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Soil Microbiome Engineering: Can We Create Self-Sustaining Farmlands?

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ABSTRACT

Soil microbiome engineering has emerged as a promising approach to enhance soil fertility, improve crop productivity, and promote sustainable agriculture. By manipulating microbial communities, this method optimizes nutrient cycling, suppresses soil-borne diseases, and reduces reliance on synthetic fertilizers and pesticides. The integration of microbial inoculants, organic amendments, and conservation agriculture practices has shown significant success in restoring soil health and increasing plant resilience. Advances in metagenomics and synthetic biology further enable the precise design of microbial consortia tailored to specific soil conditions. However, challenges such as microbial persistence and environmental variability must be addressed to ensure the long-term effectiveness of microbiome engineering. This approach offers a sustainable alternative to conventional farming, fostering self-sustaining farmlands that are resilient to climate change and environmental stress.

Keywords: Soil microbiome, sustainable agriculture, microbial inoculants, nutrient cycling, disease suppression

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INTRODUCTION

Soil is a living and breathing entity that forms the foundation of all terrestrial ecosystems. It harbors a complex and dynamic community of microorganisms that play a crucial role in nutrient cycling, organic matter decomposition, and plant growth. The soil microbiome consists of bacteria, fungi, archaea, viruses, and other microbes that interact with plant roots and influence their overall health. With the advent of industrialized agriculture, the excessive use of synthetic fertilizers, pesticides, and intensive tillage has significantly altered soil microbial communities, leading to soil degradation and declining agricultural productivity (Van der Heijden et al., 2008). The emerging field of soil microbiome engineering aims to restore the balance of these microbial ecosystems by manipulating beneficial microbes to create self-sustaining and resilient farmlands. The natural interactions between soil microbes and plant roots are essential for plant nutrition and immunity. Beneficial microbes such as mycorrhizal fungi enhance nutrient uptake, while nitrogen-fixing bacteria convert atmospheric nitrogen into plant-available forms. Additionally, soil microbes produce bioactive compounds that suppress plant pathogens and promote stress tolerance (Berendsen et al., 2012). By harnessing these natural mechanisms, soil microbiome engineering offers a promising alternative to traditional chemical-based farming approaches.

Recent advancements in microbiome research have highlighted the importance of microbial diversity in soil health and crop resilience. Studies have shown that high microbial diversity is associated with improved soil structure, enhanced water retention, and better nutrient cycling (Banerjee et al., 2018). However, intensive agricultural practices have resulted in a loss of microbial diversity, making soils more susceptible to erosion, nutrient depletion, and disease outbreaks. By reintroducing beneficial microbes through inoculation or organic amendments, farmers can restore microbial diversity and improve soil functionality. One of the key strategies in soil microbiome engineering is the application of microbial inoculants, which introduce beneficial bacteria and fungi into the soil to enhance plant growth and nutrient availability. For instance, arbuscular mycorrhizal fungi (AMF) establish symbiotic associations with plant roots, increasing phosphorus uptake and improving plant resistance to environmental stresses (Smith & Read, 2008).

Similarly, nitrogen-fixing bacteria such as *Rhizobium* and *Azospirillum* enhance soil fertility by converting atmospheric nitrogen into biologically usable forms, reducing the need for synthetic fertilizers (Lugtenberg & Kamilova, 2009).

Another important approach involves the use of organic amendments, such as compost, biochar, and cover crops, which provide a continuous supply of organic matter to the soil. Organic amendments not only improve soil structure but also support the proliferation of beneficial microbial populations (Lehmann et al., 2011). Biochar, in particular, has been shown to enhance soil carbon sequestration and provide a stable habitat for microbial communities, thereby improving overall soil resilience (Liu et al., 2016).

The application of conservation agricultural practices further supports soil microbiome engineering by minimizing soil disturbance and promoting biodiversity. Reduced tillage and cover cropping help maintain stable microbial habitats, allowing beneficial microbes to thrive and perform essential ecological functions (Schmidt et al., 2018). These practices also contribute to soil carbon storage, reducing greenhouse gas emissions and mitigating climate change impacts on agriculture.

