

Review

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Posted Date: 3 March 2025

doi: 10.20944/preprints202503.0009.v1

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Review

# Beneficial Microbes in Seed Science

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**Abstract:** Seed science has undergone transformative advancements through the integration of microbial technologies, with beneficial microorganisms emerging as critical tools for enhancing germination, seedling vigor, and crop resilience. Research demonstrates that microbial treatments improve nutrient uptake, hormonal regulation, and stress tolerance while establishing early symbiotic relationships with plants. This review synthesized recent advances in understanding the roles of beneficial microbes in seed science, focusing on their impact on seed germination, seedling growth, and plant health. We explored the composition and transmission of seed microbiomes, highlighting the vertical transfer of microbes from parent plants to seeds and the influence of environmental factors on microbial community structure. The review also discussed innovative approaches to seed microbiome engineering. Particular attention was given to seed biopriming with plant growth-promoting bacteria (PGPB), which has shown significant potential in improving germination rates, seedling vigor, and crop productivity. We examined specific microbial strains, such as *Trichoderma* species and *Pseudomonas fluorescens*, and their mechanisms of action in enhancing plant performance. The review also addressed the impact of breeding on seed microbiomes and explored emerging research directions, including the development of tailored microbial inoculants and the investigation of intracellular seed bacteria. By synthesizing these findings, this review aimed to provide a comprehensive summary of the current state of seed microbiome research and its implications in seed science for sustainable agriculture.

**Keywords:** seed microbiome; beneficial microbes; crop performance; vertical transmission; biopriming; plant growth-promoting bacteria (PGPB); sustainable agriculture; SynComs

## 1. Introduction

The seed microbiome has emerged as a crucial frontier in agricultural research, offering promising avenues for enhancing crop performance and sustainability. Recent advances in seed science, particularly in the realm of beneficial microbes, have revealed the significant impact of seed-associated microorganisms on plant health, growth, and productivity. These microbes—including bacteria, fungi, and cyanobacteria—interact with seeds through biopriming, coating, and inoculation strategies, offering sustainable alternatives to chemical inputs.

Seeds shelter diverse microbial communities, collectively termed the seed microbiome, which play vital duty in seed germination, seedling establishment, and plant development [1]. These microbial assemblages, comprising bacteria, fungi, and archaea, are unique among plant-associated microbiomes and have been shown to contain primarily beneficial microorganisms [2]. The seed microbiome serves as a critical link between plant generations, facilitating the vertical transmission of microbial resources from parent plants to offspring [3].

The importance of seed microbiomes extends beyond their role in plant development. They also contribute significantly to crop resilience against biotic and abiotic inflections, disease resistance, and overall plant fitness [2]. Furthermore, seed microbiomes have implications for grain quality and food security, making them a subject of intense research interest in recent years [1]. Studies have revealed that seed microbiomes are influenced by plant genotype, environmental factors, and developmental stages [1]. The vertical transmission of microbes from parent plants to seeds has been observed, suggesting a mechanism for passing beneficial microbes to offspring [4].

Seed-associated beneficial microbes have been shown to support seed germination, enhance seedling vigor, increase nutrient uptake, and improve plant resilience to various stresses [5]. Plant growth-promoting bacteria (PGPB) and fungi have also demonstrated their abilities to increase crop production and improve plant performance under adverse conditions [6,7]. Innovative approaches to manipulate seed microbiomes, such as the endophytic microbial introduction (EMI) [8] and synthetic microbial communities (SynComs) [9], are being developed to enhance crop performance [4]. Seed biopriming with beneficial microorganisms has also shown significant potential in improving germination rates, seedling vigor, and crop productivity [6].

The use of beneficial seed microbes offers an environmentally friendly alternative to chemical fungicides and fertilizers, aligning with the goals of sustainable agriculture [7]. These microorganisms can reduce the need for synthetic inputs while promoting natural disease suppression and nutrient cycling [10]. Research has identified various beneficial microbial strains, including species of *Bacillus*, *Pseudomonas*, and *Trichoderma*, which have shown auspicious results in enhancing plant growth and stress tolerance [11,12].

As our understanding of seed microbiomes continues to grow, so does the potential for harnessing these microbial communities to improve crop production, enhance food security, and promote sustainable agricultural practices.

## 2. Seed Microbiome Composition and Transmission

Recent research has noticeably advanced our knowledge of seed microbiome composition and transmission, revealing the complex and dynamic nature of microbial communities associated with seeds.

Seed microbiomes, composed of bacteria, fungi, and archaea, are crucial for plant health and development. They form a unique microbial community distinct from other plant-associated microbiomes [2]. Seed microbiota composition varies widely, but most seeds share a core microbiome [13]. Individual seeds are often dominated by a single bacterial taxon (>75% of reads), with high variability between and within plants [14].

Vertical transmission of microbes from parent plants to seeds is a critical process in seed microbiome assembly. Parent seed and stem endosphere fungal and bacterial communities are key sources of progeny seed microbiomes [15]. Seed-transmitted fungi and bacteria dominate juvenile crop plant microbiomes by abundance [16]. Vertical transmission allows beneficial microbes to establish early founder populations, shaping the plant microbiome from the start through priority effects [16].

Several factors contribute to the composition and transmission of seed microbiomes. Plant genotype, environmental factors, and developmental stages influence seed microbiome composition [15]. Selection is the key ecological process driving dominant taxa succession during seed filling and maturation [14]. Storage conditions can affect seed microbiota conservation, with initial seed drying before storage reducing microbial composition [4].

Seed microbiome assembly and transmission involve complex processes. Abundance-based models classify microbes, with many late colonizers dominating at ripening. Temporal patterns are shaped by niche changes and neutrality [15]. The transition from seed to seedling involves significant changes in microbial population sizes and community structure [14].

Understanding seed microbiomes is crucial as seed-associated microbes support germination, protect against pathogens, and enhance seedling nutrition and vigor [16]. The seed microbiome links

the maternal and offspring environments, influencing plant ecology and evolution [4]. Manipulating seed microbiomes offers potential for improving crop establishment and developing microbial-based solutions for agriculture [4,14,17].

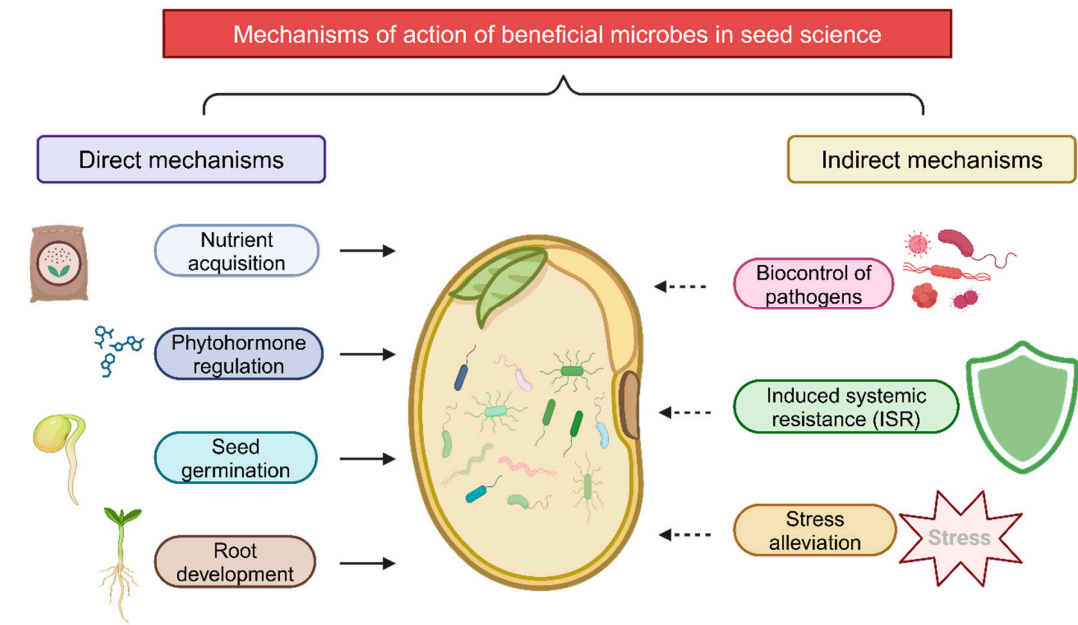
In other words, recent research has revealed the complexity of seed microbiome composition and transmission, highlighting its importance in plant health and agricultural practices. Further studies are needed to fully elucidate the mechanisms of microbial inheritance and their implications for sustainable agriculture.

3. Beneficial Effects of Seed-Associated Microbes

Seed microbes are vital for plant health, growth, and productivity. Recent research has revealed several key benefits of these microorganisms. Seed-associated microbes significantly improve germination rates and seedling vigor. Beneficial microbes boost germination, seedling vigor, biomass, and help overcome seed-related stresses during and after emergence [5,18–20].

Beneficial microbes initiate complex biochemical interactions during seed germination, priming metabolic pathways to enhance vigor and uniformity. Cyanobacteria such as *Spirulina platensis* enhance photosynthetic capacity in emerging seedlings by upregulating chlorophyll synthesis genes, particularly under cadmium stress [5]. In a study on mung bean seeds, the addition of cultured microbes from domestic soil significantly enhanced germination frequency compared to controls [21].

Seed microbes enhance plant growth, with PGPB boosting crop yields by 12-20% (Figure 1)[6]. Seed endophytic bacteria enhance crop growth and yields in plants like rice, maize, wheat, and tomatoes [22]. Microbes produce phytohormones like IAA, gibberellins, and cytokinin, boosting root growth, biomass, and plant development [22]. For example, *Bacillus subtilis* and *Pseudomonas fluorescens* secrete hydrolytic enzymes such as  $\alpha$ -amylase and invertase, which mobilize stored carbohydrates in seeds, providing energy for radicle emergence [5,23]. These bacteria produce phytohormones like IAA, promoting cell elongation and lateral root growth in wheat and maize seedlings [6].



**Figure 1.** Mechanisms of action of beneficial microbes in seed science. Beneficial microbes play a crucial role in seed science by enhancing plant growth, resilience, and productivity through diverse direct and indirect mechanisms. Their application in agricultural practices, such as seed treatments or microbiome engineering, provides sustainable solutions for improving crop performance under variable environmental conditions. Modified from [7,24].Created with BioRender.com (accessed on 1 January 2025).

Seed-associated microbes can also improve nutrient uptake and availability. Some microbes, like mycorrhizal fungi, increase the surface area of plant roots and improve nutrient uptake by creating an extensive mycelial network [25]. Certain bacteria can even solubilize phosphorus, potassium, and zinc, making these nutrients more available to plants [6].

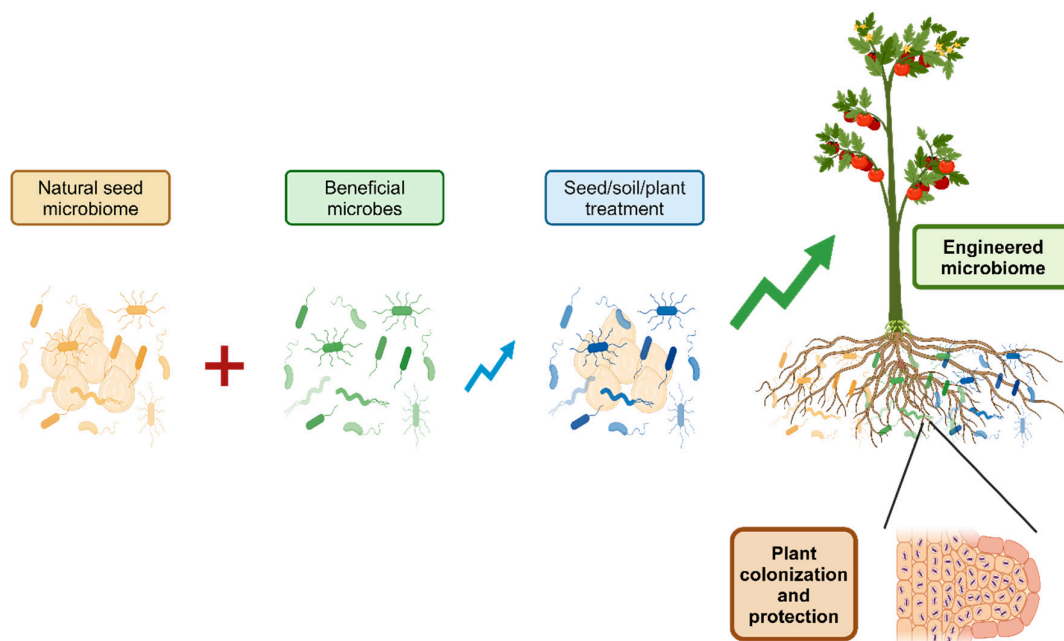
Seed microbes enhance plant resilience to environmental stresses like drought and salinity [7,22,26,27]. Seed biopriming with beneficial microorganisms increases plant resilience and effectiveness under adverse conditions [6].

Seed-associated microbes also play a crucial role in protecting plants from different pathogens. Some microbes produce compounds that inhibit the growth of harmful pathogens in the soil, reducing the risk of disease [18,19,26,27]. Seed biopriming offers eco-friendly biotic stress management, serving as an alternative to chemical fungicides [6,7,18]. The seed microbiome influences plant ecology and evolution by enhancing nutrient uptake, pathogen resilience, and abiotic stress tolerance. It also aids plant establishment, colonization, and spread, offering insights into conservation and invasion ecology [28].

In short, seed-associated microbes offer numerous benefits to plants, from improved germination and growth to enhanced stress tolerance and disease resistance. Understanding and harnessing these beneficial microbes can contribute significantly to sustainable agriculture and plant ecology.

#### 4. Innovative Approaches to Seed Microbiome Engineering

Recent advances in seed microbiome engineering have opened new possibilities for enhancing crop performance and sustainability (Figure 2).



**Figure 2.** Seed microbiome engineering. Seed microbial communities, shaped by genotype, environment, and management, are vital for sustainable agriculture. Modified from [19,26,29]. Created with BioRender.com (accessed on 1 January 2025).

SynComs offer a promising method to engineer plant microbiota. A study in *FEMS Microbiology Ecology* showed that seed inoculation with SynComs effectively shaped seedling microbiota, with SynComs comprising 80% of the community. Strain abundance on seeds was key to successful colonization [9].

A new framework integrates top-down and bottom-up strategies to engineer natural microbiomes. Using herbicides and degrader inoculation, researchers guided microbiomes toward enhanced bioremediation. They also developed SuperCC, a metabolic modeling tool, to analyze interactions and predict microbiome performance [30].

Recent research has focused on harnessing seed-associated endophytes for improved crop performance. A 2024 study used genome-wide association studies (GWAS) to identify genetic loci linked to bacterial seed endophyte diversity in fonio [31]. The results suggested that Seed endophytes promote plant growth by enhancing nutrient availability and uptake [31].

Understanding and manipulating microbial inheritance has emerged as a key approach in seed microbiome engineering. Researchers divide inheritance into three stages: plant-to-seed, seed dormancy, and seed-to-seedling [2]. Vertical transmission of seed microbiomes is crucial for forming microbial communities and defending against phytopathogens [1].

Recent studies have explored engineering seed microbiomes for targeted functions. Researchers have investigated the potential of seed microbe engineering for producing targeted metabolites or antimicrobial compounds to improve plant biomass and yield under stress conditions [31]. Studies have focused on introducing beneficial bacteria at flowering to engineer the microbiomes of progeny seeds [1].

These innovative approaches to seed microbiome engineering offer promising avenues for enhancing crop resilience, productivity, and sustainability in agriculture. As research in this field continues to advance, we can expect to see more sophisticated and targeted methods for manipulating seed microbiomes to improve crop performance.

5. Seed Biopriming with Beneficial Microorganisms

Seed biopriming with beneficial microorganisms has emerged as a promising approach for enhancing crop performance and sustainability. This technique merges seed priming with beneficial microbes, boosting plant growth, stress tolerance, and agricultural productivity.

Seed biopriming applies beneficial microbes to seeds with controlled hydration, enabling microbial colonization and metabolic activation without triggering germination [32]. Many microorganisms or beneficial microbes have shown effectiveness in seed biopriming (Table 1). For example, *Bacillus*, *Pseudomonas*, and *Azospirillum*, and *Microbacterium* species in Bacteria [32,33]; *Trichoderma* species in Fungi [34]; and *Cyanobacteria*, which is particularly useful for dryland restoration [33].

Table 1. Selected beneficial microbes in seed science.

Beneficial microbes	Beneficial effects	Mode of action	Example strains	References
<i>Trichoderma harzianum</i>	Improves seed germination and enhances plant growth;	Stress Tolerance	<i>T. harzianum</i> strain T22; <i>T. harzianum</i> strain S. INAT	[35–38]
	Provides stress tolerance against biotic and abiotic factors;			
	Alleviates physiological stresses in germinating seeds and seedlings;	Enhancement; Seed Germination and Seedling		
	Promotes root colonization and enhances disease resistance.	Vigor Enhancement		

<i>Bacillus subtilis</i>	Enhances seed germination rates and promotes plant growth;		
	Improves stress tolerance by increasing chlorophyll content and root length under saline conditions;	Seed Germination and Seedling Vigor	<i>B. subtilis</i> HS5B5; <i>B. subtilis</i> ER-08;
	Acts as a biocontrol agent against pathogens;	Enhancement; Disease Suppression;	<i>B. subtilis</i> QM3
	Stimulates plant growth through production of phytohormones and stress-related metabolites.		
			[39–42]
<i>Pseudomonas fluorescens</i>	Enhances seed germination and promotes plant growth by producing phytohormones;	Seed Germination and Seedling Vigor	<i>P. fluorescens</i> SP007S; <i>P. fluorescens</i> F113
	Improves nutrient acquisition;	Enhancement;	
	Suppresses various plant diseases through production of antimicrobial compounds.	Direct Growth Promotion	
			[43–46]
<i>Paenibacillus polymyxa</i>	Promotes plant growth and enhances stress tolerance;		
	Suppresses diseases by producing various antibiotics;	Direct Growth Promotion; Disease Suppression	<i>P. polymyxa</i> HK4; <i>Paenibacillus polymyxa</i> E681
	Improves germination and protects plants against pathogenic fungi, oomycetes, and bacteria.		
			[47–49]
<i>Variovorax</i> sp.		Direct Growth Promotion; Stress Tolerance	<i>Variovorax</i> sp. P1R9;
	Enhances wheat germination under salt stress conditions;	Enhancement; Seed Germination and Seedling Vigor	<i>Variovorax</i> sp. EBFNA2
	Improves biomass;		
	Reduces lipid peroxidation.		
		Enhancement	[50,51]

<i>Azospirillum brasilense</i>	Enhances seed germination;		<i>Azospirillum</i>	
	Increases root length;	Direct Growth	<i>brasilense</i> Ab-	
	Promotes plant growth	Promotion;	V5;	
	through auxin production,	Seed	<i>Azospirillum</i>	
	stimulating root	Germination	<i>brasilense</i> Ab-	
	development and improving	and Seedling	V6;	
	nutrient uptake;	Vigor	<i>Azospirillum</i>	
	Enhances stress tolerance in	Enhancement	<i>brasilense</i>	
	plants.		<i>Sp245</i>	[52,53]

Biopriming significantly improves seed germination rates, uniformity, and seedling vigor [6]. Studies have reported increases in germination potential by up to 60% compared to control groups [54]. Biopriming with PGPB can boost crop yields by 12-20% [6]. This improvement is attributed to enhanced nutrient uptake, hormone production, and overall plant fitness [32,33]. Bioprime seeds show increased resilience to both biotic and abiotic stresses, including improved tolerance to drought, salinity, and heavy metals [33,54] and enhanced resistance against soil-borne pathogens [32,33]. Seed biopriming offers an environmentally friendly alternative to chemical fungicides and fertilizers, aligning with sustainable agriculture goals [6].

Recent research has demonstrated the effectiveness of biopriming across a diverse range of crops, highlighting its potential to enhance agricultural productivity and resilience. In wheat, biopriming has been shown to improve drought tolerance and boost germination potential [54]. Carrot seeds treated with biopriming techniques have exhibited enhanced germination rates and overall plant growth promotion [6]. Furthermore, the benefits of biopriming extend to other major crops such as maize, barley, pea, tomato, and sunflower, where significant improvements have been observed in germination rates, seed viability, and ultimately, crop yield [33]. These findings collectively underscore the versatility and efficacy of biopriming as a promising approach in modern agriculture, offering potential solutions to various challenges faced by farmers across different crop types.

In conclusion, seed biopriming with beneficial microorganisms represents a significant advancement in sustainable agriculture, offering a multifaceted approach to improving crop performance and resilience. As research in this field continues to evolve, it holds exciting potential for addressing global food security challenges while promoting environmentally friendly farming practices.

6. Specific Microbial Strains and Their Effects

Recent studies show specific microbial strains significantly impact seed germination, plant growth, and stress tolerance (Table 1). *T. harzianum* demonstrated the greatest inhibition of seed mycoflora like *Alternaria* sp. and *Fusarium* spp. in cucumber seeds. It significantly improved seed germination (88.75%), shoot length (14.58 cm), root length (13.58 cm), and seedling vigor (2501.31) [35]. *T. viride* and *T. virens* also showed strong inhibition of seed mycoflora and improvements in seed germination and seedling growth, though slightly less than *T. harzianum* [35]. Applying *T. viride* with *Pseudomonas fluorescens* to cabbage seeds improved seedling vigor, root length, biomass, and chlorophyll content [55].

*Bacillus* strains have demonstrated plant growth-promoting effects. *B. subtilis* showed ability to solubilize potassium and phosphorus, and produce indole acetic acid (IAA), contributing to improved seed germination [6]. However, in a comparative study with other bioagents, *B. subtilis* was found to be less effective than *Trichoderma* species in improving cucumber seed germination and seedling vigor [35].

*Pseudomonas fluorescens* effectively controls soft rot caused by *Pectobacterium carotovorum* in kale [56]. When applied in combination with *T. viride* to cabbage seeds, *P. fluorescens* significantly improved seedling quality characteristics and yield-related traits [55].

Mycorrhizal fungi (*Rhizophagus irregularis*) and *Trichoderma* spp. modulate root exudation patterns, releasing metabolites that attract symbiotic microbes while suppressing pathogens. In wheat, *Trichoderma harzianum* increases phenolic compounds and peroxidase activity, reinforcing cell walls against oxidative stress. Similarly, *Azospirillum lipoferum* enhances nitrogen fixation in maize, improving radicle growth and early biomass accumulation [5] (Table 1).

Recent research in FEMS Microbiology Ecology (2024) developed a method for engineering seedling microbiota using synthetic microbial communities (SynComs). This approach successfully modulated seed microbiota composition, with SynComs contributing 80% of the seedling microbiota. Strain abundance on seeds was identified as a key driver of colonization, with *Enterobacteriaceae* and *Erwiniaceae* as strong colonizers and *Bacillaceae* and *Microbacteriaceae* as weak colonizers [9].

These findings demonstrate the diverse and significant effects of specific microbial strains on plant health and productivity. The research emphasizes the potential of these microorganisms in sustainable agriculture practices, particularly in enhancing crop resilience to biotic and abiotic stresses.

## 7. Mechanisms of Action

Recent research has revealed several key mechanisms through which beneficial seed-associated microbes promote plant growth and enhance stress tolerance. These mechanisms can be broadly categorized into direct growth promotion, stress tolerance enhancement, and disease suppression (Figure 1).

Beneficial microbes promote plant growth by producing hormones like IAA, gibberellins, and cytokinins, which enhance root elongation, biomass, and development [5]. Many beneficial bacteria can solubilize essential nutrients like phosphorus, potassium, and zinc, making them more readily available for plant uptake [57]. Certain bacteria, especially rhizobia in legumes, can fix atmospheric nitrogen, reducing the need for synthetic fertilizers [6]. For instance, *Bacillus gaemokensis* upregulates salicylic acid (SA), ethylene (ET), and jasmonic acid (JA) pathways in cucumbers, conferring systemic resistance against bacterial pathogens and insect herbivores [58]. Cyclodipeptides from *Bacillus* strains activate defense-related genes, such as *PR-1* and *PDF1.2*, even before pathogen exposure [58]. In rice, biopriming with *Paenibacillus yonginensis* alters DNA methylation patterns, enhancing drought tolerance through sustained expression of osmoprotectant genes [6,33].

Beneficial microbes boost drought tolerance by releasing osmolytes, improving seed germination and growth under water stress [57]. Microbes like *Trichoderma harzianum* improve the activities of antioxidant enzymes (SOD, APX, CAT, POD) in plants, enhancing their ability to cope with oxidative stress [5]. For example, in sunflower, *Paraburkholderia phytofirmans* upregulates superoxide dismutase (SOD) and catalase (CAT), lowering lipid peroxidation by 45% under 150 mM NaCl stress [23,33]. Some bacteria produce 1-aminocyclopropane-1-carboxylic acid (ACC) deaminase, which helps regulate ethylene levels in plants, improving stress tolerance [57]. Microbial consortia mitigate drought through osmolyte synthesis. *Bacillus subtilis* QST 713 increases proline levels in wheat seedlings, maintaining cell turgor and reducing electrolyte leakage by 30% [6].

Beneficial microbes protect plants by producing antibiotics that suppress pathogens [10,19,59]. Certain microbes activate plant defenses, enhancing resistance to various pathogens [19,20,27,60]. Beneficial microbes can outcompete pathogens for nutrients and space in the rhizosphere [6,7]. Seed treatments with *Bacillus velezensis* CMRP 4490 produce lipopeptides (e.g., surfactin) that disrupt *Fusarium graminearum* hyphae, cutting infection rates by 70% in soybean [5]. *Trichoderma harzianum* induces systemic resistance in tomatoes via  $\beta$ -1,3-glucanase and chitinase expression, reducing *Botrytis cinerea* incidence by 60% [5,23].

Beneficial microbes directly influence seed germination and seedling development. Microbes can increase the activity of germination-related enzymes like  $\alpha$ -amylase and invertase, improving

seed germination and seedling vigor [5,26]. Microbes also help mobilize seed reserves, providing energy for seedling growth [5]. *Spirulina platensis* coatings restrict cadmium translocation in maize, reducing shoot Cd accumulation by 57% through extracellular sequestration and glutathione-S-transferase activation [5]. Similarly, *Enterobacter spp.* in sunflower enhance phytochelatin synthesis, chelating lead and arsenic in root vacuoles [33,61].

These mechanisms demonstrate the multifaceted ways in which beneficial seed-associated microbes contribute to plant health and productivity. The complexity and diversity of these interactions highlight the potential for developing targeted microbial treatments to address specific agricultural challenges, from improving crop yields to enhancing resilience against climate change-induced stresses [12,57].

## 8. Conclusions and Future Directions

Recent advances in beneficial microbes for seed science have revealed promising avenues for enhancing crop performance and sustainability. This review synthesized the latest findings on microbial applications in seed science, emphasizing their role in agricultural productivity, abiotic/biotic stress mitigation, and ecosystem restoration. Several future directions are emerging.

The seed microbiome has been identified as a crucial factor in plant health, development, and resilience to environmental stresses. Research has shown that seed-associated microbes play vital roles in germination, seedling establishment, and long-term plant fitness. Microbial efficacy depends on strain-plant compatibility. For example, *Trichoderma virens* isolates exhibit 40% variability in soybean germination outcomes, necessitating tailored formulations [5]. Advances in nanoencapsulation (e.g., chitosan-coated *Pseudomonas spp.*) prolong viability during storage, maintaining 90% cell viability after 12 months [61]. Research is needed on thermotolerant strains (e.g., *Geobacillus spp.*) for seed treatments in warming climates. Preliminary trials show *Geobacillus stearothermophilus* enhances rice germination at 42°C, though field validation is pending [6].

Studies have demonstrated the vertical transmission of beneficial microbes from parent plants to seeds, suggesting a mechanism for passing advantageous traits to offspring. This finding opens new possibilities for harnessing seed microbiomes to improve crop performance across generations.

Seed biopriming with beneficial microorganisms has shown significant potential in improving germination rates, seedling vigor, and crop productivity. Research indicates that biopriming can increase crop production by ~20% and enhance plant resilience under adverse conditions.

Researchers are exploring innovative approaches to manipulate seed microbiomes for enhanced crop performance. The development of SynComs shows promise for seedling microbiota engineering. A 2024 study in *FEMS Microbiology Ecology* demonstrated that SynCom inoculation on seeds could successfully modulate seedling microbiota composition, with SynComs contributing up to 80% of the seedling microbiota [33]. There is growing interest in understanding and manipulating microbial inheritance and vertical transmission. Researchers are investigating ways to introduce beneficial bacteria at the flowering stage to engineer the microbiomes of progeny seeds [62].

Future research aims to engineer seed microbiomes to produce targeted metabolites or antimicrobials, boosting plant biomass and yield under stress [31]. The application of advanced -omics technologies is providing deeper insights into seed-microbe interactions. The 2024 study published in *Microbiome* used GWAS to identify genetic loci associated with seed endophyte diversity in fonio millet. This approach opens up new possibilities for understanding the genetic basis of plant-microbe interactions in seeds [31].

Research focuses on developing microbial inoculants to boost crop resilience to climate change by improving seed germination and seedling growth under stresses like drought, salinity, and extreme temperatures [61]. The potential of seed-associated microbes in promoting sustainable agriculture and ecosystem restoration is gaining attention. A 2024 review in *Frontiers in Plant Science* highlighted the potential of seed biopriming with beneficial microorganisms to enhance crop resilience and effectiveness under adverse conditions, potentially increasing crop production by 12-20% [62]. Improving the formulation and delivery of beneficial microbes for seed applications is

crucial. Research is needed to develop more effective seed coating technologies that ensure the viability and efficacy of beneficial microbes during storage and after planting [61].

Beneficial microbes revolutionize seed science by bridging agronomic productivity and ecological sustainability. From molecular priming to large-scale ecosystem restoration, microbial technologies offer scalable solutions to global challenges like food security and climate change. Future success hinges on interdisciplinary collaboration—integrating microbiology, genomics, and precision agriculture—to optimize microbial consortia for diverse cropping systems. As seed coatings and biopriming gain traction, they promise to redefine agricultural paradigms, ensuring resilient food systems for generations to come [23,61].

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