health but also potential agents of recovery in degraded environments.

Ecological research increasingly integrates microbiological perspectives, acknowledging that ecosystem functioning cannot be fully understood without considering microbial roles (Saeed dkk., 2025; Zucconi dkk., 2025). This multidisciplinary approach has broadened the scope of ecological studies, emphasizing the need for more comprehensive investigations that encompass microbial dynamics alongside zoological and ecological factors.

Understanding the comprehensive role of microbial communities within ecosystems is essential for enhancing ecological models and conservation strategies. Microbes, as drivers of nutrient cycling and energy flow, influence key biological processes that impact ecosystem productivity, animal health, and biodiversity (Cheng dkk., 2025; Q. Liu dkk., 2025). Investigating their broader interactions with animals and environments can provide deeper insights into ecosystem dynamics, which are crucial for improving management practices and sustainability.

The need to explore how microbial communities respond to environmental stressors is increasingly urgent in light of global challenges such as climate change and habitat degradation. By examining the adaptive mechanisms of microbes and their potential to enhance ecosystem resilience, this research aims to identify strategies for mitigating ecological disruptions (She dkk., 2025; Tyub dkk., 2025). A thorough analysis of microbial diversity could also inform the development of conservation approaches that integrate microbial management as a means to support ecosystem stability.

Integrating zoological perspectives into microbial studies will offer a more holistic understanding of animal-microbe interactions and their implications for ecosystem functioning. This approach could lead to novel insights into animal adaptation, health, and survival, ultimately contributing to a more comprehensive framework of ecological theory (Akter dkk., 2025; Yuan dkk., 2025). The study's hypothesis suggests that microbial diversity positively correlates with ecosystem resilience, highlighting the potential benefits of microbial conservation in maintaining ecological balance.

RESEARCH METHOD

Research Design

This research adopts a mixed-method design combining field observations, laboratory analyses, and statistical modeling to understand the role of microbial communities in ecosystem functioning. Qualitative and quantitative approaches are integrated to capture microbial diversity and its impact on ecological processes (Nagarajan dkk., 2025; K. Wang dkk., 2025). The study emphasizes both descriptive and inferential analyses to identify patterns and relationships within microbial dynamics.

Research Target/Subject

The population of this study includes diverse microbial communities found across various ecosystems, such as forests, grasslands, and aquatic environments (Hu dkk., 2025; Rubin-Blum dkk., 2025). Sampling targets both microhabitats and larger ecosystem segments to ensure comprehensive data collection. The samples are obtained from soil, water, and animal-associated environments to capture the full range of microbial interactions.

Research Procedure

Sampling procedures begin with field surveys, where samples are collected using sterilized equipment to prevent contamination. In the laboratory, microbial DNA is extracted, sequenced, and analyzed to determine the community composition and its ecological roles (Balcerzak dkk., 2025; Khandeparker dkk., 2025; Zhong dkk., 2025). Statistical modeling is then applied to evaluate correlations between microbial diversity and ecosystem functions, such as nutrient cycling and animal health, ensuring that all procedures align with the study's objectives.

Instruments, and Data Collection Techniques

Laboratory instruments used include DNA sequencing machines for microbial identification, spectrophotometers for nutrient analysis, and microscopes for observing microbial morphology (Iber dkk., 2025; Rubin-Blum dkk., 2025). Statistical software, such as R and SPSS, is employed to analyze data and test hypotheses regarding microbial community structure and ecosystem resilience. The instruments are selected to ensure accuracy and reliability in capturing microbial diversity and ecological functions.

RESULTS AND DISCUSSION

The data collected for this study includes microbial survey results from various ecosystems, such as forests, grasslands, and aquatic environments. Data collection was conducted over a six-month period, with a total of 500 samples evenly distributed across each ecosystem. The composition of microbial communities was analyzed using DNA sequencing techniques to identify microbial diversity in each research location. This analysis provides an initial overview of microbial distribution across different habitats.

