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Review

Biofertilizers for Enhanced Nitrogen Use Efficiency: Mechanisms, Innovations, and Challenges

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Abstract

The extensive use of synthetic nitrogen fertilizers has sustained global food production for more than a century but at high environmental and energetic costs. Improving nitrogen use efficiency (NUE) has therefore become a key objective to maintain productivity while reducing the ecological footprint of agriculture. This review synthesizes current knowledge on the biological foundations of NUE enhancement, focusing on the role of microbial biofertilizers and biostimulants. The main mechanisms through which plant-associated microorganisms contribute to nitrogen acquisition and assimilation are analyzed. In parallel, advances in genomics, biotechnology, and formulation science are highlighted as major drivers for the development of next-generation microbial consortia and biobased products. Particular attention is given to the current landscape of commercial biofertilizers and biostimulants, summarizing the principal nitrogen-fixing and plant growth–promoting products available on the market and their agronomic performance. Moreover, major implementation challenges are discussed, including formulation stability and variability in field results. Overall, this review provides an integrated perspective on how biological innovations, market evolution, and agronomic optimization can jointly contribute to more sustainable nitrogen management and reduce dependence on synthetic fertilizers in modern agriculture.

Keywords: nitrogen use efficiency (NUE); microbial biofertilizers; biostimulants; sustainable agriculture

1. Introduction

The extensive use of synthetic nitrogen (N) fertilizers has been the driving force behind the global agricultural intensification of the last century. To secure crop yields and sustain a rapidly expanding population, the Haber-Bosch process, which was commercialized in the early 20th century, allowed for the large-scale industrial fixation of atmospheric nitrogen into ammonia, revolutionizing food production. Approximately half (around 40–50%) of the current world population is sustained by food whose production depends on synthetic nitrogen [1].

Nitrogen is the most common nutrient limiting crop productivity and, after water, represents the second most critical input for plant growth [2]. Nitrogen use efficiency (NUE; the ratio of nitrogen absorbed by plants to the nitrogen applied to crops) remains notably low in most agricultural systems. Cereal crops such as rice, wheat, and maize, which collectively supply the majority of global caloric intake, require high nitrogen inputs to achieve optimal yields [3]. Yet, these systems typically recover less than 50% of applied nitrogen, with the remainder lost through gaseous emissions,

leaching, and runoff [4]. There is an urgent need for innovation because such inefficiency has farreaching consequences, contributing to climate change via nitrous oxide (N_2O) emissions, to eutrophication of aquatic ecosystems through nitrate leaching, and to soil acidification [5].

The environmental and energetic costs associated with synthetic nitrogen fertilizers are increasingly recognized as unsustainable. A large proportion of applied nitrogen is lost as nitrate, contaminating groundwater and promoting eutrophication of rivers, lakes, and coastal zones [6] . Such nutrient overloads contribute to algal blooms, hypoxic zones, and biodiversity loss in aquatic systems. In parallel, nitrogen fertilizers are a major anthropogenic source of nitrous oxide (N₂O), a greenhouse gas with a global warming potential nearly 300 times that of carbon dioxide and a significant contributor to stratospheric ozone depletion [7]. Agricultural soils, particularly those managed with intensive fertilizer regimes, account for approximately 60% of global anthropogenic N₂O emissions [8]. Moreover, the Haber–Bosch process itself consumes about 2% of global energy output and emits approximately 450 Mt of CO₂ annually, further linking fertilizer use to the climate crisis [9]. In addition to its environmental footprint, the production and use of synthetic nitrogen fertilizers are tightly coupled to the global energy market. Because the Haber-Bosch process depends heavily on natural gas as both a feedstock and an energy source, fluctuations in energy prices directly affect the cost of nitrogen fertilizers. Recent geopolitical tensions and energy crises have caused unprecedented increases in natural gas prices, which in turn have driven fertilizer prices to record highs worldwide [10]. This volatility has placed considerable economic pressure on farmers, particularly in developing regions, and has highlighted the vulnerability of modern food systems to external energy shocks. As a result, the high energy intensity and price dependency of nitrogen fertilizer production further reinforce the need for alternative, low-input biological strategies such as biofertilizers and microbial inoculants, which can provide a more resilient and sustainable basis for nitrogen management in agriculture. These impacts underscore the urgent need for a transition toward more sustainable nitrogen management strategies.

Improving NUE is one of the most promising approaches to reduce fertilizer dependency while maintaining high agricultural productivity. Efforts to achieve this goal include breeding crops with enhanced nitrogen uptake and assimilation capacities, optimizing agronomic practices such as precision fertilization, and developing biological alternatives that harness plant–microbe interactions [11]. In this context, microbial inoculants and biofertilizers have emerged as compelling candidates to complement or partially replace synthetic inputs. These biological approaches offer multiple mechanisms to improve NUE: enhancing root architecture, promoting symbiotic or associative nitrogen fixation, modulating phytohormone signalling, and facilitating nutrient mobilization in the rhizosphere [12,13].

The rise of microbial inoculants and biofertilizer innovations has been accelerated by advances in microbiome research, genomics, and synthetic biology. Plant growth-promoting rhizobacteria (PGPR), diazotrophic bacteria, and engineered microbial consortia are increasingly investigated not only for their ability to fix atmospheric nitrogen but also for their roles in regulating stress tolerance and improving nutrient uptake efficiency [14]. Furthermore, mycorrhizal fungi and endophytes contribute to enhanced plant performance under low-input conditions, underscoring the potential of microbial solutions to mitigate dependence on synthetic fertilizers [15]. The commercialization of microbial inoculants has expanded rapidly in recent years, with several biofertilizer products already available in agricultural markets worldwide [16]. Nevertheless, key challenges persist in ensuring consistent field performance, optimizing carrier formulations, and effectively integrating biological solutions into existing fertilization systems.

In parallel with microbial inoculants, biostimulants have gained increasing attention as complementary tools to enhance NUE and overall nutrient performance in crops. Biostimulants comprise a diverse group of substances and microorganisms (including humic and fulvic acids, protein hydrolysates, seaweed and algal extracts, microbial metabolites, and signaling molecules) that act on plant physiology through non-nutritional pathways [17]. Their application can improve nitrogen uptake, assimilation, and remobilization by modulating root growth, enhancing nitrate reductase activity, and stimulating the expression of genes related to nitrogen metabolism. Recent studies have demonstrated that combining biostimulants with reduced nitrogen fertilization regimes

can maintain or even improve yields while lowering N losses [18]. Furthermore, biostimulants often exhibit synergistic effects with microbial biofertilizers, creating integrated formulations that promote both microbial activity and plant responsiveness to nitrogen. This integrated biotechnological approach offers a promising strategy to increase NUE in sustainable cropping systems, particularly under conditions of abiotic stress or limited fertilizer availability [19].

The aim of this review is to synthesize current understanding of the mechanisms underlying NUE enhancement, with particular focus on microbial inoculants and biofertilizer-based strategies. We highlight key field applications, evaluate their agronomic and ecological impacts, and identify critical knowledge gaps that limit large-scale adoption. In addition, we present an overview of current market trends and industrial developments in microbial technologies aimed at improving nutrient use efficiency, emphasizing the rapid expansion of nitrogen-fixing and plant growth-promoting microbial products within the global biofertilizer sector. Finally, we outline future research priorities required to translate microbial innovations into scalable, sustainable solutions capable of reshaping nitrogen management in modern agriculture.

2. Mechanism for Nitrogen Use Efficient Improvement

Nitrogen use efficiency depends on a complex network of interactions among plant physiology, soil chemistry, and microbial activity. Biofertilizers and microbial inoculants can affect several processes that improve nitrogen uptake and retention, helping boost productivity while reducing environmental losses. Key mechanisms include enhanced nutrient mobilization, regulation of nitrogen cycling, stimulation of root development, and the alleviation of abiotic stress effects on nitrogen metabolism (Table 1).

2.1. Nutrient Mobilization: Enhanced Solubilization of Bound Nutrients Improves Nitrogen Uptake Synergy

Microbial inoculants enhance nutrient mobilization in the rhizosphere by secreting organic acids, phosphatases, and siderophores that help solubilize nutrients otherwise unavailable to plants, such as phosphorus, potassium, and various micronutrients [20]. The increased availability of these co-limiting nutrients creates a synergistic effect that improves nitrogen uptake and assimilation within plant tissues. Phosphorus, for example, is essential for adenosine triphosphate (ATP) synthesis and nitrate reduction, and its simultaneous availability supports more efficient nitrogen assimilation pathways [21]. Plant growth-promoting rhizobacteria, such as *Bacillus subtilis*, *Azospirillum brasilense* or *Bradyrhizobium japonicum*, are known to enhance plant nitrogen uptake by upregulating nitrate reductase activity and improving nitrate reduction pathways, thereby facilitating more efficient nitrogen assimilation in plant tissues [22]. By optimizing nutrient stoichiometry within the rhizosphere, these microorganisms indirectly promote greater nitrogen uptake and more efficient utilization for plant growth and yield.

Table 1. Mechanisms of Action, Key Processes, and Microbial Agents Used to Enhance Nitrogen Use Efficiency (NUE) in Crops. The table summarizes the principal processes, representative microbial groups, and the resulting beneficial effects on plant nitrogen uptake and assimilation.

