

## Soil Microbes-Plants-Insect Herbivores: A Tritrophic Interaction

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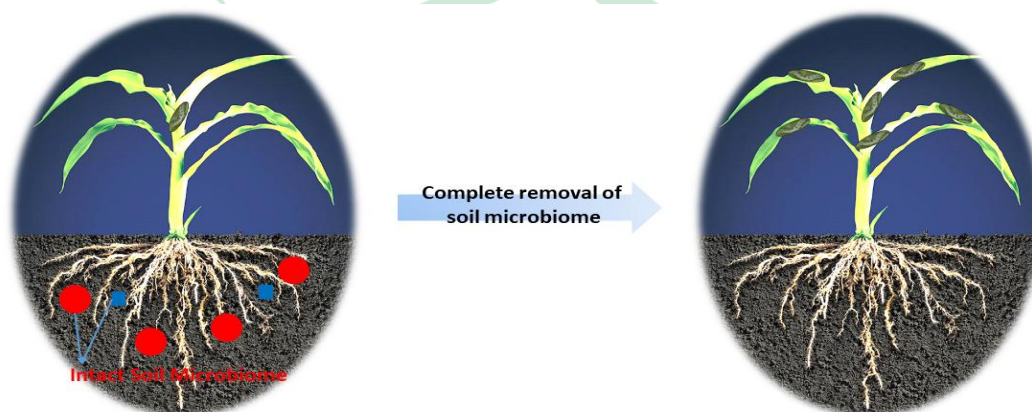
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### Introduction

Plants are acting as a link between the numerous kinds of herbivores and non-pathogenic root-associated microbes which are providing benefits to the plant against insect herbivores (Aljory et al., 2018). We can also call them mediators of multitrophic interactions between microbes and insects. Moreover, plant vegetation also interlinks below ground and above ground community members, such as soil-borne microbes and above-ground insects (Pena et al., 2016). Soil properties and soil history also have a major impact on insect herbivores (Howard et al., 2018). Sometimes a normal soil amendment can have a major detrimental effect on insect herbivores and sometimes many seasonal soil amendments are required for better control on the negative effect of insects on plants (Thomson et al., 2010). Soil properties affect the soil microbiome and this can have direct and indirect effects on the insect herbivores either aboveground or belowground (Pineda et al., 2017).



**Figure 1:** When the soil microbiome is removed completely from the soil the infestation of insect herbivores increases many folds as studied by Hubbard et al., 2019.

Genetic and ecological modifications are being used together to achieve higher crop yields whilst minimizing negative impacts on the ecosystem. Keeping these things in mind, scientists are following various strategies like improving the plant resistance against herbivores and other kinds of abiotic stresses, conservational biological management, and improving the plant growth. Here the importance of such kind of microbes comes in front of the researchers as they are naturally available and don't have any negative impacts on the agro-ecosystem. In this article, we have covered the interaction between different microbes with plants, herbivores, and the tritrophic interaction between plants, microbes, and herbivores.

### **1. Tritrophic interaction between microbes-plants and herbivores**

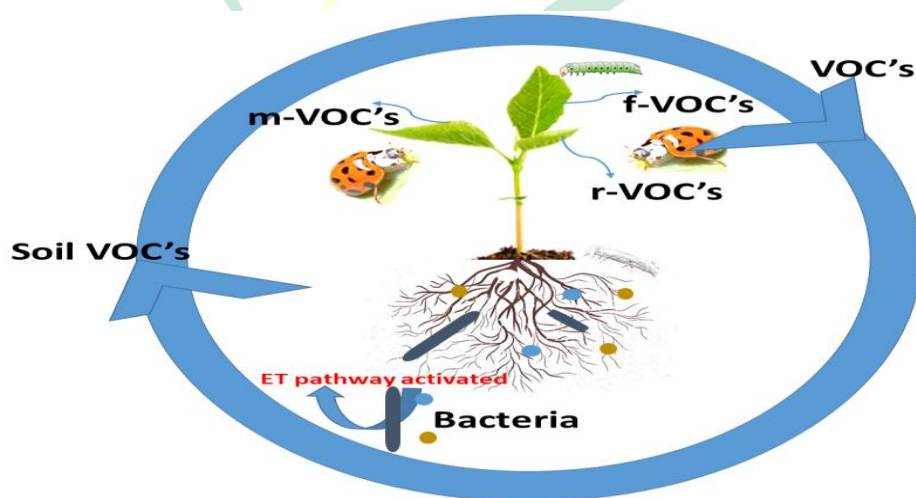
Microbes, plants, and insects are part of the same ecosystem, and in an ecosystem, these three elements can have either direct interaction or indirect interaction. Few interactions are mediated by the plants, and changes are made by insects in the plants, which affect the microbes in the soil. When a plant gets attacked by an insect, it can respond in many different ways. For instance, plants can change the quantity and quality of root exudates, and plants can also modify the soil properties, and this can lead to changes in the plant-microbe interaction. When some changes occur in the distant organs and tissues of the plant due to the attack of insects on plant tissues, it can have an effect on soil microbes and the organisms below ground (Kong et al., 2019), and these changes might not affect the soil microbes in equal proportion. This has been shown by one of the studies done by Landgraf et al. (2012) on *Medicago truncatula*. In *Medicago truncatula*, when the leaf was mechanically wounded many times more than at local level, Jasmonic acid (JA) was increased, but there was a difference in the colonization of mycorrhiza and rhizobacteria (Rivero et al., 2019). It was observed that the colony of mycorrhiza was showing an increasing trend while rhizobacteria remained unaffected.