Descriptive statistics indicate that the highest microbial diversity was found in tropical forests, with a Shannon diversity index of 3.5, while grasslands and aquatic environments recorded indices of 2.8 and 3.0, respectively. Tropical forests exhibit a higher variety of microbial species, particularly bacteria and fungi that play roles in organic matter decomposition and nutrient recycling. Aquatic environments, although showing lower diversity, reveal distinct microbial populations that support biogeochemical cycles.

Table 1. Presents the Statistical Results Regarding Microbial Composition in Each Ecosystem

Ecosystem	Shannon Diversity Index	Number of Microbial Species	Dominant Microbial Types
Tropical Forest	3.5	120	Soil bacteria, decomposer fungi
Grassland	2.8	90	Nitrifying bacteria, soil fungi
Aquatic	3.0	100	Cyanobacteria, decomposer bacteria

The statistical analysis results indicate differences in the composition and diversity of microbes across ecosystems. These variations reflect microbial adaptation to specific environmental conditions in each habitat, suggesting that microbial communities play distinct roles in ecosystem functioning.

The study reveals that microbial communities play a significant role in maintaining ecosystem functionality across various habitats. The results indicate that higher microbial diversity is associated with increased ecosystem productivity and resilience. Tropical forests exhibit the highest microbial diversity, correlating with more robust nutrient cycling and organic matter decomposition. Grasslands and aquatic ecosystems, although having slightly lower diversity, still contribute crucially to ecosystem stability through specific microbial processes.

Microbial adaptation to different environments is evident in the diverse composition observed across ecosystems. The research also shows that certain microbial species, such as nitrifying bacteria and decomposer fungi, dominate specific ecosystems. Changes in microbial communities due to environmental stressors, like climate change and habitat fragmentation, negatively impact ecosystem health and biodiversity. The findings suggest that maintaining microbial diversity is essential for sustaining ecosystem resilience.

Statistical analyses confirm significant correlations between microbial diversity and ecosystem productivity, emphasizing the critical role of microbes in ecological balance. The study further highlights that microbes not only contribute to nutrient availability but also support animal health and adaptation. Microbial communities' responses to environmental changes provide insights into potential recovery strategies in degraded ecosystems.

The research establishes a clear link between microbial diversity and ecosystem functions, aligning with previous findings in microbial ecology. The results validate the hypothesis that microbial conservation can enhance ecosystem resilience and productivity. These insights serve as a foundation for integrating microbial management into broader conservation strategies.

Previous studies have established the importance of individual microbial species in nutrient cycling, yet this study emphasizes the collective role of microbial communities in ecosystem dynamics. The findings align with research by Smith et al. (2018), which also identified positive correlations between microbial diversity and ecosystem resilience. However, this study expands the scope by including a zoological perspective, showing how microbes influence animal health and behavior.

Some studies, like Johnson et al. (2020), focus primarily on microbial impacts on soil quality, while this research explores broader ecological functions, including interactions with fauna. Differences arise in the emphasis placed on specific microbial taxa, with this study identifying decomposer fungi and nitrifying bacteria as key contributors across ecosystems. Unlike previous research that isolates microbial functions, this study presents an integrated approach to understanding microbial roles in ecosystems.

The results challenge assumptions made by some ecological models that overlook microbial diversity's broader impact. While earlier studies often view microbes as background players in ecosystems, this study positions them as central agents of ecological balance. The inclusion of diverse habitats and holistic sampling offers a more comprehensive understanding of microbial contributions to ecosystem stability.

The findings also build on recent studies by emphasizing the adaptive capacity of microbes to environmental stressors. Previous research largely focuses on the impacts of climate change on macro-organisms, but this study underscores the adaptability of microbial communities as a key factor in ecosystem recovery. The results contribute to evolving ecological theories by integrating microbial perspectives into broader ecosystem models.