Mechanism	Key Processes	Microbial Agents / Examples	Main Effects on Nitrogen Use Efficiency (NUE)	Reference
	Solubilization of P, K,		Increased availability of	
	and micronutrients	Bacillus subtilis,	co-limiting nutrients (P,	
Nutrient	through secretion of	Azospirillum	K), enhanced nitrate	[20–22]
Mobilization	Mobilization organic acids,	hrasilense	reductase activity,	[20-22]
	phosphatases, and	orustiense	improved nitrogen	
	siderophores		assimilation	

Nutrient cycling regulation	Suppression of nitrification and denitrification; enhancement of Biological Nitrification Inhibition (BNI) and N retention as NH4+	Nitrosomonas inhibition by Brachiaria humidicola, Sorghum spp.; microbial consortia promoting anammox or partial nitrification— denitrification	Reduced N2O emissions, minimized nitrate leaching, improved synchronization of N availability with plant demand	[23,24]
Root Growth Stimulation	Hormonal modulation (auxin, cytokinin, gibberellin), enhanced root hair formation, mycorrhizal associations	Azospirillum brasilense, Bacillus spp., arbuscular mycorrhizal fungi (AMF)	Greater root surface area and soil exploration, improved N absorption and assimilation, enhanced plant vigor	[25–27]
Stress Mitigation	Activation of antioxidant enzymes, osmolyte accumulation, photosynthetic stabilization, stressresponsive gene expression	Bacillus spp., Pseudomonas spp., AMF	Maintained N metabolism under salinity/drought, improved nitrate reductase and glutamine synthetase activity	[28,29]

2.2. Nutrient Cycling Regulation: Suppressing Nitrification or Denitrification Reduces Nitrogen Losses

Regulation of the nitrogen cycle is paramount for enhancing NUE in agricultural systems and mitigating environmental pollution. Nitrification and denitrification are major loss pathways that convert plant-available nitrogen forms (NH_4^+ , NO_3^-) into gaseous emissions (N_2O , N_2). Strategies focused on suppressing these pathways, using synthetic nitrification inhibitors (e.g., DCD) or exploiting Biological Nitrification Inhibition (BNI), are crucial for maintaining N in its less mobile ammonium form (NH_4^+) within the root zone [23] . For example, biological nitrification inhibition (BNI) is a plant-mediated mechanism, observed in species like *Brachiaria humidicola* and sorghum, where root exudates inhibit *Nitrosomonas spp.* activity, thus reducing nitrate leaching and N_2O emissions [24] . Inoculants that promote ammonium retention or facilitate its gradual conversion to nitrate help synchronize nitrogen availability with plant demand, minimizing losses through gaseous emissions or runoff [25]. Moreover, microbial consortia that favour anaerobic ammonium oxidation (anammox) or partial nitrification–denitrification pathways have been proposed as innovative bioengineering solutions for sustainable nitrogen management [26].

2.3. Root Growth Stimulation: Larger and Healthier Roots Improve Nutrient Absorption

A strong root system improves the plant's ability to explore a larger soil volume and access water and nutrients, directly enhancing nitrogen use efficiency. Biofertilizers containing plant growth-promoting rhizobacteria and arbuscular mycorrhizal fungi (AMF) can stimulate root development via modulation of host hormonal balance, particularly via auxin, cytokinin, and gibberellin pathways [27]. A

clear example is *Azospirillum brasilense*, known for synthesizing indole-3-acetic acid (IAA), a hormone that promotes lateral root formation and root hair elongation, thereby increasing the nutrient absorption surface area [28]. Meanwhile, mycorrhizal symbioses expand the root's reach to immobile nutrients and water, indirectly improving nitrogen uptake efficiency under limiting conditions [29]. A more vigorous root system also strengthens the plant's resilience to fluctuations in nitrogen availability, supporting steady assimilation and translocation processes during critical growth stages.

2.4. Stress Mitigation: Biofertilizers Help Plants Cope with Abiotic Stress (Salinity, Drought), Maintaining Nitrogen Metabolism Efficiency

Environmental stresses like salinity, drought, and extreme temperatures often interfere with how plants absorb and process nitrogen, mainly by disrupting enzyme activity and transport functions [30]. Biofertilizers help counter these effects by triggering tolerance mechanisms that allow nitrogen metabolism to continue even under stress. Plant growth–promoting rhizobacteria can boost the activity of antioxidant enzymes, increase the buildup of cell osmolytes (like proline), maintain photosynthetic performance, which keeps key enzymes such as nitrate reductase and glutamine synthetase functioning properly and consistent up-regulation of stress responsive genes [31]. For example, inoculation with PGPR consortia including *Bacillus* and *Pseudomonas* species has been shown to mitigate drought-induced declines in photosynthetic rate and biomass, and increase nitrogen accumulation in shoots, relative to uninoculated controls [32]. Likewise, mycorrhizal fungi help plants cope with salinity stress by improving ion balance and water absorption [33]. Altogether, these microbial inoculants act as biostimulants that help plants withstand environmental fluctuations, maintaining nitrogen use efficiency even under challenging conditions.

3. Biological Nitrogen Fixation

Biological nitrogen fixation (BNF) is a fundamental biogeochemical process by which diazotrophic prokaryotes reduce atmospheric nitrogen (N₂) to ammonia (NH₃), thereby supplying bioavailable nitrogen to plants [34,35]. The reaction is catalysed by the nitrogenase complex, composed of Fe and MoFe proteins, which mediate the energetically demanding six-electron reduction of N₂ [36]. However, the reaction is strongly inhibited by oxygen and requires substantial ATP investment, making BNF both biologically costly and physiologically constrained [37]. Despite these limitations, BNF remains ecologically indispensable, continuously replenishing soil nitrogen pools and supporting the productivity of both natural ecosystems and agricultural systems.

From an agronomic standpoint, BNF reduces reliance on synthetic nitrogen fertilizers, lowering input costs and mitigating environmental risks such as nitrate leaching and greenhouse gas emissions [38]. Yet, terrestrial BNF estimates across spatial and temporal scales remain highly uncertain due to methodological constraints and environmental variability [39]. Recent global analyses suggest that BNF in croplands and cultivated pastures accounts for approximately 56 Tg N annually, representing a 64% increase in terrestrial BNF and a 60% increase in total nitrogen inputs compared to pre-industrial levels [40]. Importantly, its benefits extend beyond legumes: fixed nitrogen can support companion species in intercrops (e.g., soybean—wheat) and contribute residual fertility to subsequent rotations [11,41,42].

BNF occurs across three ecological modes [25]. Symbiotic fixation is the most efficient, involving specialized bacteria such as *Rhizobium* or *Bradyrhizobium* that form root nodules on legumes and channel fixed nitrogen directly to the host [43] (Table 2). Associative fixation involves diazotrophs like *Azospirillum* or *Herbaspirillum* that colonize root surfaces or intercellular spaces of cereals, fixing nitrogen while utilizing host-derived carbon [44]. Free-living fixation occurs in the rhizosphere or bulk soil under microaerobic or anaerobic conditions, where diazotrophs such as *Klebsiella*, *Pseudomonas*, and *Burkholderia* contribute modest but ecologically significant nitrogen inputs. N-fixing bacteria can be found on the surface of plant leaves (epiphytic), in the rhizosphere (close to the plant roots), or as endophytes inside of the plant tissues [45]. In the following sections, we examine recent advances in BNF research and identify key limitations that constrain its agronomic potential mainly focus on symbiotic and associative nitrogen fixation.

Table 2. Representative genera for biological nitrogen fixation, ecological adaptations, and agricultural relevance.

Genus	Representative species	Principal host	Ecological distribution	Agricultural relevance	References
Rhizobium	R. leguminosarum bv. viciae R. leguminosarum bv. phaseoli R. leguminosarum bv. trifolii R. leguminosarum bv. etli	Vetch, peas, lentils Common bean Clover Beans, common bean	Temperate, metal soils Sensitive to acidity, drought Pasture-adapted, cool climates Prevalent in Latin America	European grain legumes Requires inoculation in many regions Important in forage systems Widespread use in smallholder systems	[46–49]
Bradyrhizobium	B. japonicumB. diazoefficiensB. elkaniiB. yuanmingenseB. liaoningenseB. canariense	Soybean Soybean Common bean, soybean Lespedeza spp., soybean Soybean, wild legumes Lupinus, Genista	Acid-tolerant; tropical/subtropical Efficient, microaerobic adaptation Heat- and drought-tolerant Acid-soil adaptation Cold-adapted; found in China Mediterranean legumes	Major soybean inoculant worldwide Widely used in commercial inoculants Valuable in tropical regions Relevant for subtropical forage legumes Soybean cultivation in cooler climates Reforestation and marginal lands	[50–52]
Sinorhizobium	S. meliloti S. fredii	Melilot, alfalfa, fenugreek Soybean, bean, legumes	Alkaline soils; drought tolerance Broad host range; stress-adaptable	Model for molecular symbiosis Nodulates >200 legumes; Asia	[53,54]
Mesorhizobium	M. loti M. huakuii	Lotus, lupin Chinese milk vetch	Acid soils; low fertility East Asia distribution	Model with Lotus japonicus Manure crop in rice systems	[55,56]
Cyanobacteria	Nostoc punctiforme Anabaena azollae (N. azollae) Nostoc commune Nostoc gunnerae (symbiont)	Hornworts, liverworts, Aquatic fern <i>Azolla</i> Free-living associations <i>Gunnera</i> (angiosperm)	Cosmopolitan; oligotrophic soils Aquatic habitats, flooded rice Tolerance to desiccation and UV Tropical wet environments Subtropical to tropical soils	Nitrogen input in pioneer ecosystems Rice systems Nitrogen input in arid ecosystems Ecological model for co-evolution Cycad survival in oligotrophic soils	[57–60]
Azospirillum	A. brasilense	Maize, wheat, rice, sorghum	Rhizosphere colonizer; drought and nutrient stress tolerant	Commercial inoculant	[61]

