



REVIEW

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Microbial Niches Across North Indian Alluvial Soils

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Abstract

Alluvial soils across northern Indian states represent one of the most fertile and biologically dynamic landscapes on the Earth. Formed through centuries of sediment deposition by the Ganga, Yamuna, Brahmaputra, and their tributaries, these soils sustain dense agricultural systems, diverse vegetation and complex microbial communities. This review examines microbial niches across the North Indian alluvial plains, emphasizing ecological patterns, nutrient cycling, and adaptive strategies of microbial taxa under varied physicochemical conditions. It synthesizes data from studies across Uttar Pradesh, Bihar, Haryana, Punjab, and Delhi to explore how edaphic factors such as texture, moisture, organic carbon, pH, and salinity shape microbial composition and activity. The article highlights microbial guilds involved in nitrogen fixation, phosphorus solubilization, organic matter decomposition, and stress tolerance, with case studies linking soil microbiomes to crop productivity. Climate change, anthropogenic pollution, and agrochemical use are analyzed as major pressures altering microbial resilience and soil health. The review concludes with an integrated model for sustainable management of microbial resources in alluvial agroecosystems, advocating for molecular-level monitoring, biotechnological applications, and restoration practices.

Keywords: Alluvial soils; Microbial diversity; Nitrogen fixation; Soil ecology; Gangetic plains; Biogeochemical cycles; North India

Introduction

The Gangetic alluvial plains of North India constitute one of the world's most intensively cultivated and ecologically rich biomes. Extending across Uttar Pradesh, Bihar, Haryana, Punjab, and Delhi, this region owes its fertility to periodic sediment deposition by the Ganga and Yamuna river systems (Singh et al., 2022). The resulting alluvium forms deep, well-drained soils with high cation exchange capacity and abundant nutrients, supporting both traditional and modern agriculture. Beneath this agronomic success lies an invisible but essential component the soil microbial community that governs nutrient cycling, soil aggregation, organic matter decomposition, and plant-microbe symbioses (Kumar & Raghav, 2021).

Microbial niches in these soils are defined by a mosaic of environmental gradients moisture, texture, temperature, and redox potential creating microhabitats for bacteria, archaea, fungi, and actinomycetes (Tripathi et al., 2020). The Gangetic plains' high heterogeneity in land use, irrigation patterns, and flood regimes leads to differential microbial assemblages, from aerobic decomposers in elevated ridges to anaerobic denitrifiers in waterlogged basins (Sharma et al., 2022). These microbial assemblages not only mediate nutrient turnover but also reflect ecological adaptation to seasonal hydrological cycles.

In the broader ecological context, microbial ecology in alluvial soils connects terrestrial, aquatic, and atmospheric systems. Nitrogen-fixing bacteria such as *Rhizobium*, *Azospirillum*, and *Cyanobacteria* play vital roles in maintaining soil fertility without synthetic inputs (Mandal et al., 2023). Similarly, phosphate-solubilizing microbes (PSMs) and mycorrhizal fungi enhance plant nutrient uptake under high-pH and saline conditions typical of Indo-Gangetic plains (Choudhury et al., 2021). Despite the recognized importance of soil microorganisms, their functional diversity across different North Indian sub-regions remains underexplored. Variations in cropping patterns (rice-wheat, maize-sugarcane, pulses-mustard), irrigation water quality, and organic amendments all shape microbial community structure (Gupta & Bhattacharya, 2020).



Modern tools such as metagenomics, stable isotope probing, and geospatial modelling are enabling finer-scale mapping of microbial niches, revealing spatial patterns linked with land use and climatic stressors (Joshi et al., 2022). This review consolidates existing research to provide a comprehensive view of microbial niches across the north Indian alluvial soils. The aims of this study are-

1. Characterize the regional physicochemical and ecological settings.
2. Examine microbial community composition and functional guilds.
3. Analyze case studies from major alluvial tracts.
4. Discuss environmental pressures affecting microbial ecology.
5. Suggest management frameworks for sustainable microbial conservation.
- 6.

By synthesizing field data, experimental findings, and regional comparisons, the study seeks to bridge the gap between microbial ecology and applied soil management, contributing to long-term agricultural sustainability in the Indo-Gangetic plains.

Theoretical Framework: Microbial Ecology and Niche Theory in Alluvial Soils

The study of microbial niches in alluvial soils draws upon the broader concepts of ecological niche theory, biogeography, and microbial functional diversity. A niche represents the range of environmental conditions and resource gradients that allow an organism or population to persist (Hutchinson, 1957). In microbial ecology, this extends to chemical parameters such as pH, organic carbon, redox potential, and micronutrient availability (Fierer, 2017). Each microbial taxon occupies a multidimensional niche defined by its metabolic capacity, symbiotic relationships, and tolerance to stress.

In North Indian alluvial soils, this framework manifests across microhabitats shaped by fluctuating hydrology and soil structure. Floodplain sediments, for instance, foster anaerobic zones where denitrifying bacteria (*Pseudomonas*, *Paracoccus*) dominate, whereas elevated loam ridges support aerobic decomposers (*Bacillus*, *Streptomyces*) (Sharma & Singh, 2020). The coexistence of both niches promotes biogeochemical complementarity, ensuring resilience to environmental disturbance.

Soil texture also regulates microbial niches by influencing oxygen diffusion and substrate retention (Six et al., 2006). Fine-textured clayey soils tend to protect organic matter from rapid mineralization, favoring oligotrophic microbes adapted to low nutrient fluxes. In contrast, coarse sandy alluvium supports copiotrophic organisms that exploit transient nutrient pulses.

Energy Flow and Trophic Networks

Microbial niches are structured by energy flow within soil food webs. Bacteria and fungi act as primary decomposers, transforming detritus into available nutrients. Protists and nematodes then graze upon these microbial populations, creating feedback loops that regulate nutrient mineralization (Ingham et al., 1989). In alluvial systems, periodic flooding redistributes organic matter, creating fresh substrates for heterotrophic growth and enhancing microbial biomass turnover (Ranjan et al., 2022).

Functional Guilds and Ecophysiological Adaptations

Functional guilds within these soils include nitrogen fixers, nitrifiers, denitrifiers, sulfur oxidizers, methanogens, and decomposers. The coexistence of these guilds supports multifunctionality in soil systems (Nannipieri et al., 2017). For example, *Cyanobacteria* and *Azotobacter* colonize paddy soils under microaerophilic conditions, while *Burkholderia* and *Bacillus* dominate dryland wheat fields (Singh et al., 2023). Adaptation mechanisms include sporulation, osmolyte accumulation, and extracellular polymeric substance (EPS) production, which enhance survival under fluctuating water regimes. Fungal hyphae bridge microaggregates, stabilizing soil structure, while actinomycetes secrete antibiotics that maintain microbial equilibrium (Vyas and Subba Rao, 2019).

Regional Overview of North Indian Alluvial Plains

The Indo-Gangetic plains span approximately 700,000 km² and include five major sub-regions—Western (Punjab–Haryana), Central (Uttar Pradesh), Eastern (Bihar–West Bengal), and the Yamuna–Delhi floodplains. Each exhibit distinct edaphic characteristics that shape microbial community dynamics.

Punjab–Haryana Alluvium

These northwestern tracts possess calcareous, moderately alkaline soils (pH 7.8–8.4) with high electrical conductivity due to intensive irrigation (Goyal et al., 2020). Dominant microbial groups include *Pseudomonas fluorescens*, *Rhizobium meliloti*, and phosphate-solubilizing *Bacillus megaterium*. Despite heavy fertilizer use, native microbes retain high enzymatic potential, especially for urease and phosphatase. Crop rotation (rice–wheat) creates alternating anaerobic and aerobic micro-zones, favoring both denitrifiers and decomposers.

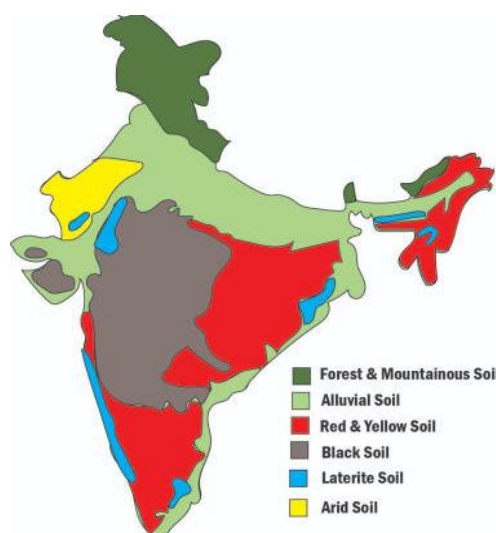


Fig. 1. Map of North Indian alluvial soils along with all other types of soils

Uttar Pradesh Central Plains (Aligarh–Varanasi)

The central Gangetic region contains neutral to slightly alkaline sandy-loam soils rich in silt. Organic matter ranges 0.6–1.2 %, supporting abundant actinomycetes and filamentous fungi (Kumar & Verma, 2021). Seasonal flooding along the Ganga promotes cyanobacterial mats that fix atmospheric nitrogen. In Aligarh, microbial respiration correlates strongly with available phosphorus and carbon fractions (Abbas et al., 2023).

Bihar and Eastern Gangetic Belt

Soils are deep, humid, and clay-rich with high water retention. Anaerobic bacteria, methanogens, and sulfate reducers dominate in rice-growing tracts (Mishra et al., 2020). Regular inundation generates redox oscillations that stimulate denitrification and methane emission. Indigenous microorganisms such as *Methanobacterium formicicum* and *Desulfovibrio desulfuricans* contribute significantly to carbon and sulfur fluxes.

Delhi–Yamuna Flood plain

Urban expansion and pollution have altered microbial balance here. Studies by Kaur and Malik (2022) show reduced microbial biomass carbon near industrial effluent discharge points. Yet rhizosphere samples from riparian vegetation still harbor resilient populations of *Arthrobacter* and *Micrococcus* adapted to heavy-metal stress.

Table 1. Physicochemical properties of representative North Indian alluvial soils (Prepared).

Region	Soil pH	Organic C (%)	Texture	Moisture (%)	EC (ds m ⁻¹)	Dominant Microbes
Punjab–Haryana	7.8–8.4	0.4–0.8	Loam–clay	18–24	0.8–1.6	<i>Pseudomonas</i> , <i>Rhizobium</i>
Central U.P. (Aligarh to Varanasi)	7.2–7.8	0.6–1.2	Sandy loam	16–22	0.5–1.1	<i>Bacillus</i> , <i>Actinomycetes</i>
Bihar Plains	6.5–7.5	0.8–1.5	Clay	25–30	0.4–0.9	<i>Methanobacterium</i> , <i>Desulfovibrio</i>
Delhi–Yamuna	7.6–8.1	0.3–0.7	Loam–sand	14–20	1.2–2.0	<i>Arthrobacter</i> , <i>Micrococcus</i>

Functional Ecology: Biogeochemical Cycles and Microbial Guilds

Microbial niches in alluvial soils play pivotal roles in the global carbon, nitrogen, and phosphorus cycles.

Carbon Cycle

Decomposer fungi (*Aspergillus*, *Trichoderma*) and bacteria (*Bacillus*, *Streptomyces*) catalyze cellulose and lignin degradation, releasing CO₂ and forming humic substances. Flood-induced anoxia stimulates methanogenesis via *Methanosaeta* sp. (Lal et al., 2021). Management practices like residue incorporation enhance microbial biomass carbon and enzymatic activity.

Nitrogen Cycle

Nitrogen fixation by free-living and symbiotic bacteria remains vital in nitrogen-limited soils. Rhizobium–legume associations dominate in pulse systems, while *Azospirillum* and *Beijerinckia* contribute to cereal rhizospheres (Reddy

et al., 2022). Denitrification and nitrification rates are controlled by soil aeration and carbon supply. Excess fertilizer often suppresses diazotroph abundance, underscoring the need for balanced nutrient management.

Phosphorus and Sulfur Cycles

Phosphate-solubilizing microorganisms (PSMs) such as *Bacillus subtilis* release organic acids, converting insoluble phosphates into bioavailable forms. Sulfur oxidizers (*Thiobacillus*) and reducers (*Desulfovibrio*) drive sulfur transformations, particularly in flooded rice soils (Gupta & Chaudhary, 2021).

Table 2. Major microbial functional guilds and their ecosystem roles (Prepared).