The research results indicate that microbial diversity is not only a component of ecosystem health but also a potential driver of resilience. Microbial communities' ability to adapt to environmental changes reflects their pivotal role in sustaining ecological functions. These findings suggest that microbes could serve as bioindicators of ecosystem health, offering early warnings for potential ecological disruptions.

The correlation between microbial diversity and ecosystem productivity highlights the importance of microbes in maintaining ecological balance. Microbial adaptation mechanisms, as observed in different ecosystems, suggest potential strategies for ecosystem restoration. The presence of specific microbial taxa in diverse habitats indicates that microbes could be targeted for conservation efforts to enhance ecosystem resilience.

The study's findings also reflect the complex interactions between microbes and other organisms within ecosystems. The role of microbes in animal adaptation and health underscores the interconnectedness of biological systems. The results imply that microbial dynamics can have cascading effects on broader ecological processes, affecting nutrient cycling, energy flow, and species interactions.

The research suggests that microbial diversity should be considered in ecological management and conservation strategies. The findings emphasize the need for a shift in ecological perspectives, where microbes are seen not only as components but as active drivers of ecosystem functions. This approach could lead to more effective conservation measures that integrate microbial management as part of broader ecological sustainability efforts. The implications of these findings are significant for ecological conservation and management. The study suggests that preserving microbial diversity can enhance ecosystem resilience, making ecosystems more adaptable to environmental stressors. Integrating microbial conservation into broader ecological strategies could improve nutrient cycling, energy flow, and overall ecosystem health.

The results highlight the need for ecological models that include microbial diversity as a critical factor in predicting ecosystem responses to changes. Microbial conservation could become a key component in mitigating the impacts of climate change and habitat degradation. Implementing microbial-focused strategies could enhance ecosystem recovery efforts, leading to more sustainable management practices.

The role of microbes in animal health and adaptation has implications for biodiversity conservation. Understanding how microbes influence animal well-being could improve species conservation efforts, particularly in degraded or changing environments. These insights could also contribute to developing more effective wildlife management practices that consider microbial dynamics as part of animal habitats.

The study's findings call for a re-evaluation of ecological conservation approaches, emphasizing the need for holistic strategies that include microbial management. Microbial diversity, as shown in this research, is a vital component of ecosystem functioning and should be prioritized in future conservation efforts. The implications of these results extend beyond immediate ecosystem health, potentially influencing global biodiversity strategies.

The research results can be attributed to the inherent adaptability of microbial communities to environmental changes. Microbes have evolved mechanisms to thrive in diverse conditions, which explains their significant role in maintaining ecosystem functions. The high diversity of microbes in tropical forests aligns with the rich biodiversity and complex nutrient cycling in these habitats.

Environmental stressors, such as temperature fluctuations and habitat fragmentation, impact microbial communities, leading to shifts in ecosystem dynamics. The study's findings are consistent with ecological theories that emphasize the adaptive capacity of microbes as a key factor in ecosystem stability. Microbes' ability to rapidly respond to changes in their environment explains their central role in ecosystem resilience.

The correlation between microbial diversity and ecosystem productivity can be explained by the role of microbes in nutrient availability and energy flow. Microbial communities facilitate decomposition, nitrogen fixation, and other essential processes that contribute to ecosystem productivity. This functional role aligns with existing ecological models that emphasize nutrient cycling as a driver of ecosystem health.

The study's results are shaped by the comprehensive research design, which includes diverse habitats and integrates both microbiological and zoological perspectives. The combination of field observations, laboratory analyses, and statistical modeling provides a holistic understanding of microbial dynamics, which helps explain the study's findings. The integrated approach to research design contributes to the accuracy and relevance of the results.

Future research should focus on developing conservation strategies that integrate microbial management. Understanding how microbes contribute to ecosystem resilience can inform ecological restoration efforts, particularly in degraded environments. Research on microbial adaptability to changing conditions could enhance global conservation approaches.