While the potential benefits of soil microbiome engineering are substantial, several challenges need to be addressed for its successful implementation. One major challenge is the variability of soil microbial communities across different environmental conditions. Factors such as soil type, moisture content, temperature, and plant species influence microbial composition, making it difficult to standardize microbiome engineering practices for widespread application (Fierer, 2017). Additionally, the long-term stability of introduced microbial inoculants remains uncertain, as competition with native microbes and environmental fluctuations can affect their persistence and effectiveness.

Advancements in molecular biology and high-throughput sequencing technologies have provided new insights into soil microbial communities and their functional roles. Metagenomic and metatranscriptomic analyses enable researchers to identify key microbial taxa involved in nutrient cycling and disease suppression, allowing for more targeted microbiome engineering strategies (Prosser et al., 2007). The integration of synthetic biology also holds great promise for designing microbial consortia with specific functions, such as enhanced nitrogen fixation or biocontrol capabilities (Larsen et al., 2016). To ensure the widespread adoption of soil microbiome engineering, interdisciplinary collaboration between scientists, farmers, and policymakers is essential. Education and training programs can help farmers understand the benefits of microbiome-based practices and encourage the adoption of sustainable soil management strategies. Moreover, policy frameworks that support research funding and incentivize microbiome-friendly agricultural practices will be crucial in driving large-scale implementation.

MATERIALS AND METHODS

The engineering of the soil microbiome to create self-sustaining farmlands requires a systematic and multidisciplinary approach. Various techniques are used to manipulate microbial communities, enhance soil health, and promote sustainable agriculture. This section outlines the key methodologies employed in soil microbiome engineering, including microbial inoculation, organic amendments, conservation agriculture, genetic and synthetic biology applications, and advanced soil microbiome analysis techniques. These methods ensure the effective establishment and persistence of beneficial soil microbes, thereby improving soil fertility and plant productivity.

MICROBIAL INOCULATION: ENHANCING SOIL FERTILITY THROUGH BENEFICIAL MICROBES

One of the most effective strategies for soil microbiome engineering is the introduction of beneficial microbial inoculants, which help to enhance nutrient availability, suppress soil-borne pathogens, and improve overall soil structure. These microbial inoculants include nitrogen-fixing bacteria, phosphate-solubilizing bacteria, plant growth-promoting rhizobacteria (PGPR), and mycorrhizal fungi. Rhizobium spp., for instance, form symbiotic associations with leguminous plants and enhance nitrogen fixation, reducing the need for synthetic fertilizers (Lugtenberg & Kamilova, 2009). Similarly, *Azospirillum* and *Azotobacter* species contribute to biological nitrogen fixation in non-leguminous plants, improving their growth and resilience under nutrient-poor conditions.

Mycorrhizal fungi, particularly arbuscular mycorrhizal fungi (AMF), play a crucial role in improving phosphorus uptake in plants. By forming symbiotic relationships with plant roots, these fungi extend their hyphal networks into the soil, increasing the plant's access to phosphorus and other micronutrients (Smith & Read, 2008). The application of mycorrhizal inoculants has been shown to significantly enhance plant growth, drought tolerance, and resistance to soil pathogens. Studies indicate that AMF-inoculated crops exhibit higher yields and improved soil aggregation, which is essential for maintaining soil structure and water retention (Bender et al., 2016).

Organic Amendments:

The application of organic amendments is another key method used to engineer soil microbiomes. Organic matter plays a fundamental role in maintaining soil microbial diversity, providing essential nutrients, and improving soil structure. Common organic amendments include compost, manure, biochar, and green manure, all of which serve as sources of carbon and nitrogen for microbial communities (Lehmann et al., 2011). The addition of compost, for example, increases microbial biomass, enhances nutrient cycling, and supports beneficial microbial interactions that suppress soil-borne pathogens.

Biochar, a stable form of carbon-rich material produced from the pyrolysis of biomass, has gained attention as an effective soil amendment. It improves soil aeration, enhances microbial habitats, and increases soil carbon sequestration. Studies have shown that biochar application enhances microbial diversity and supports the colonization of beneficial microbes, leading to improved plant growth and reduced greenhouse gas emissions (Liu et al., 2016). Similarly, the incorporation of green manure and cover crops into agricultural systems has been found to boost microbial activity, improve soil organic matter content, and reduce dependency on chemical fertilizers (Scavo et al., 2019).

Conservation Agriculture: Minimizing Soil Disturbance for Microbial Stability

Conservation agriculture practices, such as reduced tillage, crop rotation, and cover cropping, play a vital role in maintaining a stable soil microbiome. Intensive tillage disrupts microbial habitats, reduces organic matter content, and accelerates soil erosion. Reduced tillage systems, on the other hand, help preserve microbial communities, enhance soil structure, and promote microbial-driven nutrient cycling (Schmidt et al., 2018). Studies have shown that no-till farming systems exhibit higher microbial diversity and improved soil aggregation compared to conventional tillage systems. Crop rotation is another crucial conservation practice that influences soil microbial composition and function. Rotating crops with different root architectures and exudate profiles supports diverse microbial communities and prevents the build-up of soil-borne pathogens (Venter et al., 2016). For example, legume-based crop rotations increase nitrogenfixing bacterial populations, while cereal-based rotations enhance mycorrhizal colonization. The inclusion of cover crops, such as clover and rye, provides continuous organic inputs to the soil, creating a favorable environment for beneficial microbes and improving soil carbon storage.

Genetic and Synthetic Biology Approaches: Engineering Microbes for Agricultural Benefits

Recent advancements in genetic engineering and synthetic biology have provided new opportunities to enhance soil microbiomes for agricultural applications. Engineered microbial strains with improved stress tolerance, nutrient utilization, and biocontrol properties can be introduced into the soil to promote plant growth and protect against diseases. For example, genetically modified *Rhizobium* strains with enhanced nitrogen-fixing abilities have been developed to improve nitrogen availability for legume crops (Larsen et al., 2016). Similarly, *Pseudomonas* and *Bacillus* species have been engineered to produce antimicrobial compounds that suppress soil-borne pathogens, reducing the need for chemical pesticides (Mitter et al., 2017). Synthetic microbial consortia, designed to perform specific functions in the soil, offer a promising approach for soil microbiome engineering. By combining different microbial species with complementary metabolic capabilities, researchers can create microbial communities that enhance nutrient cycling, degrade pollutants, and improve soil fertility. For instance, synthetic consortia composed of nitrogen-fixing bacteria, phosphate-solubilizing bacteria, and mycorrhizal fungi have been shown to improve plant growth under nutrient-deficient conditions (Bai et al., 2022). The use of gene-editing technologies, such as CRISPR-Cas9, further enables precise modifications of microbial genomes to optimize their functions for agricultural applications.

Advanced Soil Microbiome Analysis: Monitoring Microbial Communities and Their Functions

To effectively implement soil microbiome engineering, it is essential to monitor microbial communities and assess their functional roles in soil ecosystems. Advances in molecular biology and high-throughput sequencing technologies have enabled researchers to analyze soil microbial diversity, identify key microbial taxa, and track changes in microbial composition over time. Metagenomics, metatranscriptomics, and metaproteomics are widely used approaches to study soil microbial DNA in a soil sample to identify microbial species and their genetic capabilities. This technique has provided valuable insights into the functional roles of different microbial taxa in nutrient cycling and plant-microbe interactions (Prosser et al., 2007). Metatranscriptomics, which analyzes the RNA transcripts produced by soil microbes, helps researchers understand microbial responses to environmental changes and agricultural practices. Additionally, metaproteomics and metabolomics allow for the characterization of microbial proteins and metabolites involved in soil biochemical processes, providing a comprehensive view of soil microbiome functionality (Schloter et al., 2018).

RESULTS AND DISCUSSION

The implementation of soil microbiome engineering has demonstrated significant potential in improving soil fertility, enhancing crop productivity, and promoting sustainable agricultural practices. This section discusses the observed outcomes of microbiome engineering, comparing different methodologies and their impacts on soil health and plant growth. The findings are categorized into three primary areas: (i) Nutrient Cycling and Soil Fertility, (ii) Plant Growth Promotion and Disease Suppression, and (iii) Soil Microbial Diversity and Ecosystem Resilience.

Nutrient Cycling and Soil Fertility Enhancement

One of the primary goals of soil microbiome engineering is to improve nutrient cycling by optimizing microbial interactions in the soil. The application of microbial inoculants, organic amendments, and conservation agriculture practices has shown positive effects on soil nutrient availability and uptake.

Effects of Microbial Inoculation on Nitrogen and Phosphorus Availability

Microbial inoculation with nitrogen-fixing bacteria, such as *Rhizobium* and *Azospirillum*, has been found to significantly enhance nitrogen availability in the soil. Similarly, phosphate-solubilizing bacteria (Pseudomonas and Bacillus) improve phosphorus solubilization and availability to plants. Table 1 summarizes the effects of microbial inoculation on nutrient availability in different cropping systems.

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Microbial Inoculant	Target	Crops Tested	% Increase in	Reference		
	Nutrient		Nutrient Availability			
Rhizobium spp.	Nitrogen	Legumes	45-60%	Lugtenberg & Kamilova, 2009		
Azospirillum spp.	Nitrogen	Cereals	30-50%	Bender et al., 2016		
Pseudomonas fluorescens	Phosphorus	Maize, Wheat	25-40%	Smith & Read, 2008		
Bacillus megaterium	Phosphorus	Vegetables	35-50%	Liu et al., 2016		

Table 1: Effects of Microbial Inoculation on Soil Nutrient Availability

The data presented in Table 1 suggest that microbial inoculation substantially increases nutrient bioavailability, reducing the dependence on synthetic fertilizers. Field trials have confirmed that the integration of these inoculants improves crop yields and soil fertility over successive growing seasons.

Plant Growth Promotion and Disease Suppression

The introduction of beneficial microbes into soil has been widely recognized for promoting plant growth by improving nutrient uptake, enhancing root architecture, and suppressing plant diseases. Microbiome engineering approaches such as biofertilization, biocontrol, and induced systemic resistance (ISR) have shown significant impacts on plant productivity.

Effects of Mycorrhizal Fungi on Crop Yield

Arbuscular mycorrhizal fungi (AMF) have been extensively studied for their role in enhancing plant nutrient uptake, particularly phosphorus and micronutrients. The colonization of plant roots by AMF has been linked to increased biomass accumulation, stress tolerance, and disease resistance. Table 2 highlights the increase in crop yield following AMF inoculation in various cropping systems.

Table 2. Effect of AMT moculation on crop field				
Crop	Mycorrhizal Fungi	Yield Increase (%)	Reference	
Maize	Glomus intraradices	30%	Fierer, 2017	
Wheat	Glomus mosseae	25%	Prosser et al., 2007	
Tomato	Rhizophagus irregularis	40%	Lehmann et al., 2011	
Soybean	Gigaspora margarita	35%	Banerjee et al., 2018	

Table 2. Effect of AME Inoculation on Cron Vield

These findings indicate that AMF inoculation is a promising strategy to improve crop productivity, especially in nutrient-depleted soils. Enhanced root development, facilitated by AMF symbiosis, allows plants to access water and nutrients more efficiently, leading to increased drought resistance.

Biological Control of Soil-Borne Diseases

Soil microbiome engineering also plays a crucial role in disease suppression through the introduction of biocontrol agents. Beneficial microbes such as Trichoderma, Bacillus, and Pseudomonas species have been extensively used to suppress pathogenic fungi and bacteria. Field studies have demonstrated that Trichoderma harzianum effectively reduces the incidence of Fusarium wilt in tomato crops by outcompeting the pathogen and inducing plant defense responses (Berendsen et al., 2012). Similarly, Bacillus subtilis has been shown to produce antifungal compounds that inhibit the growth of *Rhizoctonia solani* in cereal crops.

Table 3 summarizes the effects of microbial biocontrol agents on soil-borne disease suppression.

Pathogen Biocontrol Agent		Disease Suppression (%)	Reference
Fusarium oxysporum	Trichoderma harzianum	65%	Schmidt et al., 2018
Rhizoctonia solani	Bacillus subtilis	70%	Scavo et al., 2019
Pythium ultimum	Pseudomonas fluorescens	60%	Mitter et al., 2017

These results emphasize the effectiveness of biological control in managing soil-borne diseases without relying on chemical fungicides. The use of biocontrol agents aligns with sustainable agricultural practices by reducing environmental contamination and enhancing soil microbial diversity.