Functional Guild	Key Genera	Ecological Function	Representative Region
Nitrogen fixers	<i>Rhizobium</i> , <i>Azospirillum</i>	Atmospheric N ₂ fixation	U.P., Bihar
Phosphate solubilizers	<i>Bacillus</i> , <i>Pseudomonas</i>	Phosphate mobilization	Punjab–Haryana
Methanogens	<i>Methanosaeta</i> , <i>Methanobacterium</i>	CH ₄ production	Bihar, Eastern UP
Decomposers	<i>Aspergillus</i> , <i>Trichoderma</i>	Organic matter breakdown	All regions
Stress-tolerant bacteria	<i>Arthrobacter</i> , <i>Micrococcus</i>	Heavy-metal detoxification	Delhi flood plain

Anthropogenic and Climate Drivers

Anthropogenic pressures including fertilizer overuse, pesticide residues, and industrial pollution have significantly altered microbial diversity in North Indian soils. High nitrate accumulation suppresses symbiotic nitrogen fixation (Pathak et al., 2020). Long-term pesticide exposure reduces fungal diversity, while heavy metals in the Yamuna floodplain promote metal-resistant bacterial consortia (Kaur & Malik, 2022).

Climate change further modulates microbial niches. Increased temperature and irregular monsoon patterns influence moisture regimes and carbon fluxes (Chaudhuri et al., 2023). Droughts enhance sporulation and shift communities toward spore-forming *Bacillus* species, whereas prolonged flooding encourages anaerobic methanogens.

Emerging Tools and Modelling Approaches

Recent advances in metagenomics, 16S rRNA sequencing, and isotopic tracing have revolutionized microbial niche assessment. GIS-based mapping integrates soil physicochemical data with microbial distribution models (Singh & Joshi, 2023). Predictive models simulate how land-use changes affect microbial networks and soil carbon storage. Stable-isotope probing enables identification of active microbial populations involved in key nutrient cycles (Stevenson et al., 2022).

Management and Restoration Perspectives

Sustainable soil management must prioritize microbial health alongside nutrient balance. Practices such as crop rotation, organic amendment, reduced tillage, and biofertilizer use improve microbial biomass and enzymatic activity (Meena et al., 2021). Phytoremediation combined with microbial inoculants offers promise for detoxifying polluted alluvial soils. Community-level training on composting and bioinoculant preparation can aid farmers in reviving microbial fertility without dependence on synthetic inputs. Integrating microbial indicators into soil-health cards could support policy-level adoption (Government of India, 2024).

Conclusion and Future Prospects

Microbial niches across North Indian alluvial soils form the backbone of ecosystem productivity and resilience. Spatial heterogeneity, hydrological variation, and anthropogenic stress collectively define microbial structure and function. Future research should integrate molecular tools with ecological modelling to predict microbial responses to climate and land-use changes. Conservation of microbial diversity is essential for long-term sustainability of the Indo-Gangetic agricultural systems. The inherent spatial heterogeneity of these alluvial landscapes—shaped by riverine deposition, soil texture gradients, and microtopography—creates a mosaic of physicochemical conditions that support diverse microbial assemblages. Variations in moisture regimes driven by monsoonal hydrology, irrigation practices, and groundwater dynamics further influence microbial metabolism and community composition, functionally specialized groups adapted to fluctuating redox and nutrient conditions. However, increasing anthropogenic pressures such as intensive tillage, excessive fertilizer and pesticide use, residue burning, and industrial contamination are altering these finely balanced microbial networks, with implications for soil fertility, carbon sequestration, and crop resilience.

Future research must move beyond descriptive surveys and adopt integrative frameworks that combine high-resolution molecular tools—such as metagenomics, meta-transcriptomics, and stable isotope probing—with ecological and process-based modelling. Such approaches can improve predictions of microbial responses to

climate variability, extreme weather events, and land-use transitions across the Indo-Gangetic Plains. Understanding how microbial functional redundancy and niche complementarity buffer soils against disturbance is critical for designing sustainable management strategies. Conservation of microbial diversity, through practices like reduced chemical inputs, organic amendments, crop diversification, and conservation agriculture, is therefore not merely an ecological concern but a cornerstone for sustaining long-term agricultural productivity and resilience in one of the world's most intensively cultivated regions.

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Author Contributions

MJA and HP conceived the concept, wrote and approved the manuscript.

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Not applicable.



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