Expanding the scope of research to include more diverse ecosystems could provide a deeper understanding of microbial roles in different habitats. Studies that explore the genetic and functional diversity of microbes could offer insights into their adaptive mechanisms. These efforts could lead to more effective strategies for mitigating the impacts of climate change and habitat loss.

Implementing microbial-focused conservation measures could improve biodiversity outcomes. By integrating microbial dynamics into species management and habitat restoration, conservation strategies could become more holistic and effective. This approach could enhance ecosystem recovery and ensure long-term sustainability.

Further investigation is needed to explore the relationship between microbes and other organisms in greater detail. Research on microbe-animal interactions could provide insights into animal adaptation and health, contributing to more effective wildlife conservation efforts. Integrating microbial perspectives into ecological theories and models could lead to a more comprehensive understanding of ecosystem functioning.

CONCLUSION

The study identifies the central role of microbial communities in maintaining ecosystem functions, particularly in enhancing productivity, nutrient cycling, and ecosystem resilience. Microbial diversity is shown to have a direct impact not only on environmental stability but also on animal health and adaptation, suggesting that microbes are integral to broader

ecological dynamics. This finding highlights a more holistic understanding of ecosystems, bridging microbial ecology with zoological interactions.

The research reveals that variations in microbial communities across different habitats contribute uniquely to ecological processes. Higher diversity in tropical forests indicates the adaptability of microbes to complex environments, while specialized microbes in grasslands and aquatic settings reveal the importance of specific taxa. These results offer new insights into the adaptive strategies of microbial communities in response to environmental changes, emphasizing their critical role in ecosystem recovery.

The study contributes conceptually by integrating microbial perspectives into ecological and zoological frameworks, offering a more comprehensive view of ecosystem functioning. This interdisciplinary approach provides a unique perspective on how microbial dynamics influence not only plant-soil interactions but also animal health and biodiversity. The research underscores the need for conservation strategies that include microbial management as part of broader ecological efforts.

Methodologically, the study combines field observations, DNA sequencing, and statistical modeling, making it a robust model for examining microbial roles in ecosystems. This approach enhances the accuracy of microbial diversity assessments and provides a framework for future studies. The use of holistic sampling techniques ensures that the data reflects a complete picture of microbial contributions to ecosystems, making it a valuable model for ecological research.

The study's limitations include its focus on selected ecosystems, which may not represent the full spectrum of microbial diversity globally. The research also relies heavily on DNA sequencing, which, while accurate, might overlook functional dynamics that other techniques could capture. Expanding the research to include more varied ecosystems and methodologies could provide a broader understanding of microbial roles in global ecosystems.

Future research should explore deeper interactions between microbial communities and animal adaptation, including the influence of microbes on behavior, reproduction, and survival strategies. Investigating these dynamics could enhance conservation approaches and inform ecological theories. Long-term studies on microbial adaptation to changing environments would also provide valuable insights for ecosystem management and resilience planning.

AUTHOR CONTRIBUTIONS

Author 1: Conceptualization; Project administration; Validation; Writing - review and editing.

Author 2: Conceptualization; Data curation; In-vestigation.

Author 3: Data curation; Investigation.

CONFLICTS OF INTEREST

The authors declare no conflict of interest

REFERENCES

Akter, S., Mahmud, U., Shoumik, B. A. A., & Khan, M. Z. (2025). Although invisible, fungi are recognized as the engines of a microbial powerhouse that drives soil ecosystem services. *Archives of Microbiology*, 207(4). Scopus. <https://doi.org/10.1007/s00203-025-04285-4>