Herbaspirillum	H. seropedicae	Sugarcane, maize, rice	Endophytic; tropical soils	Improves N uptake in grasses; tested as inoculant for sugarcane	[62]
Methylobacterium	M. oryzae	Rice, wheat	Leaf and root colonizer; utilizes methanol; moderate stress adaptation	Enhances germination, growth, and yield in cereals	[63]
Bacillus	B. subtilis	Maize, soybean, vegetables	Endospore-forming; drought, salinity, and heat tolerant	Biofertilizer/biocontrol agent; improves nutrient uptake and stress resilience	[64]

3.1. Symbiotic: Rhizobium-Legume Interactions, Bradyrhizobium, Sinorhizobium and Cyanobacteria

3.1.1. Rhizobium-Legume System

Symbiotic nitrogen fixation (SNF) represents a central model for mutualistic plant–microbe interactions and stands as a cornerstone of sustainable agriculture [65–67]. In this system, the legume host must tightly regulate nodule development and activity, since nitrogen fixation is highly energy intensive. It consumes large amounts of photosynthate and must therefore be carefully balanced against the plant's overall carbon and energy budget [68]. The interaction is governed by a sophisticated molecular dialogue. Flavonoids secreted by roots activate rhizobial *nod* genes via the regulator NodD, triggering the production of Nod factors that initiate root hair curling, infection thread formation, and cortical cell divisions to form the nodule primordium. In some legumes, additional host signalling is modulated by effectors delivered through a type III secretion system [69,70].

A defining feature of the rhizobium–legume symbiosis is its specificity [71,72]. Typically, each rhizobial strain is compatible with a narrow set of legumes, and conversely, each legume species selectively recruits rhizobial partners [72,73]. Promiscuous strains such as *Sinorhizobium* sp. NGR234 can nodulate more than 230 legume species, whereas others, like *Rhizobium leguminosarum* bv. *viciae*, are compatible with only a limited subset of pea cultivars. When incompatibility arises, it often results in aborted nodules or structures incapable of fixing nitrogen. [74,75]

This specificity carries important agronomic implications. Farmers often inoculate crops such as soybean or peanut with elite rhizobial strains to ensure effective nodulation, particularly in soils where compatible symbionts are absent. Crop rotations with legumes further enrich soil nitrogen for subsequent cereals, frequently contributing tens of kilograms of fixed nitrogen per hectare. However, the success of these practices depends on the precise alignment of host genotype, inoculant strain, and environmental conditions. Soil pH, salinity, temperature, and nutrient availability all influence infection, nodule development, and symbiotic efficiency [76–78]. Because legumes satisfy most of their nitrogen needs through SNF, fertilization is generally restricted to small starter doses applied before sowing, which support seedlings until nodulation is established [79]. Taken together, the rhizobium–legume symbiosis exemplifies a highly coordinated and evolutionarily refined mutualism while simultaneously serves as a practical model for sustainable agriculture, thanks to its ability to naturally enhance soil fertility.

3.1.2. Bradyrhizobium

Within the diversity of rhizobia, the genus *Bradyrhizobium* is notable for its physiological resilience and prominent symbiotic role [80]. A classic example is *Bradyrhizobium japonicum* (and the closely related *B. diazoefficiens*), which are the principal symbionts of soybean (*Glycine max*), one of the world's most important legume crops [80]. In agricultural practice, soybean seeds are routinely inoculated with *Bradyrhizobium* strains to ensure effective nodulation and high nitrogen fixation [81,82]. Besides soybean, different *Bradyrhizobium* strains nodulate groundnut (peanut), cowpea, various tree and shrub [82–87].

These bacteria thrive under environmental stresses that limit other rhizobia [88,89]. For instance, in acidic tropical soils (low pH, high aluminum), indigenous *Bradyrhizobium* outperform introduced rhizobia, nodulating legumes like groundnut more effectively [50,90]. Moreover, inoculation with stress-tolerant *Bradyrhizobium* can help legume crops withstand drought, as seen in pot experiments where inoculated soybeans had higher nodule function and antioxidant enzyme activity during drought stress [91,92]. Their adaptations include production of extracellular polysaccharides that protect cells under drought or acidity and the presence of hopanoid lipids that stabilize membranes [93]. *Bradyrhizobium* stands out among rhizobia for its versatility, it associates with a wide range of legumes (including major crops), withstands harsh soil conditions (acid, drought, heat) making it a reliable inoculant in challenging environments, and even challenges the canonical rules of nodulation

through Nod-independent mechanisms. As climate change and soil degradation intensify, the resilience of *Bradyrhizobium* symbioses will be increasingly valuable, and lessons learned from their adaptability could inform future efforts to enhance symbiotic efficiency in agriculture.

3.1.3. Sinorhizobium

Sinorhizobium (also known as Ensifer) is another major lineage, best represented by S. meliloti in alfalfa. One striking feature of Sinorhizobium is its broad range of compatible hosts across different species [94]. While S. meliloti is relatively specific (effectively nodulating Medicago and a few related genera), other Sinorhizobium species like S. fredii are far more promiscuous [95]. From a comparative genomics viewpoint, Sinorhizobium meliloti has one of the best-characterized pangenomes among rhizobia [96]. Compared with other rhizobia, Sinorhizobium exhibits extensive genetic plasticity. Its genome is multipartite, consisting of a chromosome and large plasmids carrying many symbiotic genes. Core functions are conserved, but accessory plasmid genes vary widely among strains, shaping ecological fitness, stress responses, and host compatibility [97]. This genomic mosaic explains why strains of the same species may differ in nodulation efficiency or competitiveness. For example, some S. meliloti strains possess enhanced systems for iron acquisition or microaerobic respiration, conferring advantages during nodule colonization [96]. Another strain might fix nitrogen more efficiently because it has an enhanced system for microaerobic respiration inside the nodule (like an extra high-affinity cytochrome oxidase). These differences matter for agriculture, as there are variations in symbiotic performance of different rhizobial strains associated with the same host [98]. From an agronomic standpoint, identifying and deploying the most effective strains can significantly enhance yields in forage and grain legumes.

3.1.4. Cyanobacteria

Symbiotic nitrogen fixation is not confined to legume roots; certain non-leguminous plants have forged mutualisms with nitrogen-fixing cyanobacteria (often of the genus Nostoc or closely related genera). Cyanobacteria are photosynthetic, often multicellular (filamentous) bacteria capable of fixing N_2 , and they represent some of Earth's oldest and most self-reliant organisms [99]. These cyanobacteria frequently differentiate heterocysts-specialized, thick-walled cells that create a microoxic environment for nitrogenase activity-allowing N₂ fixation to occur alongside oxygenic photosynthesis [100] . Molecular and phylogenomic studies further show that Nostoc lineages display broad host competence across bryophytes, ferns, gymnosperms, and angiosperms, suggesting conserved symbiotic programs among distant plant taxa [101,102]. Unlike legumerhizobium nodules, plant-cyanobacteria symbioses do not typically involve the formation of a new organ analogous to a nodule (with one notable exception in Gunnera). In Gunnera, Nostoc colonizes stem glands and intracellular channels forming a highly integrated association; in cycads, cyanobacteria inhabit specialized coralloid roots; and in bryophytes such as Anthoceros, colonization occurs in mucilage cavities [103,104]. These associations often take place in pre-existing plant structures (like specialized cavities or glands), and the symbiont often remains less integrated than rhizobia in nodules. Nonetheless, they are true symbioses, the cyanobacteria provide fixed nitrogen to the plant, and the plant offers habitat and nutrients to the cyanobacteria. Three well-known examples of such symbioses are those involving Azolla (water fern), Gunnera (tropical herbaceous plant), and certain bryophytes (primitive plants like hornworts and mosses), as well as the symbiosis in cycads (ancient gymnosperm trees)[105]. Recent reviews reaffirm these systems: (i) Azolla-Anabaena azollae fixes nitrogen at high rates and contributes to sustainable rice systems; (ii) Gunnera-Nostoc punctiforme is a rare angiosperm intracellular symbiosis; (iii) bryophytes and (iv) cycads harbor cyanobionts with a conserved Nostocales core that coevolved with their hosts [101,103,106]. Metagenomic surveys also reveal that coralloid roots of cycads host additional microbial partners that influence cyanobacterial performance and nitrogen fixation efficiency [106].

Cyanobacteria contribute fixed nitrogen while often continuing photosynthesis, providing both carbon and nitrogen benefits to their hosts. Thus, they have even been proposed as biofertilizer

strategies in modern agriculture, inoculating crop fields or orchards with free-living cyanobacteria that could colonize soil or plant surfaces and fix nitrogen in situ [99]. Cyanobacteria-based biofertilization provides a promising sustainable alternative for low-input farming, especially in flooded crops where cyanobacteria naturally thrive. From an agronomic point of view, free-living or associative cyanobacteria such as *Nostoc*, *Anabaena*, *Tolypothrix*, *Calothrix*, and even *Arthrospira* are being developed as biofertilizers. In rice fields, cyanobacterial inoculation or Azolla co-cultivation enhances soil nitrogen availability and improves yields (up to 20% in some trials); in wheat, Anabaena cylindrica-based formulations increased biomass and soil fertility under open-field conditions [107,108]. Beyond flooded systems, cyanobacteria are being applied for soil restoration and carbon sequestration in degraded drylands [99].