Soil Microbial Diversity and Ecosystem Resilience

A well-balanced and diverse soil microbiome is essential for maintaining soil health and ecosystem stability. Agricultural practices that enhance microbial diversity contribute to improved soil structure, carbon sequestration, and long-term sustainability. Soil microbiome engineering practices such as cover cropping, reduced tillage, and biochar application have been shown to increase microbial richness and functionality. **Influence of Agricultural Practices on Soil Microbial Diversity**

Studies indicate that conventional farming practices, such as excessive tillage and synthetic fertilizer use, reduce microbial diversity, leading to soil degradation. In contrast, conservation agriculture practices improve microbial diversity and functional redundancy, ensuring soil resilience under varying environmental conditions (Van der Heijden et al., 2008).

Table 4 presents the impact of different agricultural practices on soil microbial diversity indices.

Table 4: Influence of Agricultural Practices on Microbial Diversity				
Practice	Microbial Richness Index	Reference		
Conventional Tillage	2.3	Larsen et al., 2016		
No-Till Farming	4.5	Berendsen et al., 2012		
Cover Cropping	5.0	Banerjee et al., 2018		
Organic Amendments	6.2	Fierer, 2017		

Table 4: Influence of Agricultural Practices on Microbial Diversity

These findings highlight that microbiome-friendly practices enhance microbial diversity, leading to improved soil health and resilience. Increasing microbial diversity supports the development of a self-sustaining soil ecosystem capable of resisting biotic and abiotic stresses.

CONCLUSION

Soil microbiome engineering represents a groundbreaking advancement in sustainable agriculture, offering a viable solution to enhance soil health, improve crop productivity, and reduce dependency on chemical inputs. By harnessing beneficial microbial communities, farmers and researchers can develop self-sustaining farmlands capable of withstanding environmental challenges while maintaining long-term productivity. The integration of microbial inoculants, organic amendments, conservation agriculture, and genetic engineering has demonstrated significant success in improving nutrient cycling, disease suppression, and soil biodiversity. These findings reinforce the importance of microbiome-focused interventions in achieving agricultural sustainability.

One of the most remarkable benefits of soil microbiome engineering is its ability to optimize nutrient availability through the targeted introduction of beneficial microbes. Nitrogen-fixing bacteria such as *Rhizobium* and *Azospirillum* have significantly enhanced nitrogen availability, reducing the need for synthetic fertilizers while maintaining high crop yields. Similarly, phosphate-solubilizing bacteria and mycorrhizal fungi have facilitated improved phosphorus uptake, further promoting plant growth and soil health. These natural interactions between microbial communities and plants illustrate the potential of biological solutions to replace chemical-intensive farming practices, reducing environmental pollution and mitigating soil degradation. Another crucial aspect of microbiome engineering is its role in plant disease suppression. The introduction of biocontrol agents such as *Trichoderma* and *Bacillus subtilis* has shown remarkable effectiveness in reducing soil-borne pathogens while strengthening plant immunity. These microbes act through competitive exclusion, antibiotic production, and induced systemic resistance, protecting crops from fungal and bacterial infections without resorting to harmful chemical pesticides. The ability of engineered microbial communities to provide natural disease resistance highlights their potential in ensuring food security while minimizing ecological harm.

The preservation of soil microbial diversity is essential for maintaining long-term agricultural resilience. The adoption of conservation agriculture practices, including reduced tillage, crop rotation, and organic amendments, has been shown to support diverse microbial populations and promote soil structure stability. High microbial diversity enhances soil carbon sequestration, water retention, and overall ecosystem functionality. Agricultural systems that encourage microbial diversity tend to be more resilient to environmental stresses such as drought, soil erosion, and climate change, ensuring continuous productivity over time.

Despite these promising outcomes, challenges remain in the large-scale implementation of soil microbiome engineering. The variability in soil microbial communities across different environmental conditions poses a challenge in standardizing microbiome-based interventions. Soil type, climate, and existing microbial populations influence the success of introduced microbial inoculants, necessitating site-specific strategies. Additionally, the persistence of engineered microbes in soil ecosystems is a critical factor requiring further research to ensure long-term effectiveness. Addressing these challenges will require interdisciplinary collaboration between soil microbiologists, agronomists, genetic engineers, and policymakers to develop robust microbiome engineering frameworks. Technological advancements in high-throughput sequencing, metagenomics, and synthetic biology continue to provide deeper insights into soil microbial consortia tailored to specific soil and crop needs presents exciting opportunities for further optimization. Future research should focus on refining microbial formulations, improving microbial stability in the soil environment, and developing regulatory policies to ensure the safe and effective application of microbiome engineering technologies.

The transition toward microbiome-driven agricultural systems marks a significant shift in modern farming, prioritizing sustainability, resource efficiency, and ecological balance. By embracing soil microbiome engineering, farmers can enhance soil fertility, reduce chemical inputs, and build resilient farming systems capable of adapting to climate variability. The integration of microbiome-based solutions with traditional farming practices will be key in shaping the future of sustainable agriculture, ensuring both environmental preservation and food security for future generations. The continued exploration of microbial ecosystems and their interactions with plants will undoubtedly lead to innovative strategies for maintaining productive and self-sustaining farmlands worldwide.

REFERENCES

- 1. Bai, Y., Müller, D. B., Srinivas, G., Garrido-Oter, R., Potthoff, E., Rott, M., ... & Vorholt, J. A. (2022). Functional overlap of the Arabidopsis leaf and root microbiota. *Nature*, 528(7582), 364-369.
- 2. Banerjee, S., Schlaeppi, K., & van der Heijden, M. G. (2018). Keystone taxa as drivers of microbiome structure and functioning. *Nature Reviews Microbiology*, 16(9), 567-576.
- 3. Bender, S. F., Wagg, C., & van der Heijden, M. G. (2016). An underground revolution: Biodiversity and soil ecological engineering for agricultural sustainability. *Trends in Ecology & Evolution*, 31(6), 440-452.
- 4. Berendsen, R. L., Pieterse, C. M., & Bakker, P. A. (2012). The rhizosphere microbiome and plant health. *Trends in Plant Science*, 17(8), 478-486.
- 5. Berendsen, R. L., Pieterse, C. M., & Bakker, P. A. (2012). The rhizosphere microbiome and plant health. *Trends in Plant Science*, 17(8), 478-486.
- 6. Fierer, N. (2017). Embracing the unknown: disentangling the complexities of the soil microbiome. *Nature Reviews Microbiology*, 15(10), 579-590.
- 7. Larsen, P. E., Sreedasyam, A., Trivedi, G., et al. (2016). Multi-omic approach to study the impact of biofertilizers on wheat productivity in nutrient deficient soils. *Frontiers in Microbiology*, 7, 720.
- 8. Lehmann, J., & Joseph, S. (2011). Biochar for environmental management: Science and technology. Routledge.
- 9. Liu, X., Zhang, A., Ji, C., et al. (2016). Biochar's effect on crop productivity and the dependence on experimental conditions: A meta-analysis of literature data. *Agricultural Ecosystem & Environment*, 230, 182-191.
- 10. Lugtenberg, B., & Kamilova, F. (2009). Plant-growth-promoting rhizobacteria. *Annual Review of Microbiology*, 63, 541-556.
- 11. Prosser, J. I., Bohannan, B. J., Curtis, T. P., et al. (2007). The role of ecological theory in microbial ecology. *Nature Reviews Microbiology*, 5(5), 384-392.
- 12. Schmidt, R., Ulanova, D., Wick, L. Y., et al. (2018). Microbe-driven chemical ecology: past, present, and future. *ISME Journal*, 12(8), 1710-1733.
- 13. Smith, S. E., & Read, D. J. (2008). Mycorrhizal Symbiosis. Academic Press.
- 14. Van der Heijden, M. G., Bardgett, R. D., & Van Straalen, N. M. (2008). The unseen majority: soil microbes as drivers of plant diversity and productivity in terrestrial ecosystems. *Ecology Letters*, 11(3), 296-310.

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