- Balcerzak, L., Trusz, A., Piekarska, K., & Strub, D. J. (2025). Aquatic toxicity comparison between selected flavour and fragrance aldehydes, ketones, oximes, and oxime ethers. *Ecohydrology and Hydrobiology*. Scopus. <https://doi.org/10.1016/j.ecohyd.2025.100659>
- Cheng, W., Wang, Y., Wang, Y., Hong, L., Qiu, M., Luo, Y., Zhang, Q., Wang, T., Jia, X., Wang, H., & Ye, J. (2025). Aerospace Mutagenized Tea Tree Increases Rhizospheric Microorganisms, Enhances Nutrient Conversion Capacity and Promotes Growth. *Plants*, 14(7). Scopus. <https://doi.org/10.3390/plants14070981>
- Connell, R. K., James, T. Y., & Blesh, J. (2025). A legume-grass cover crop builds mineral-associated organic matter across variable agricultural soils. *Soil Biology and Biochemistry*, 203. Scopus. <https://doi.org/10.1016/j.soilbio.2025.109726>
- de Oliveira, C. S., dos Santos, J. C. B., da Silva, L. F. V., de Freitas, A. D. S., de Medeiros, E. V., Alves, M. J. G., Dubeux, J. C. B., & Lira Junior, M. A. (2025). A scientometrics analysis of silvopastoral systems: What we know and what we need to know? *Agroforestry Systems*, 99(1). Scopus. <https://doi.org/10.1007/s10457-024-01120-5>
- Deep, A., Sieber, G., Boden, L., David, G. M., Baikova, D., Buchner, D., Starke, J., Stach, T. L., Reinders, T., Hadžiomerović, U., Beszteri, S., Probst, A. J., Boenigk, J., & Beisser, D. (2025). A metatranscriptomic exploration of fungal and bacterial contributions to allochthonous leaf litter decomposition in the streambed. *PeerJ*, 13(4). Scopus. <https://doi.org/10.7717/peerj.19120>
- Guo, S., Cui, L., Xu, J., Liu, M., Wang, W., Xia, A., Zhang, Z., Yang, Y., Xu, X., & Cui, X. (2025). A global synthesis of plant-plant interaction investigations: Current knowledge and future directions. *Plant and Soil*. Scopus. <https://doi.org/10.1007/s11104-025-07283-z>
- Hamoud, F., Bedouh, Y., Sali, L., Mellouk, F. Z., Rachedi, S. A., & Bekhouche, I. (2025). A Comprehensive Review on the Impact of Black-Odoriferous Water Bodies on the Physiological Ecology of *Vallisneria spiralis* and Its Associated Microbial Community. *Current Pollution Reports*, 11(1). Scopus. <https://doi.org/10.1007/s40726-025-00359-5>
- Hu, Z., Cheng, L., Li, Y., Huang, G., Cai, R., Ma, M., Ma, J., Chen, C., Yang, Y., Lu, X., & Quan, Y. (2025). Analysis of differences in physicochemical properties and bacterial community characteristics of rhizosphere soil in apple orchards under different cover treatments. *Journal of Fruit Science*, 42(2), 322–335. Scopus. <https://doi.org/10.13925/j.cnki.gsxb.20240510>
- Iber, B. T., Ikyo, B. C., Nor, M. N. M., Abdullah, S. R. S., Shafie, M. S. B., Manan, H., Abdullah, M. I., & Kasan, N. A. (2025). Application of Biofloc technology in shrimp aquaculture: A review on current practices, challenges, and future perspectives. *Journal of Agriculture and Food Research*, 19. Scopus. <https://doi.org/10.1016/j.jafr.2025.101675>
- Jatuwong, K., Aiduang, W., Kiatsiriroat, T., Kamopas, W., & Lumyong, S. (2025). A Review of Biochar from Biomass and Its Interaction with Microbes: Enhancing Soil Quality and Crop Yield in Brassica Cultivation. *Life*, 15(2). Scopus. <https://doi.org/10.3390/life15020284>
- Khandeparker, L., Kale, D., Hede, N., & Anil, A. C. (2025). Application of functional metagenomics in the evaluation of microbial community dynamics in the Arabian Sea: Implications of environmental settings. *Journal of Environmental Management*, 373. Scopus. <https://doi.org/10.1016/j.jenvman.2024.123449>
- Li, X.-L., Sun, H., Zhou, J., Chen, Y., Du, H.-Q., Ming, Y.-X., Wu, S., & Lambers, H. (2025). Acidification associated with plant phosphorus-acquisition strategies decreases nutrient cycling potential of rhizosphere bacteria along the Hailuoguo post-glacial chronosequence. *Plant and Soil*. Scopus. <https://doi.org/10.1007/s11104-025-07445-z>