Recent advances in formulation technology including alginate-based carriers, polymer encapsulation, and biofilm matrices have improved cell viability, stress tolerance (UV, desiccation), and persistence in the rhizosphere. For instance, alginate composites can act as water-retaining carriers that prolong cyanobacterial survival during establishment in soil [109,110].

Cyanobacteria-based biofertilization provides a promising sustainable alternative for low-input farming, especially in flooded crops where cyanobacteria naturally thrive. However, challenges remain in open-field applications, such as competition with other microbes and maintaining cyanobacterial activity in target areas. Current research highlights that the main bottlenecks are carrier shelf-life, production cost, photodamage, and strain specific adaptation issues that can be addressed using omics-based selection and synthetic biology approaches to design next generation cyanobacterial inoculants [109,111]. Despite these limitations, the potential is considerable: cyanobacterial biofertilizers can enhance soil organic carbon and nitrogen levels while reducing reliance on chemical nitrogen fertilizers.

3.2. Associative: Azospirillum, Herbaspirillum, Methylobacterium, Bacillus

Unlike symbiotic nitrogen fixers or strictly free-living soil diazotrophs, associative nitrogenfixing bacteria establish loose, non-nodulating relationships with plants [112]. These microorganisms colonize the rhizosphere or internal plant tissues (endosphere) of various crops, particularly grasses and non-legumes, without forming specialized nodules. Associative diazotrophs can contribute to the nitrogen requirements of their hosts, providing an advantage under low-fertility conditions [112]. However, their contributions are generally smaller and less direct than those of symbiotic rhizobia. Numerous studies suggest that plant growth promotion by associative bacteria is mediated largely through mechanisms other than atmospheric N2 fixation, such as phytohormone production or nutrient solubilization as it was discussed in section 1 of this review [113,114]. Recent evaluations indicate that evidence for substantial atmospheric N2 fixation by most free-living or endophytic diazotrophs remains limited [115]. To rigorously confirm associative N2 fixation, several criteria must be met: (i) the inoculant bacterium carries a complete set of nif genes required for nitrogenase synthesis; (ii) nitrogenase activity is protected from oxygen inactivation; (iii) the bacterium successfully colonizes plant tissues; (iv) increased respiration linked to N2 fixation is detectable in colonized tissues; (v) plants inoculated and grown under nitrogen-free conditions exhibit enhanced growth and nitrogen accumulation; and (vi) multiple independent methods confirm significant inputs of fixed N₂ under field conditions [115].

3.2.1. Azospirillum: A Model Associative Diazotroph

Azospirillum spp. are Gram-negative, motile bacteria that serve as well-established associative nitrogen fixers in the rhizosphere of cereals and grasses. Although they carry functional nitrogenase enzymes and activate nif gene expression during root colonization, their direct nitrogen contribution to host plants is generally modest compared with that of symbiotic partners. Long-term inoculation studies in maize, wheat, rice, and sorghum consistently report growth and yield enhancement, but these effects are largely attributed to additional plant growth–promoting mechanisms in addition to nitrogen transfer [116]. A defining feature of *Azospirillum*–plant interactions is the stimulation of root

system development through phytohormone production [61]. Some strains also generate nitric oxide signals that enhance root branching or produce ACC deaminase, which lowers stress-induced ethylene levels and further encourages root elongation [61]. Collectively, these responses increase root surface area and thereby improve nutrient and water uptake, indirectly enhancing nitrogen nutrition. Recent evidence shows that seed inoculation with *A. brasilense* Ab-V5 improves maize growth under nitrogen-limiting conditions by modulating biochemical and physiological pathways linked to nutrient assimilation [117]. Similarly, potato inoculation with nitrogen-fixing *Azospirillum* strains has been associated with increased plant nitrogen content, reflecting enhanced nitrogen uptake and utilization efficiency [118]. These effects are consistent with the modulation of transcriptional responses associated with the uptake, assimilation, and efficient use of nitrogen [119].

From an agronomic perspective, *Azospirillum* has been successfully developed as a commercial inoculant in multiple countries. Field trials demonstrate that seed or soil inoculation can boost yields of cereals, particularly under low-nitrogen input systems [120,121]. However, outcomes are variable: responses depend on soil fertility, crop genotype, and environmental conditions. To enhance reliability and consistency, current strategies include co-inoculation with other PGPR or mycorrhizal fungi to synergize growth promotion and nitrogen delivery in the field.

3.2.2. Herbaspirillum: Endophytic Diazotrophs in Grasses

Herbaspirillum spp. are another important group of associative nitrogen fixers, best known for their endophytic colonization of tropical grasses. First identified in sugarcane, Herbaspirillum (e.g., H. seropedicae, H. rubrisubalbicans) colonizes roots, stems, and leaves of crops such as sugarcane, rice, maize, and sorghum [122]. Entry into the plant occurs through cracks at emerging lateral roots or other openings, after which the bacteria spread intercellularly within the cortex and vascular tissues [122]. Carrying nif genes, Herbaspirillum actively fixes nitrogen inside host plants. Along with Gluconacetobacter diazotrophicus, it has been implicated as a major contributor to the high levels of biological nitrogen fixation (BNF) observed in certain Brazilian sugarcane varieties that require little or no fertilizer [62]. In maize field trials, inoculation with H. seropedicae has provided substantial nitrogen inputs; for example, Alves et al. (2021) reported that a peat-based inoculant supplied approximately 30% of the nitrogen demand in inoculated maize [123]. Similarly, inoculation of the rice rhizosphere at the seedling stage with a Herbaspirillum strain significantly increased nitrate nitrogen, ammonium nitrogen, and available phosphorus in the rhizosphere compared with control [124]. These results demonstrate tangible nitrogen savings and yield maintenance through endophytic BNF. Beyond nitrogen fixation, Herbaspirillum promotes plant growth through multiple mechanisms. It produces phytohormones that stimulate root elongation and enhance nutrient uptake, like Azospirillum [125,126]. One challenge for its use as a biofertilizer lies in formulation and shelf life: as a non-spore-forming Gram-negative bacterium, Herbaspirillum is typically delivered fresh, and maintaining viability during storage and under field conditions remains a focus of ongoing research. Nevertheless, field evaluations with Herbaspirillum inoculants have shown promising results [123], underscoring their potential role in sustainable agriculture.

3.2.3. Methylobacterium: Pink Phyllospheric Partners

Methylobacterium spp., often recognized by their pink pigmentation, are ubiquitous facultative methylotrophs that live on plant surfaces and as endophytes. While known mostly for consuming methanol from plant exudates, certain Methylobacterium are also diazotrophic and plant-growth-promoting [127]. They colonize a broad range of hosts (from cereals to trees) and are especially common in the phyllosphere and seeds [128]. Remarkably, Methylobacterium can fix nitrogen in the aerial parts of plants. Madhaiyan et al. (2015) isolated Methylobacterium strains from Jatropha leaves and found ~30% of the leaf endophytes were nifH-positive diazotrophs. One strain, Methylobacterium L2-4, actively fixed N₂ on the leaf surface and inside leaves, improving Jatropha growth and seed yield under low-nitrogen conditions [128]. This was the first report of significant bacterial N-fixation in the phyllosphere, suggesting these microbes may help plants in nutrient-poor environments by

providing a small but critical trickle of nitrogen. In addition to N-fixation, *Methylobacterium* spp. exert various plant growth promoting effects. They are known to produce cytokinins and auxins, and to modulate ethylene levels via ACC deaminase, thereby influencing plant development [129]. In field applications, *Methylobacterium* has gained attention in recent years: foliar sprays of N-fixing *Methylobacterium* are being tested to enhance crop nitrogen use efficiency [130]. Furthermore, a strain *Methylobacterium sp*. 2A was found to alleviate salt stress and even suppress fungal disease in potato, demonstrating bioprotective roles beyond nutrition [131]. While not yet as common in commercial biofertilizers, *Methylobacterium* represents a versatile genus whose multi-faceted plant interactions (from nitrogen fixation to stress mitigation) could be harnessed in future crop inoculant formulations.

3.2.4. Bacillus: Versatile PGPR with Nitrogen-Fixing ability

Species of Bacillus (including related genera such as Paenibacillus) are among the most widely used PGPR in agriculture, valued for their durable spores and broad-spectrum benefits. Traditionally, Bacillus spp. have been recognized for traits such as phosphate solubilization, antibiotic production against pathogens, and the induction of systemic resistance in plants [132]. However, many Bacillus and Paenibacillus strains also carry nitrogenase genes (nif) and are capable of associative nitrogen fixation [64,133]. In addition to nitrogen fixation, Bacillus species promote plant growth through the production of phytohormones [134,135]. Moreover, numerous Bacillus spp. produce ACC deaminase, which helps plants tolerate drought and salinity by reducing stress-related ethylene levels [136]. From an agronomic perspective, Bacillus-based biofertilizers and biopesticides are already widely used. Their ability to form stress-resistant spores gives them an advantage in formulation and storage compared with non-sporulating PGPR. Field trials often report improved root and shoot growth, particularly under stress; for example, B. amyloliquefaciens strain EB2003A enhanced corn and soybean germination under salinity stress [137,138]. Key challenges include ensuring effective root colonization and selecting strains suited to specific crops and soils. Nonetheless, thanks to their combined benefits (modest N2 fixation, hormone production, nutrient solubilization, and stress protection) Bacillus and Paenibacillus remain central to the development of next-generation biofertilizers.

4. Recent Advances in Biofertilizer Development

The last decade has witnessed substantial progress in the design and application of biofertilizers aimed at improving nitrogen use efficiency (NUE). Traditional inoculants based on single microbial strains are progressively being complemented, or even replaced, by more sophisticated approaches that combine microbial diversity, innovative carriers, and biotechnology-driven improvements (Figure 1). These innovations respond to the urgent need for biofertilizers that remain effective across diverse soils, climates, and crop systems.

The successful integration of microbial biofertilizers to enhance Nitrogen Use Efficiency (NUE) hinges entirely on the precise understanding of their underlying mechanisms of action. Without clear mechanistic knowledge, the efficacy of biofertilizer application becomes highly variable, failing to account for critical environmental factors like native soil microbial competition or specific crop physiological needs [139]. Therefore, elucidating the specific pathways involved is crucial for moving away from generic inoculation practices towards a targeted, strain-specific strategy that ensures consistent field results and optimizes the contribution of microbial inputs to sustainable nitrogen management.

4.1. Multifunctional Strains and Microbial Consortia

Beyond individual strains, there is a clear shift from single-strain inoculants to *multi-strain* consortia. Research shows that microbial consortia (combinations of complementary bacteria) often outperform single strains in promoting plant growth [140–142]. Recent efforts have focused on the identification of strains capable of performing multiple functions beyond nitrogen fixation. Certain

Azospirillum, Burkholderia, and Pseudomonas isolates not only fix nitrogen but also produce phytohormones, solubilize phosphorus, and confer tolerance to abiotic stress. The integration of such strains into microbial consortia allows synergistic effects, where complementary species sustain more stable colonization and broaden the range of beneficial functions [141,142]. Co-inoculation strategies, for example combining Rhizobium with plant growth-promoting rhizobacteria (PGPR), have shown to enhance nodulation efficiency and nutrient uptake simultaneously [142]. These findings align with the emerging consensus that multispecies biofertilizers are often more robust and effective under field conditions than single-strain products [140]. However, designing optimal consortia is complex, strains must be compatible and not antagonistic. Recent reviews emphasize the importance of understanding intermicrobial interactions to intentionally design synthetic communities with predictable functions [142]. Advanced strategies are being explored to model and intentionally assemble consortia that deliver multiple plant benefits in tandem. Overall, harnessing multifunctional PGPB and well-coordinated consortia represents a major advance in biofertilizer development, enabling more comprehensive improvements in soil fertility and crop resilience.

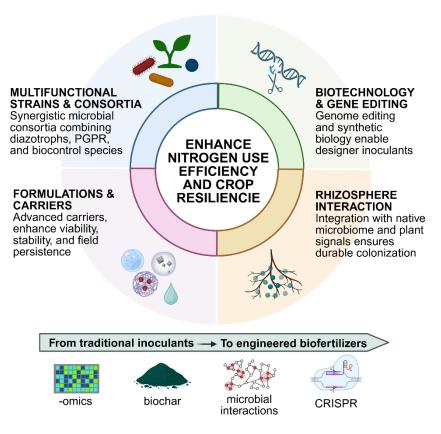


Figure 1. Overview of principal biostimulant trends for improving nutrient use efficiency. Current biostimulant strategies aim to enhance nitrogen nutrition in agriculture through multiple complementary approaches. These include the design of microbial consortia or strain "cocktails" with diverse modes of action, specifically tailored to target crops to improve colonization and effectiveness. Significant efforts are also directed toward the optimization of product formulations to extend shelf life and ensure accurate delivery of active ingredients to plants. In addition, advances in rhizosphere ecology, biotechnology, and gene editing are being leveraged to increase the efficiency and consistency of biostimulant performance under field conditions (*Created in BioRender. Diaz Santos*, E. (2025) https://BioRender.com/3fnfod6).

4.2. Formulation and Carriers

Innovations in biofertilizer formulation technology are improving the viability, shelf stability, and field performance of bacterial inoculants. Traditional carrier-based formulations used peat as a solid carrier for inoculant microbes due to its large surface area, high water-holding capacity, and nutrient content [143]. While peat-based powder inoculants can effectively deliver bacteria (e.g.

coated onto seeds), they typically have limited shelf life (6–12 months) and variable quality [144]. Recent efforts focus on alternative carriers and formulation types that prolong microbial survival. Liquid biofertilizers have gained popularity by suspending bacteria in nutrient-rich broth or oil-based solutions with added protective agents [144]. Unlike dried peat powders, liquid formulations allow inclusion of growth stimulants and osmoprotectants (glycerol, sugars, etc.) to maintain high cell viability. Understanding the complex interactions between carrier materials and microbial communities is essential for optimizing nutrient cycling in agricultural systems. With proper storage (cool, dark conditions), commercial liquid inoculants can achieve shelf lives of 9–13 months [145].

Another major advance is the use of polymeric and novel carriers to protect bacteria from desiccation and stress. Encapsulation of plant growth promoter in natural polymers like alginate, starch, or chitosan can create gel beads that shelter cells and release them slowly into soil [146]. Alginate is a biocompatible, non-toxic matrix that maintains moisture around the microbes and buffers against harsh soil conditions, thereby increasing desiccation resistance and longevity [147]. Additives like skim milk powder, trehalose, and glycerol are often incorporated into capsules to further improve stability by serving as nutrients or stress protectants [148]. Such nanocarrier-enhanced formulations illustrate how coupling beneficial microbes with nanomaterials can improve inoculant efficacy. Biochar-based carriers have also emerged as a promising solid matrix; biochar's porous structure provides microsites for microbial attachment and shields cells from predators and drying [149]. Studies indicate biochar inoculant formulations can extend bacterial survival in soil and even contribute additional soil benefits (e.g. improving nutrient retention).

Furthermore, advanced drying techniques are used to prepare *dry* biofertilizer formulations without compromising viability. Technologies like air-drying, freeze-drying and spray-drying produce powdered inoculants that are easier to handle and apply. To counter viability losses during drying, cryo- or lyoprotectants (e.g. sucrose, mannitol, milk powder) are added to preserve cell membranes [150,151]. Taken together, these formulation advances – whether through optimized liquids, polymer encapsulation, nanoparticle additives, or improved drying methods – have greatly increased the shelf life and field reliability of bacterial biofertilizers. By providing a protective microenvironment, modern carriers ensure more bacteria survive storage and reach plant roots alive, ultimately translating to more consistent plant growth responses in the field.

4.3. Interaction with the Soil and Rhizosphere Microbiota

Understanding the dynamics between introduced inoculants and native microbial communities is central to the success of biofertilizers. Recent research has shown that biofertilizers can modulate rhizosphere microbial networks, either by stimulating beneficial groups or by altering competitive relationships that determine nutrient cycling [152,153]. Some formulations include prebiotic components (such as organic acids or polysaccharides) that selectively enhance the growth of diazotrophs and other beneficial microbes, thereby reinforcing the long-term effectiveness of the inoculant [154]. Biofilm-producing bacteria can anchor to roots and survive fluctuations in moisture or nutrients, thereby outcompeting less resilient microbes [155,156]. Formulation approaches described earlier (e.g. encapsulation in protective polymers or biochar) also enhance persistence by shielding inoculants from desiccation.

The dynamic with the plant itself is equally critical. Plants actively shape their rhizosphere microbiome through root exudation of sugars, amino acids, and other metabolites that selectively recruit beneficial microorganisms [157]. Introduced biofertilizer bacteria must be capable of utilizing these exudates and responding to plant signals to effectively colonize the root zone. Strains that are well-adapted to a crop's exudate profile (sometimes achieved by isolating PGPB from the same crop's rhizosphere) tend to establish more successfully. In turn, a successful inoculant can modulate the native community: studies show that adding beneficial *Bacillus* or *Trichoderma* via biofertilizer can enrich those genera in the soil and even increase the abundance of other advantageous taxa while suppressing some pathogens [158]. Biofertilizer application tends to enhance soil microbial network complexity and stability, meaning beneficial organisms form more robust communities that resist

pathogen invasion [158]. In essence, a well-designed bacterial inoculant does not act in isolation but triggers a cascade of shifts in the soil microbiome that improve overall soil health (e.g. boosting nutrient-cycling microbes or antagonists of plant diseases).

Nevertheless, ensuring that inoculated strains persist long enough to exert their effects remains a challenge. Approaches to improve persistence include repeated or high-dose applications, coinoculating supportive companion microbes, and even breeding plants (or engineering root exudates) that favor the introduced bacteria ("rhizosphere engineering") [159,160]. Some biofertilizer strategies now consider native microbiome compatibility, choosing inoculant strains that complement rather than disrupt indigenous communities [159]. There is also interest in monitoring introduced strains after application using molecular tools to understand their colonization patterns and interactions. Overall, current research recognizes that a biofertilizer's performance hinges on the complex *soil-microbe-plant* interplay. By accounting for factors like competition with native biota, root signaling, and soil physico-chemical conditions, scientists are developing methods to improve colonization (for example, seed coatings that adhere bacteria to roots, or inoculating at a plant growth stage when niches are available). A successful bacterial biofertilizer will integrate into the resident soil microbiome in a way that reinforces beneficial processes without upsetting the ecological balance. The goal is a *persistent*, *self-sustaining population* of the introduced microbes that continues to promote plant growth long after initial inoculation.

4.4. Biotechnology and Gene Editing

Recent advances in biotechnology are revolutionizing bacterial biofertilizer development by enabling precise manipulation of plant growth-promoting bacteri. Omics-guided approaches allow researchers to identify genetic determinants of nitrogen fixation, phosphate solubilization, and stress tolerance, while CRISPR-Cas tools can enhance or delete specific traits such as ACC deaminase production or competing metabolic pathways. Recent studies have successfully applied CRISPR interference (CRISPRi) systems in the model diazotroph Azotobacter vinelandii, enabling targeted modulation of regulatory genes controlling nitrogenase expression and demonstrating precise control of nitrogen fixation under variable environmental conditions [161]. At the same time, omicsdriven analyses of PGPR have revealed key gene clusters related to nitrogen fixation, siderophore production, phytohormone biosynthesis, and oxidative stress protection, which are now being used to guide genome editing and synthetic pathway reconstruction for enhanced field performance [162,163]. Synthetic biology complements this by introducing modular toolkits for constructing or refactoring biosynthetic pathways, as demonstrated in nitrogen-fixing genera like Azotobacter, Pseudomonas stutzeri, and Klebsiella, where inducible nitrogenase circuits and heterologous metabolite pathways have been engineered [163]. In a recent example, a synthetic biology toolkit was established for nitrogen-fixing bacteria (Azotobacter, Pseudomonas stutzeri, Klebsiella), enabling the introduction of regulatory circuits for inducible nitrogenase activity [163]. These innovations illustrate the potential to create "designer" strains optimized for nitrogen fixation, stress resilience, and compatibility with diverse crops. Although regulatory, environmental and safety challenges remain, the integration of multi-omics data, genome editing, and synthetic circuits provides a powerful toolkit for tailoring next-generation biofertilizers with enhanced agronomic performance.

5. Bottlenecks and Challenges

Biological nitrogen fixation offers a sustainable alternative by supplying crops with bioavailable nitrogen naturally. In theory, effective inoculants could reduce fertilizer needs, lowering input costs and mitigating pollution. In practice, however, translating BNF potential into consistent field gains has proven challenging. It is important then, to analyse the major biological, environmental, technological, and regulatory bottlenecks limiting the success of N-fixing bacterial inoculants and biostimulants under field conditions and discusses evidence-based solutions to overcome these hurdles. By addressing these challenges, from strain-host compatibility to formulation and farmer

adoption, we can enhance the contribution of microbial BNF to crop NUE and sustainable agriculture (Figure 2).

Host Specificity and Symbiotic Compatibility: Symbiotic nitrogen-fixing systems are often highly specific. Classical rhizobia form nodules only on compatible legume hosts, with many bacteria and legumes restricted to one another in near one-to-one partnerships [72]. This specificity means an inoculant strain highly effective on one crop may be useless on another. Even with the right pairing, competitive interactions in soil can interfere indigenous rhizobia, if present, often outcompete introduced strains for nodule occupancy, sometimes leading to suboptimal symbiosis despite inoculation. For non-legume crops, the challenge is greater, most cannot form nodules at all, and must rely on looser associations with endophytic or rhizosphere bacteria. Achieving tight integration of N-fixers with non-legume roots remains an unresolved biological hurdle.

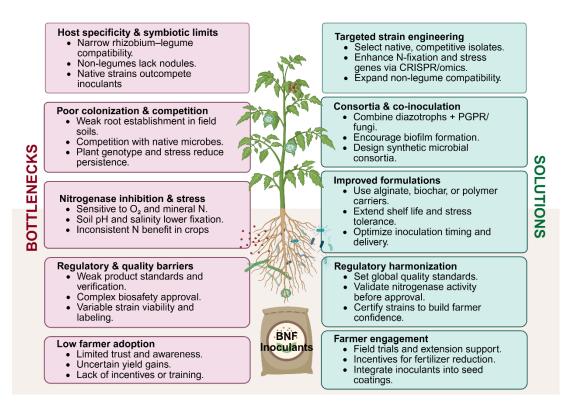


Figure 2. Main gaps and practical solutions for the development of biostimulants focused on improving nitrogen use efficiency in agriculture. The left panel highlights current limitations in large-scale development, including host specificity, regulatory and quality barriers, and low farmer adoption. The right panel presents practical guidelines for product selection and deployment, emphasizing targeted strain development, regulatory harmonization, and farmer engagement. Together, these perspectives illustrate the key challenges and practical considerations required for effective strain selection and successful product implementation (*Created in BioRender. Diaz Santos, E.* (2025)https://BioRender.com/04ldavn).

Colonization Efficiency and Competition: Associative diazotrophs like *Azospirillum* and *Herbaspirillum* can colonize cereals and grasses, but establishing a robust population in the field is inconsistent. These bacteria often provide only a modest fraction of the plant's N needs under real-world conditions. One reason is competition from the native microbiome: the soil and root habitat is crowded with microbes adapted to local conditions. An introduced inoculant may struggle to compete for root attachment sites and nutrients. Crop genotype and exudates also influence colonization; certain cultivars simply recruit inoculants better than others. Moreover, environmental stresses limit colonization. Cyanobacterial biofertilizers (e.g. *Nostoc* or *Anabaena* species used in flooded rice fields) face similar issues: they must remain active against competition from other microbes and not be displaced in open-field conditions.

Nitrogenase Regulation and Environmental Inhibition: A further limitation is that nitrogenase, the enzyme complex for N_2 fixation, is exquisitely sensitive to environmental factors. Oxygen inactivates nitrogenase, so even associative bacteria must find or create micro-anaerobic niches (e.g. inside root tissues or biofilms) to fix N – conditions that may not consistently occur in field soil [115]. Additionally, the presence of mineral nitrogen (ammonium or nitrate) strongly represses nitrogenase gene expression in most diazotrophs, as the microbes downregulate BNF when external N is available. In practical these bacteria tend to benefit crops more via other plant-growth-promoting mechanisms (phytohormone production, etc.) than through N provision [115]. The result is often inconsistent yield respons. Finally, many symbiotic systems are sensitive to soil conditions: acidity, salinity, or temperature extremes can impair infection and nodulation. These environmental stressors frequently limit the field performance of inoculants by disrupting either the microbial partner or the plant's receptivity.

Inoculant Formulation and Viability: Technological challenges begin with formulating microbial inoculants that remain viable and effective from factory to field. Seed-applied inoculants face additional insults: desiccation on the seed surface and exposure to seed-applied chemicals (e.g. fungicide or insecticide treatments) can kill the bacteria. Consequently, a significant bottleneck is ensuring a sufficient live dose of the inoculant reaches the soil and root at planting time.

Biosafety and Regulatory Approval: Bacterial biofertilizers inhabit a gray area between fertilizers and live biological agents, which poses regulatory complexities. In many jurisdictions, standard rhizobial inoculants are lightly regulated (treated as soil amendments), but newer products, especially those with novel strains or genetic modifications, face stricter oversight. Biosafety approval is a barrier for regulation. Native strains may require evidence that they are safe (non-pathogenic to plants, animals, and ecosystems) and not invasive. Currently, regulatory frameworks often lag behind the biofertilizer industry's growth. This can either allow dubious products on the market or, conversely, hinder the introduction of genuinely beneficial innovations.

Quality control is a related issue: absent stringent standards, some commercial inoculants have been found to contain fewer viable bacteria or different strains than claimed. Giller et al. (2025) argue that regulations should require robust evidence of an inoculant's N-fixing efficacy before it is sold, to prevent spurious claims [115]. They suggest criteria such as demonstrating the strain possesses nitrogenase genes and remains present in sufficient numbers throughout the crop life. Implementing such standards would raise product reliability but also increase development costs and time to market.

Farmer Awareness and Adoption: Lastly, socio-economic factors influence the uptake of microbial inoculants. Farmers are understandably cautious about replacing a portion of synthetic N fertilizer with a biological product whose effects may be less predictable. The benefits of inoculation, often a moderate yield increase or fertilizer saving under specific conditions, can be hard to see year-to-year, especially if weather or soil variability masks their impact. In contrast, the cost of failure (yield loss from N deficiency) is high. Therefore, many growers stick to conventional fertilization unless given strong evidence and guidance to adopt biofertilizers. Building farmer trust will require not only better products but also extension efforts, demonstration trials, decision-support tools, and possibly economic incentives (e.g. credits for reducing fertilizer use). In addition, the convenience of application matters: inoculants that integrate seamlessly into farming operations (such as pre-coated seeds or easily applied liquids) are more likely to be adopted. In short, overcoming the human and regulatory hurdles is as important as the biological ones for the success of BNF inoculants.

Despite the persistent challenges that limit the widespread success of bacterial inoculants, interdisciplinary research across microbiology, agronomy, biotechnology, and policy is driving important advances that could transform their role in agriculture and NUE.

One central approach involves the careful selection and genetic improvement of strains, where native isolates adapted to local soil and climate conditions are prioritized because of their competitive advantage and greater resilience, while modern biotechnology, is enabling targeted enhancement of nitrogen fixation capacity, stress tolerance, and host compatibility. Complementary to strain

development, there is growing recognition that single-strain inoculants rarely perform optimally in the complex soil–plant–microbe continuum; thus, microbial consortia combining diazotrophs with other plant growth-promoting rhizobacteria or fungi are increasingly being developed to provide synergistic benefits ranging from improved nodulation and nutrient uptake to enhanced stress resilience, although these multi-component formulations also raise additional challenges for regulatory approval, product quality assurance, and ecological predictability.

At the same time, the role of regulatory frameworks and farmer engagement cannot be overstated: implementing minimum quality standards such as viable cell counts and proof of nitrogen-fixing efficacy, streamlining the approval of well-characterized and non-genetically modified strains, and enforcing measures against spurious or ineffective products will build confidence in biofertilizer markets, while farmer-oriented outreach through on-farm trials, demonstration plots, decision-making guidelines, and integration into certification or subsidy programs will be crucial for mainstream adoption. Evidence from soybean production, where rhizobial inoculation has become routine and highly profitable, shows that when economic, agronomic, and environmental incentives align, farmers readily embrace microbial technologies, and similar success could be replicated in cereals and other non-leguminous crops once consistency and reliability are improved.

Importantly, microbial inoculants need not operate in isolation; integration with non-microbial biostimulants such as humic and fulvic acids, seaweed extracts, and amino acid hydrolysates can generate additional synergies by enhancing soil structure and water retention, stimulating beneficial microbial activity, prolonging the availability of mineral nitrogen in the rhizosphere, chelating micronutrients essential for nitrogenase activity, and directly modulating plant metabolism and stress tolerance. Indeed, humic substances have been shown since the 1970s to improve the growth and nitrogen-fixation efficiency of free-living diazotrophs, while more recent studies highlight their capacity to act as fertilizer synergists by slowing nitrogen losses and improving uptake efficiency. When used together, microbial inoculants can supply nitrogen and phytohormones, while biostimulants enhance root development, microbial colonization, and nutrient assimilation, creating a multilayered system that maximizes NUE. Taken together, these developments illustrate that scaling up the impact of BNF inoculants on global agriculture requires not only scientific innovation in strain engineering, formulation, and consortia design, but also supportive policies, farmercentered extension, and integration with complementary inputs, offering a realistic pathway toward reducing synthetic nitrogen fertilizer dependence by 20-30% in major cropping systems while safeguarding productivity, profitability, and environmental sustainability.

6. Market landscape and Future Perspectives

The positive impact of diazotrophic microorganisms on agriculture has driven rapid growth in the has opened the biofertilizer market. Over the past decade, the global biofertilizer market has expanded considerably, reflecting increasing demand for sustainable alternatives to chemical fertilizers. Numerous nitrogen-fixing microorganisms, such as *Rhizobium*, *Azospirillum*, and *Azotobacter* are now commercialized as biofertilizer products (Table 3).

Table 3. Overview of commercial biofertilizer products designed to enhance nitrogen use efficiency (NUE). The table summarizes representative formulations, their declared microorganisms, concentration, target crops, and manufacturing companies.

Name	Microorga	Conce	Crop (s)	Compa	Reference
	nism	ntra.		ny	
Pivot	Kosakonia	≥ 1x10 ⁹	Maize	Pivot	https://www.pivotbio.com/product-
Bio	sacchari 6-	CFU	(Corn)	Bio Inc	proven40-corn (accessed on 27/10/2025)
PROVE	5687 (Ks6-	g ⁻¹		(USA)	

N®40	5687)				
OS	/Klebsiella				
	variicola				
	137-2253				
	(Kv137-				
	2253)				
Pivot	Klebsiella	4x10 ⁸	Wheat	Pivot	https://www.pivotbio.com/product-return-
Bio	variicola	CFU		Bio Inc	wheat (accessed on 27/10/2025)
RETUR	137-1036	g^{-1}		(USA)	
N®	(Kv137-				
	1036)				
Pivot	Kosakonia	≥ 1x10 ⁹	Cotton	Pivot	https://www.pivotbio.com/product-cert-n-
Bio	sacchari 6-	CFU		Bio Inc	cotton (accessed on 27/10/2025)
CERT-	5687 (Ks6-	g^{-1}		(USA)	
$N^{\scriptscriptstyle TM}$	5687) /				
	Klebsiella				
	variicola				
	137-2253				
	(Kv137-				
	2253)				
Always-	Proprietar	Not	Row Crops	BioCons	https://bioconsortia.com/bioconsortia-
N^{TM} /	у	declare		ortia	expands-nitrogen-fixing-microbial-products-
FixiN 33	microbial			Inc.	for-row-crops/ (accessed on 27/10/2025)
	consortiu			(USA /	
	m (N-			NZ)	
	fixing)				
BlueN®	Methylobac	$3x10^{7}$	Row	Corteva	https://www.corteva.co.uk/products-and-
/Utrisha	terium	CFU	crops,Hortic	Agriscie	solutions/biologicals/bluen.html (accessed on
$^{\circledR}N$	symbioticu	g^{-1}	ultural,	nce /	27/10/2025)
	m (SB23)		Olives and	Symbor	
			Grapes	g	
Nutri-	Azotobacter	Not	Broadacre &	Nutri-	https://nutri-tech.com.au/products/bio-n
Life Bio-	vinelandii	declare	horticultural	Tech	(accessed on 27/10/2025)
$N^{\scriptscriptstyle TM}$		d	crops	Solution	
				S	
				(Austral	
				ia)	
Life	Azotobacter	$1x10^{8}$	Broadacre &	Nutri-	https://regenagsolutions.ca/biologicals/life-
Force	chroococcu	CFU	horticultural	Tech	force-bio-n/ (accessed on 27/10/2025)
$Bio\text{-}N^{\scriptscriptstyle TM}$	m	ml ⁻¹	crops	Solution	
				S	
				(Austral	
				ia)	

AZOFIX	Azotobacter	1x10 ⁹	Grassland,	UAB	https://vatzum.lrv.lt/media/viesa/saugykla/20
®	vinelandii	CFU	arable, Fruit	Bioener	24/2/xMNuAnrcjXA.pdf (accessed on
	MVY-010	ml^{-1}	and	gy LT	27/10/2025)
			vegetables		
Azofix	Paenibacill	1.2x10 ⁹	Grassland,	UAB	https://www.bioenergy.lt/en/product/azofix-
Plus	us	CFU	arable, Fruit	Bioener	plus/ (accessed on 27/10/2025)
	polymyxa	ml ⁻¹	and	gy LT	
	MVY-024		vegetables		
N-	Methylobac	1.2x10 ⁹	Grassland,	UAB	https://vatzum.lrv.lt/media/viesa/saugykla/20
FOLIAR	terium	CFU	arable, Fruit	Bioener	24/2/xMNuAnrcjXA.pdf (accessed on
	phyllosphae	ml ⁻¹	and	gy LT	27/10/2025)
	rae MVY-		vegetables		
	033				
	Azotobacter	1x10 ⁷	Grassland,	Syngent	https://www.syngenta.co.uk/product/crop-
Nientris	salinestris	CFU	arable, Fruit	a	protection/biostimulant/nientris (accessed on
®	strain	g^{-1}	and		27/10/2025)
	CECT 9690	1x10 ⁷	vegetables		
	Wickerham	CFU			
	omyces	g^{-1}			
	anomalus				
	strain				
	CECT				
	13172				
Accolad	Azospirillu	1x10 ⁹	Row Crops	Verdesi	https://vlsci.com/products/accolade/
e®	m	CFU		an	(accessed on 27/10/2025)
	brasilense	g^{-1}			
NEOFO		1x10 ⁸	Grassland,	Fertiberi	https://www.fertiberia.com/en/products/agric
RCE N	Azospirillu	CFU	arable, Fruit	a	ulture/neoforce-n-fixer/ (accessed on
FIXER	m	ml ⁻¹	and		27/10/2025)
	brasilense	$1x10^{8}$	vegetables		
		CFU			
	Azotobacter	ml ⁻¹			
	vinelandii	$1x10^{8}$			
	Rhizobium	CFU			
	leguminosa	ml ⁻¹			
	rum				
N-Leaf	Methylobac	1x10 ⁹	Grassland,	De	https://desangosse.com.au/produit/n-leaf/
	terium	CFU	arable, Fruit	Sangoss	(accessed on 27/10/2025)
	brachiatum	g ⁻¹	and	e	
	AGN12	1x10 ⁹	vegetables		
	Methylobac	CFU			
	terium	g ⁻¹			
	pseudosasic	1x10 ⁹			
	•				

	Arthrobacte	CFU			
	r	g^{-1}			
	globiformis				
	AGN14				
Simbioti	Bacillus	2x10 ⁸	Grassland,	CERES	https://ceresbiotics.com/en/products/simbioti
c®	megaterium	CFU	arable, Fruit	Biotics	c/ (accessed on 27/10/2025)
	CECT	g ⁻¹	and		
	30994		vegetables		
Symbius	Bacillus	4.853×	Horticultura	Nostoc	https://nostoc.es/producto/fertilizante-
	altitudinis	10^{10}	l and woody	Biotech	simbius/ (accessed on 27/10/2025)
	strain	CFU	crops		
	NTC/BC/0	ml ⁻¹			
	1				
Amylis	Bacillus	5x10 ⁸	Horticultura	De	https://www.desangosse.co.nz/produit/amyli
	amyloliquef	CFU	l and woody	Sangoss	s/ (accessed on 27/10/2025)
	aciens cepa	ml^{-1}	crops	e	
	I4995	5x10 ⁸			
	Bacillus	CFU			
	amyloliquef	ml^{-1}			
	aciens				
	strainI4996				
Cell-	Bradyrhizo	1x10 ¹⁰		Narrana	https://www.novonesis.com/en/biosolutions/
Tech®	bium	CFU	Soybean	Novone	animal-and-plant/plant/soybeans/cell-tech-xc
XC	japonicum	ml^{-1}		sis	(accessed on 27/10/2025)
LALFIX	D1.:1.:			T -11	
®	Rhizobium	8x10 ⁸	Pea, lentil,	Lallema	https://www.lallemandplantcare.com/en/usa/
LIQUID	leguminosa	CFU	and faba	nd	products/lalfix-liquid-pea-lentil/ (accessed on
Pea &	rum biovar	ml ⁻¹	bean	PLant	27/10/2025)
Lentil	viciae			Care	
NI adoler	Bradyrhizo				
Nodulat or® PRO / IP Plus	bium	$1x10^{10}$			https://agriculture.basf.ca/east/en/products/so
	јаропісит	CFU/	Soybean	BASF	lutions/nodulator-ip-plus.html (accessed on
	(strain	ml			27/10/2025)
	532C)				

Several formulations have demonstrated strong potential to improve crop growth, yield, and soil health while reducing reliance on synthetic nitrogen inputs and, most importantly, lowering farmers' fertilizer costs In few years the biofertilizer market has grown and at present, many nitrogen-fixing microorganisms are marketed as biofertilizers. Different products are available and some of them have shown great potential by improving crop growth and yield and could significantly reduce a farmer's fertilizer bill [164].

Industry analyses estimate that the nitrogen-fixing biofertilizer segment reached approximately USD 1.03 billion in 2024, with projected annual growth rates of 12–13 % through 2030, reflecting the rapid maturation of this niche within the broader biofertilizer market (GrandViewResearch,2024; https://www.grandviewresearch.com/industry-analysis/biofertilizers-industry ; accessed on the

27/10/2025). Economically, Brazil represents a benchmark case: the partial substitution of urea by biological nitrogen fixation in soybean cultivation generated an estimated USD 15.2 billion in savings during the 2019-2020 season, underscoring the macroeconomic potential of large-scale adoption [165]. India is also emerging as a key growth market, supported by government incentives and public programs, with double-digit expansion expected over the next decade (IMARC Group, 2024; https://www.imarcgroup.com/india-biofertilizer-market, accessed on the 27/10/2025). In the European Union, nitrogen-fixing biofertilizers represent one of the fastest-growing product categories within the biologicals sector, driven by environmental policies promoting the reduction of synthetic nitrogen fertilizers and the transition toward regenerative agricultural practices. The European market for biofertilizers was valued at approximately USD 679 million in 2024, with projections reaching USD 1.1-2.1 billion by 2030-2033, and nitrogen-fixation products accounting for the largest revenue share (Mordor Intelligence, 2024; https://www.mordorintelligence.com/industryreports/europe-biofertilizers-market; accessed on the 27/10/25; IMARC Group, https://www.imarcgroup.com/europe-biofertilizer-market; accessed on the 27/10/25; Grand View Research, 2024; https://www.grandviewresearch.com/horizon/outlook/biofertilizers-market/europ; accessed on the 27/10/25).

Current commercial formulations encompass symbiotic rhizobia, free-living diazotrophs that colonize plants endophytically or epiphytically within the rhizosphere or phyllosphere, and coinoculants combining multiple microbial functional groups. These products appear in both solid and liquid formulations, typically maintaining viable counts above 10^6 CFU g⁻¹ or mL⁻¹ and ensuring high microbial viability. Many formulations now include bioactive compounds that enhance propagule activation and rapid plant–microbe interaction after application.

On the symbiotic side, companies such as Rizobacter (Argentina) continue to lead with traditional rhizobial inoculants for legumes, where *Rhizobium* or *Bradyrhizobium* species are applied at planting or as seed treatments to induce nodulation and supply biologically fixed nitrogen.

On the associative and free-living side, both microbial diversity and crop scope are significantly broader. Unlike conventional rhizobial inoculants limited to legumes, these products target cereals, oilseeds, and horticultural crops by exploiting microorganisms capable of colonizing roots, rhizosphere, or internal plant tissues without nodule formation. Among established commercial examples, Gluconacetobacter diazotrophicus is employed in Envita® (Azotic Technologies), an endophytic inoculant applied as foliar or in-furrow treatment that supplies biologically fixed nitrogen throughout the crop cycle (Azotic Technologies, 2024). Similarly, BlueN™ (also marketed as Utrisha N by Corteva Agriscience) contains Methylobacterium symbioticum SB23, a phyllosphere endophyte that converts atmospheric N₂ into ammonium directly within leaf tissues, designed to replace a fraction of mineral fertilization in high-input systems (Corteva Agriscience, 2023). Nutribion® N (marketed as Vixeran® in some European regions) by Syngenta Biologicals is based on Azotobacter salinestris CECT 9690, a versatile diazotroph capable of both epiphytic and rhizospheric colonization, enhancing nitrogen assimilation and stress tolerance in cereals and vegetables (Syngenta Biologicals, 2023). The most innovative frontier within this segment is represented by Pivot Bio's PROVEN® 40, which integrates proprietary engineered diazotrophs—Kosakonia sacchari 6-5687 (Ks6-5687) and Klebsiella variicola 137-2253 (Kv137-2253)—designed to stably colonize maize roots and deliver consistent, measurable nitrogen through biological fixation. Company field data indicate that microbial N delivery via PROVEN® 40 can replace up to 40 lb N acre-1 while maintaining yields (Pivot Bio, 2024). Beyond its agronomic function, this product embodies the shift toward strain engineering, on-seed delivery, and digital traceability, illustrating the convergence of biotechnology and precision nutrient management.

Despite these encouraging outcomes, independent assessments consistently show that the performance of associative and free-living nitrogen-fixing inoculants remains highly variable across environments. Yield responses depend strongly on soil nutrient status, existing microbial communities, and climatic conditions [166]. According to the field study by Arrobas, Correia, and Rodrigues [167] conducted on lettuce, the commercial product BlueNTM (based on *Methylobacterium*

symbioticum) showed limited or non-significant effects on nitrogen fixation and plant biomass accumulation under open-field conditions, indicating a strong dependence on environmental and management factors. Similarly, reviews of *Azotobacter spp.* emphasize that agronomic responses are highly context-specific and influenced by strain compatibility, soil fertility, and stress tolerance [168].

As Giller et al.[169] critically argue in Science Losing Its Way: Examples from the Realm of Microbial N₂-Fixation in Cereals and Other Non-Legumes, there is no unequivocal evidence that current microbial inoculants enable non-leguminous crops to fix agriculturally meaningful quantities of atmospheric nitrogen. The authors emphasize that much of the existing literature is based on weak experimental design, non-replicated studies, or indirect proxies of nitrogen fixation, making it difficult to quantify true N contributions at field scale. Moving forward, progress in associative and free-living diazotroph technologies will depend on reconciling scientific rigor, regulatory oversight, and industrial credibility. Establishing a unified evidence framework—integrating isotopic tracing, multi-environment field trials, and nutrient-balance modeling—is essential to distinguish truly nitrogen-fixing inoculants from general plant-growth-promoting formulations. Only through transparent, science-based evaluation can the industry move beyond anecdotal claims and deliver microbial nitrogen solutions that are both agronomically relevant and scientifically defensible.

7. Conclusions

Microbial biofertilizers and biostimulants represent powerful, nature-based solutions for improving nitrogen use efficiency and reducing dependency on synthetic fertilizers. Significant advances in microbial genomics, synthetic biology, and formulation science are transforming the development of inoculants from empirical tools into precision-designed agricultural technologies. However, their large-scale adoption remains constrained by variability in field performance, lack of regulatory harmonization, and limited farmer awareness. Integrating microbial solutions with biostimulants, precision fertilization, and crop-specific management can bridge these gaps and deliver more consistent outcomes. Recent progress in multi-omics approaches (metagenomics, transcriptomics, metabolomics, and phenomics) is allowing a deeper understanding of the molecular dialogue between plants and beneficial microbes, revealing key regulatory networks that determine nutrient acquisition efficiency and stress resilience. Likewise, optimizing carrier materials, microencapsulation techniques, and co-formulation strategies with organic amendments or signaling molecules is significantly enhancing the viability, stability, and persistence of microbial inoculants across diverse soil types and climatic conditions. Future efforts should focus not only on understanding plant-microbe-soil interactions under real agricultural conditions but also on integrating this knowledge into predictive models and digital agriculture frameworks that support site-specific management. Establishing standardized evaluation protocols and harmonized regulatory frameworks across regions will be essential to ensure product quality, traceability, and farmer confidence. Furthermore, fostering interdisciplinary collaborations among microbiologists, agronomists, soil scientists, will accelerate the transition toward a new generation of microbial-based fertilizers that are both effective and environmentally safe. Ultimately, enhancing NUE through biofertilizers and biostimulants offers not only a path toward more sustainable agricultural systems but also a strategic response to the rising costs of fertilizers, energy dependence, and the environmental footprint of global nitrogen use. In this context, microbial technologies stand out as one of the most promising tools for building more resilient and efficient food systems. By working with nature rather than against it, they help restore soil vitality, reduce environmental pressures, and make crop production less dependent on external inputs. Strengthening their development and practical implementation will be key to achieving a sustainable balance between productivity, profitability, and environmental care.

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