In addition, acetoin and 2, 3-butanediol are the two volatile organic compounds (VOCs) that are released by bacillus species and were the first to be known for the initiation of induced systemic resistance (ISR), and this initiation is dependent on the ethylene (ET) pathway rather than the salicylic acid (SA) and JA pathways in Arabidopsis plant (Kong et al., 2018, Li et al., 2019). When some insects infest the plant, some of the VOCs are also produced by the plant, and these are termed "herbivore-induced plant volatiles" (HIPVs). The

schematic illustration is shown in figure 3. These volatiles can affect the plant directly or indirectly. Directly, they can either poison the herbivore or repel it. Indirectly, they can help attract the natural enemies of the herbivores (Walters et al., 2017, Lackner et al., 2019).

After the initial study by Ryu et al. (2004), many more studies related to these chemicals have been done. In one of the studies, the expression of terpene genes was initiated by the leaf-chewing herbivores via the JA signalling pathway. This can have a role in insect-plant interaction (Erb et al., 2019). In one of the similar kinds of studies, scientists have shown that when *P. fluorescens* WCS417r (currently called *Pseudomonas simiae* WCS417r) colonizes the roots of Arabidopsis, it leads to the increment of plant volatiles which are dependent on JA signalling pathway genes when the caterpillar feeds on the plant (Disi et al., 2019).

In one of the other studies, when rhizobacterium was added, the expression of some of the genes like TPS03 and TPS04 was decreased due to the suppression of the aromatic compounds like terpene (*E*)- $\alpha$ -bergamotene when a caterpillar attacked the plant. Due to this suppression, an increasing number of parasitoids are attracted towards the caterpillar and it leads to an indirect defence against insect herbivores (Block et al., 2019). These kinds of studies provide evidence that these chemicals can be of great use in controlling insect herbivores in an eco-friendly and better way. We can apply much more innovative ideas and research in this field.



**Figure 2:** Soil microbes like *Bacillus* species initiate the ISR with the help of volatile organic compounds (VOC's) like acetoin and 2,3-butanediol, and this is dependent on the initiation of

the ET pathway. There are VOCs which are induced by the herbivore infestation and are termed "herbivore induced plant volatiles" (HIPVs). The emission of these VOC's attracts the natural enemies.

## 2. Host immunity modulations through soil microbes

Every microorganism has two ways to save itself from the immune system of the host: either to hide or to modulate the immune system of the host through some chemicals or some molecules. Every microorganism in the body of the host is recognized as an alien by the immune system of the host, whether the microorganism is beneficial or detrimental to the host body (Thakur et al., 2019). For the establishment of a relationship between the host and the microbe, the modulation of the immune system is a necessary step. Compounds like flagellin and chitin are normally identified by pattern recognition receptors (PRRs) in the host immune system, and these patterns are normally known as microbe associated molecular patterns (MAMPs) and pathogen associated molecular patterns (PAMPs). Basically, identification of these patterns is the initiating step in plant defense (Shehzadi et al., 2019). For the initiation of ISR, beneficial soil microbes must have a colony near the root system of the plant. MAMPs and PAMPs together are known as MAMP-triggered immunity (MTI) (Van der et al., 2019). And avoidance of MTI can lead the microbes to successful and effective colonization of the host's root (Bulgari et al., 2019).

Soil microbes which are beneficial to the plants have co-evolved with the plant and have developed strategies which can overcome the host immune system, especially the MTI for the colony formation in the roots and the mutual benefits of plants and microbes both. When soil microbes play the role of mediator in ISR, it mostly depends on the priming. Priming is a good way to suppress the immune system in the roots because it saves a lot of energy (Ling et al., 2019). When ISR-insensitive insects attack the leaves of the plants, then priming becomes irrelevant because priming takes place only when the insect initiates the ISR. In one of the studies done by Song et al., when mycorrhizacolonises the roots of tomato plants, it leads to the over expression of genes like protease inhibitor (PI-1) and allene oxide cyclase (AOC), which are involved in the defence of the plant. This is called priming based defence in tomato plants (Du et al., 2019). A similar kind of study was done by Ahn et al. in arabidopsis priming and its effects were shown by the *Pseudomonas putida* LSW17S gene, which was based on NPR1, ET, and the JA pathway (Jain et al., 2019).

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