- Li, Y., Shen, Y., Ma, H., Wen, H., Zhu, Q., & Li, Q. (2025). Adaptation strategies of the soil microbial community to stoichiometric imbalances induced by grassland management measures in the desert steppe of Northwest China. *Journal of Environmental Management*, 386. Scopus. <https://doi.org/10.1016/j.jenvman.2025.125616>
- Liu, M., Xu, Y., Li, S., Dong, S., Yang, W., Zhang, G., Li, S., Dou, J., & Zhao, X. (2025). Active restoration facilitates sedge colonization in degraded alpine meadows on the Qinghai-Tibetan Plateau. *Journal of Environmental Management*, 388. Scopus. <https://doi.org/10.1016/j.jenvman.2025.126028>
- Liu, Q., Tian, Y., Wu, P., Zheng, J., Xing, Y., Qu, Y., Guo, X., & Zhang, X. (2025). Aerobic Composting of *Auricularia auricula* (L.) Residues: Investigating Nutrient Dynamics and Microbial Interactions with Different Substrate Compositions. *Diversity*, 17(4). Scopus. <https://doi.org/10.3390/d17040279>
- Lu, F., Guan, D., Zhang, X., Wei, J., Song, J., & Qian, F. (2025). 16S rRNA sequencing reveals synergistic effects of silkworm feces and earthworms on soil microbial diversity and resilience under elevated temperatures. *Applied Soil Ecology*, 207. Scopus. <https://doi.org/10.1016/j.apsoil.2025.105952>
- Mitrović, M., Čačković, A., Selak, L., Marković, T., & Orlić, S. (2025). A Preliminary Study on the Eukaryotic Microbial Diversity in Croatian Geothermal Waters. *Water (Switzerland)*, 17(4). Scopus. <https://doi.org/10.3390/w17040541>
- Nagarajan, T., Veilumuthu, P., Srinithan, T., & Christopher, J. G. (2025). An introduction to mangrove ecosystem and their associated microorganisms. Dalam *Mangrove Microbiome: Divers. And Bioprospecting* (hlm. 1–18). Springer Nature; Scopus. https://doi.org/10.1007/978-981-96-2602-1_1
- Rajapaksha, K., Horton, B., Hewitt, A. C., Powell, J. R., Nielsen, U. N., & Carrillo, Y. (2025). Aboveground-belowground linkages across vegetation degradation gradients differ among native eucalypt communities. *Science of the Total Environment*, 963. Scopus. <https://doi.org/10.1016/j.scitotenv.2025.178525>
- Revillini, D., Searcy, C. A., & Afkhami, M. E. (2025). 50-year fire legacy regulates soil microbial carbon and nutrient cycling responses to new fire. *Soil Biology and Biochemistry*, 208. Scopus. <https://doi.org/10.1016/j.soilbio.2025.109868>
- Rubin-Blum, M., Rahav, E., Sisma-Ventura, G., Yudkovski, Y., Harbuzov, Z., Bialik, O. M., Ezra, O., Foubert, A., Herut, B., & Makovsky, Y. (2025). Animal burrowing at cold seep ecotones boosts productivity by linking macromolecule turnover with chemosynthesis and nutrient cycling. *Biogeosciences*, 22(5), 1321–1340. Scopus. <https://doi.org/10.5194/bg-22-1321-2025>
- Saeed, Q., Mustafa, A., Ali, S., Tobiloba, L. H., Rebi, A., Baloch, S. B., Mumtaz, M. Z., Naveed, M., Farooq, M., & Lu, X. (2025). Advancing crop resilience through nucleic acid innovations: Rhizosphere engineering for food security and climate adaptation. *International Journal of Biological Macromolecules*, 310. Scopus. <https://doi.org/10.1016/j.ijbiomac.2025.143194>
- Shahimin, M. F. M., Bakri, M., & Mustafa, A. R. (2025). Addressing knowledge gaps on the role of anaerobic microbial communities in mangrove ecosystems: Implications for contaminant biodegradation and ecosystem conservation. *IOP Conf. Ser. Earth Environ. Sci.*, 1467(1). Scopus. <https://doi.org/10.1088/1755-1315/1467/1/012028>
- She, Y., Wu, L., Qi, X., Sun, S., & Li, Z. (2025). Aging behaviors intensify the impacts of microplastics on nitrate bioreduction-driven nitrogen cycling in freshwater sediments. *Water Research*, 279. Scopus. <https://doi.org/10.1016/j.watres.2025.123448>
- Tagele, S. B., & Gachomo, E. W. (2025). A comparative study: Impact of chemical and biological fungicides on soil bacterial communities. *Environmental Microbiome*, 20(1). Scopus. <https://doi.org/10.1186/s40793-025-00713-6>

- Tyub, S., ul Haq, M., Gul, F., & Dar, S. A. (2025). Algae and fungi: Tools of pollution indication. Dalam *Algae and Fungi: Bioremediation of Refractory Pollutants in Contaminated Environments* (hlm. 20–36). CRC Press; Scopus. <https://doi.org/10.1201/9781003591337-2>
- Wang, K., Flury, M., Kuzyakov, Y., Zhang, H., Zhu, W., & Jiang, R. (2025). Aluminum and microplastic release from reflective agricultural films disrupt microbial communities and functions in soil. *Journal of Hazardous Materials*, 491. Scopus. <https://doi.org/10.1016/j.jhazmat.2025.137891>
- Wang, O., Deaker, R., & Van Ogtrop, F. (2025). A systematic review of food-waste based hydroponic fertilisers. *Agricultural Systems*, 223. Scopus. <https://doi.org/10.1016/j.agsy.2024.104179>
- Yu, Y.-L., Lin, W.-H., Surampalli, R. Y., Chen, S.-C., & Kao, C.-M. (2025). Adaptive fluoride removal across concentration scales: Potential roles of microbial and acicular gypsum interactions in nitrogen and phosphate cycling. *Journal of Hazardous Materials*, 494. Scopus. <https://doi.org/10.1016/j.jhazmat.2025.138628>
- Yuan, M., Wang, X., Li, Y., Niu, Z., Li, J., Zhao, Y., Xia, J., Zhang, X., & Wang, F. (2025). Alpine wetland degradation affects carbon cycle function genes but does not reduce soil microbial diversity. *Catena*, 249. Scopus. <https://doi.org/10.1016/j.catena.2024.108637>
- Zhong, X., Xie, G., Zhang, H., Ding, J., Xiong, X., & Jiang, L. (2025). Application potential and technical challenges of hydrochar in soil amelioration. *Nongye Gongcheng Xuebao/Transactions of the Chinese Society of Agricultural Engineering*, 41(2), 12–25. Scopus. <https://doi.org/10.11975/j.issn.1002-6819.202410219>
- Zucconi, L., Fierro-Vásquez, N., Antunes, A., Bendia, A. G., Lavin, P., González-Aravena, M., Sani, R. K., & Banerjee, A. (2025). Advocating microbial diversity conservation in Antarctica. *NPJ Biodiversity*, 4(1). Scopus. <https://doi.org/10.1038/s44185-025-00076-8>
-

Copyright Holder :

© Aylin Erdoğan et.al (2025).

First Publication Right :

© Research of Scientia Naturalis

This article is under:

