**Functional Role of Rhizospheric Endophytes as Plant Growth Promoters**

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**Abstract**

An eco-friendly organic agricultural system is a significant prospect for sustainable agriculture. To overcome the drawbacks of the existing conventional methods using agrochemicals and to increase crop productivity, the application of plant-associated microbes is gaining impetus globally. The plant rhizosphere with its incredibly rich microbial diversity exhibits a harmonious plant-microbe interaction. This plant-microbe interaction can facilitate microbial colonization into the root tissues. These root-colonizing active microbes, categorized as endophytes, are beneficial showing positive relationships with plants. They directly or indirectly benefit plants by promoting growth through the various phytohormones, secondary compounds or signal molecules secreted or modulated by them. The symbiotic interaction between the root-colonizing endophytic microbes enables them to establish an association with the crop plants to overcome nutrient deficiencies and environmental stresses. Integration of beneficial endophytes in agricultural practices as biofertilizers could efficiently neutralize the effect of agrochemicals, thereby reducing their harmful environmental impact. Endophytic microbial biofertilizers could also become a cost-effective and eco-friendly approach to acquire higher crop yields. Knowledge regarding the importance of endophytes in agriculture ensures various possibilities for formulating novel endophytic bioproducts such as potent bioinoculants for their prospective application in larger agricultural fields. This comprehensive review is a compilation of research on the functional role of root-colonizing endophytic microbes and highlights the mechanism of colonization of the endophytic rhizomicrobiome. The review also discusses the functional details of endophytes involved in plant growth promotion and their significant use in agrobiotechnology for crop improvement programs to ensure future food security.

**Keywords**: Endophytes, plant growth promoter, biofertilizer, rhizosphere, sustainable agriculture.

**1.**   **Introduction**

The rhizosphere- the most versatile and highly intricate entity of the soil surrounding the root zone (Manoharan *et al*., 2017; Gouda *et al*., 2018) is unique with its rich rhizodeposition that attracts a distinctive microbial community. These root exudates are sources of microbial energy capable of initiating a strong plant root-microbial interaction (Philippot *et al*., 2013; Bai *et al*., 2015; Cole *et al*., 2017; Bhupendra*et al*., 2019). These interactions often promote a range of beneficial associations between the plant and the microbe thereby effectively encouraging the colonization of the microbes inside the roots. Such root-colonizing active microbes are categorized as rhizospheric endophytes (Lacava *et al*., 2022). The strong root tissue-endophyte interactions equip the rhizospheric endophytes to establish a steady association with the host plant than the free-living rhizo-microbes. These mutualistic or trophobiotic associations with the host plants confer rhizospheric endophytes with competence to subsequently enhance plant growth and health (Ryan *et al*., 2008; Santoyo *et al*., 2016; Afzal *et al*., 2019).

Rhizospheric endophytes are considered a distinct group of soil microorganisms involved in plant growth promotion. These plant growth-promoting endophytes (PGPE) are associated with the internal root tissues of every plant species and exhibit an alternate diphasic life cycle between root tissues and soil (Ryan *et al*., 2008; Ma *et al*., 2011; Sessitsch *et al*., 2012). With easier access to nutrients, PGPE proliferates in the root apoplast region with less competition and stays protected from environmental stress (Senthilkumar *et al*., 2011). The diverse group of rhizospheric PGPE are highly beneficial to plants and provide numerous agriculturally promising traits. They are gaining attention for their environmentally benign approach to sustainable agriculture.

The root colonizing PGPE modulates plant growth via direct mechanisms and accelerates plant nutrient absorption and distribution, secretion of phytohormones and plant growth-promoting enzymes. Indirectly they facilitate plant disease resistance, enable allelopathic responses, and mediate host stress tolerance thereby maintaining overall plant health (Fescue, 1990; Mei and Flinn, 2010; Cipollini *et al*., 2012; Chukwuneme *et al*., 2020; Adeleke and Babalola*,*2020). These qualities of the PGPE could be utilized to formulate potential bio-inoculants to efficiently substitute synthetic plant growth regulators and agrochemicals for enhancing the productivity of crops and practising safe agriculture.

Endophytes associated with crop roots are being taken into consideration worldwide in view of their active role in bio-fertilization and bio-controlling of various plant diseases. Developing the rhizosphere endophytic community as a potential tool for sustainable agriculture, and improved crop management aptitude for the future, depends on comprehensive scientific research on PGPE-mediated plant growth stimulation. Such in-depth research would provide a deep insight into the mechanism of colonization and microbe-plant root interactions leading to plant growth promotion. This would expedite the formulation of appropriate isolation methods for developing potent microbial strains that can endure varied agronomic environments.

**2.**   **Colonization and interaction of plant-associated rhizospheric endophytes**

Endophytes are ranked as a subcategory of the rhizospheric microbial system, that invades and inhabits the internal tissues of the host plant (Frommel *et al*., 1993; McInroy and Kloepper 1995; Hallmann and Berg, 2007; Marquez-Santacruz *et al*., 2010). The colonization of endophytes varies according to the environmental condition, plant variety and its developmental phase, microbial species, and the abundance of the microbial population (Tan *et al*., 2003). The physical, chemical, and biological environment of the rhizosphere also attributes to endophytic colonization. Endo-rhizosphere colonization is highly affected by physical factors like temperature, water availability (including precipitation) and soil (type, texture, pH, and topography). Chemical characteristics like the presence of mineral elements, inorganic and organic substances, exudates, and other metabolites also significantly contribute towards endophytic colonization. The colonization, abundance, and diversity of root colonizing endophytes are determined by biological factors. The distribution of other competitive and inhibitory microorganisms like bacteria, fungi, viruses, protozoans, nematodes, and other plant-beneficial growth promoters (Ojuederie *et al*., 2019) can positively or negatively regulate the survival and colonization of microbes in the rhizosphere.

Endophytic microbes adopt various survival strategies for colonizing their host based on which they are catogorized into three groups- obligate, facultative, and passive endophytes (Hardoim *et al*., 2008). Obligate endophytes can thrive only within the plant tissues and get disseminated through seeds, while facultative endophytes are free-living soil microbes whose entry into the host tissues is mediated by signalling molecules. Both of these categories of endophytes play a prominent role in plant growth regulation. The third group of endophytes enter the host tissues via minute wounds on the root hairs without any signalling cues and have little effect on plant growth modulation (Verma *et al*., 2004; Swarnalakshmi *et al*., 2019).

Each plant microbiome is a hub of diverse microbial communities. Endophytic microbial establishment within the plants follows a series of events akin to rhizomicrobial colonization. This is influenced by an array of microbial traits and intricate plant-microbial communication (Afzal *et al*., 2019). The endophytic microorganisms commonly establish their entry through the root zone and immediately after the penetration, the microbes colonize the adjacent root tissues (Zinniel *et al*., 2002; Afzal *et al*., 2019). The colonization process is initiated by the plant root by secreting endophyte-specific root exudates that signal the root colonizing endophytes (de Weert *et al*., 2002; Rosenblueth and Martinez-Romero, 2006). These signalling molecules accelerate the establishment of a strong association between the endophytic microbes and the plants. This lays the foundation for the beneficial plant-endophyte interaction which is believed to promote plant growth and plant disease management (Compant *et al*., 2005) as well as various other functionalities.

**2.1 Rhizosphere colonization of endophytic microbes**

The rhizo-microbial diversity is distinct from the ordinary soil microbial community.  Plant root metabolic exudates like amino acids, sugars, alkaloids, and phenolic compounds attract microorganisms including endophytes towards the rhizosphere making it the most viable region surrounding the plant root (Vandana *et al*., 2021). These organic nutrients and plant transudes act as signals that drive microbial growth and development, where they support 10 to 50 times more bacteria and 5 to 10 times more fungi than normal soil (Ortiz-Castro *et al*., 2009). These root exudates enrich the rhizome-microbiome by serving as a rich carbon source within the rhizosphere thereby sustaining a wide range of the root-associated microbial population.

Microbial communities of the root microbiome interact and compete for their nutrients (Pervaiza, 2020). The competitive rhizospheric microbe interactions are crucial for efficacious root colonization. Endophytic microbes strive fiercely to occupy the rhizosphere for procuring nutrients (Azevedo, 1998, Raaijmakers *et al*., 2002, Oliveira *et al*., 2003). Only the endophytes that successfully colonize rhizospheres will flourish in this space and contribute towards plant growth and development. Such endophytes may subsequently colonize the entire rhizoplane and certain extracellular cells, resulting in the development of rhizodermal microbial colonies (Benizri *et al*., 2001).

Successful colonization of rhizospheric endophytes is dependent on microbial traits like signal recognition, attachment, motility, nutrient utilization, growth rate, antagonistic substance and enzyme secretion, besides polysaccharide synthesis (Compant *et al*., 2010; Santoyo *et al*., 2016; Santos *et al*., 2018). At a later stage, the rhizospheric colonizers secrete metabolites that alter the root cell wall, conceivably enabling the initiation of endophytic growth within the root cortex. These structural alterations in root cell walls are induced by the rhizospheric microbe-mediated change in plant gene expression. The metabolic regulations also associated with altered plant gene expression following rhizosphere colonization by endophytes pave the way to their entry into the root tissues.

**2.2 Root colonization of the endophytic microbes**

Rhizospheric endophytes enter the internal root tissues after colonizing the rhizoplane and establish within the intracellular region as an endophytic sub-population (Hallmann, 2001). Chemo-attraction mediated by the root exudates often stimulates the colonization by plant-specific endophytic species (Sessitsch *et al*., 2002; Krechel *et al*., 2004; Berg *et al*., 2005; Hardoim *et al*., 2008). The rhizosphere specificity of the microbes is determined by the extent of root-microbial interaction. For example, rhizobial colonization in Fabaceae is initiated via root tips leading to root nodule formation (Desbrosses and Stougaard, 2011). Furthermore, rhizospheric endophytes initially infect the host plant by colonizing the breaks at places from where lateral roots emerge. Later they invade the cortical cells of the roots, where they exhibit plant-beneficial properties (Chi *et al*., 2005; Pothier *et al*., 2007; Combes-Meynet *et al*., 2011).

Plant-microbial interaction is initiated through the adhesion of the microbes with the extracellular region of the root tissues. Specific receptors and signalling molecules modulate this, thereby providing a tissue-specific colonization strategy which is unique for each endophyte (Zachow *et al*., 2010). Following their establishment in the extracellular region, rhizospheric endophytes use specialized approaches for invading the internal root tissues.

Rhizospheric endophyte colonization within the root tissues would be either active or passive. Active colonization gets modulated via systematic events like signal recognition, attachment, and colonization while, passive colonization of rhizospheric endophytes happens along the root tips or the wounds formed at the root arising region, or through the injuries produced by the pathogenic microbes (Reinhold-Hurek and Hurek, 1998; Hardoim *et al*., 2008).

Bacterial colonization and movement within the internal root tissues are mediated through receptor-ligand signalling, chemotaxis, quorum sensing, twitching motility, lipopolysaccharides, flagella, and pili (Duijff *et al*., 1997; Dörr *et al*., 1998; Böhm *et al*., 2007, Suárez-Moreno*et al*., 2010). Bioactive secondary metabolites like alkaloids, phenolics, and enzymes help the endophytic fungal strains to recognize the host tissue. Fungi secrete specific enzymes like cellulases and pectinases to degrade the root cell wall for colonization. Similarly, cell-divider debasing compounds (CWDEs) secretion is associated with microbial colonization within the cortical cells (Lodewyckx *et al*., 2002).

**3.**   **Mechanism of plant growth promotion by PGPE**

Rhizospheric endophytes are a distinct subpopulation of rhizospheric microorganisms that colonize the internal root tissues and adopt plant growth promotion strategies. In nature, the interaction between the endosphere colonizing microbes and their host plant is established by the colonization potential inherited within the plant and the microbe. However, microbial growth may be hampered by variations in soil conditions like pH, availability of water, and temperature. Additionally, rhizospheric bacteria also compete with the endophytes and strive to colonize the plant roots (Santoyo *et al*., 2016). Once established, the PGPEs coexist within the plant tissues, supporting the overall physiological and biochemical functioning of their host plant.

Plant-beneficial rhizospheric endophytes could be bacteria or fungi seen associated with a wide range of crop plants (Table 1, Table 2). Bacterial genera like *Arthrobacter*, *Actinobacteria*,*Azospirillum,* *Azotobacter, Escherichia coli, Bacillus, Enterobacter,  Pantoea, Paenibacillus, Pseudomonas, Rhizobium, Serratia*are the common endorizhal endophytes (Hungria *et al*., 2013, Patel and Archana, 2017; Issa *et al*., 2018; Ali *et al*., 2019; Dubey *et al.,* 2021; Cun *et al.,* 2022). Fungal endophytes, which include *Aspergillus, Paecilomyces, Penicillium, Trichoderma,*mycorrhizal fungi, and the non-pathogenic strains of *Alternaria, Fusarium, Verticillium*actively colonize the root tissue andexpedite plant growth promotion and trigger the host plant defense system (Doni *et al*., 2014, Lone *et al*., 2015; Ikram *et al*., 2018; Ali *et al*., 2019).

These root-colonizing PGPEs exist in every agronomic ecosystem. They regulate both root-endophyte association and microbial interactions between the rhizomicrobes. They hold metabolically active traits that promote plant growth and development either directly or indirectly (Fig.1). The rhizospheric PGPEs directly facilitate nutrient acquisition, phytohormone production, and enzyme synthesis which are essential in plant growth under both normal and stressed environments (Wakelin *et al*., 2004; Bhardwaj *et al*., 2014). PGPEs mediate nutrient acquisition through biological nitrogen fixation and solubilization of minerals like phosphorus, potassium, iron, zinc, and other micronutrients which are vital for plant growth promotion.

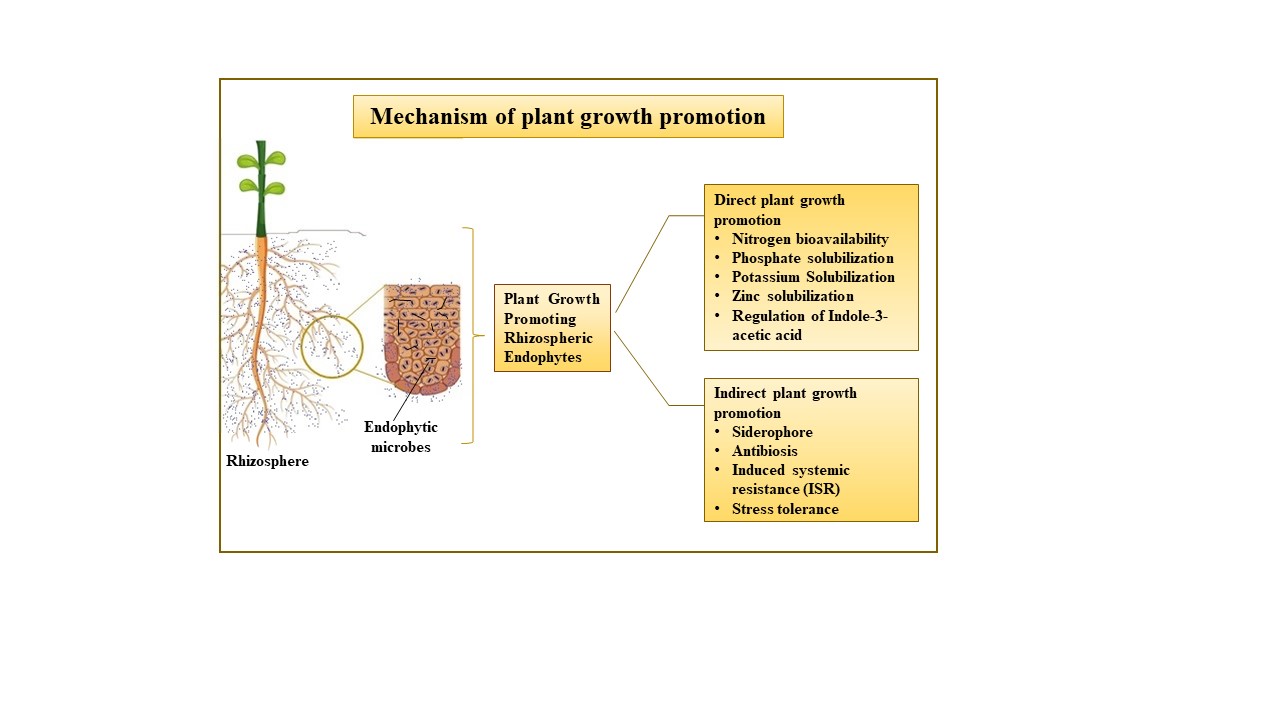
Indirect enhancement of plant growth by PGPEs is affected chiefly by suppressing the growth of phytopathogens. This is achieved through the synthesis of lytic enzymes, and antimicrobial compounds which improve plant defense mechanisms to protect the host from further pathogen attack (Miliute *et al*., 2015). The strategies exhibited by the rhizospheric endophytes coincide, depending on their nature and provenance. All these features act concurrently for plant growth promotion and disease resistance, thus ensuring higher crop yield for achieving agricultural sustainability (Omomowo and Babalola, 2019).

**Table 1 Effective plant growth-promoting strains of rhizospheric bacterial endophytes and their host crops**

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| **Species** | **Crops** | **References** |
| *Acetobacter diazotrophicus* | Sugarcane  (*Saccharum officinarum*) | Dong *et al*., 1994 |
| *Achromobacter sp.* | Wheat (*Triticum aestivum*) | Patel and Archana, 2017 |
| *Acinetobacter sp.* | Wheat (*Triticum aestivum*) | Patel and Archana, 2017 |
| Maize (*Zea mays*) | Cun *et al.,* 2022 |
| *Arthrobacter sp.* | Maize (*Zea mays*) | Cun *et al.,* 2022 |
| *Azoarcus sp.* | Rice (*Oryza sativa*) | Hurek *et al*., 1994; Reinhold-Hurek *et al*., 2006 |
| *Azospirillum sp.* | Sorghum (*Sorghum bicolor*) | Pereira *et al*., 1988; Dobbelaere *et al*., 2001; Molla *et al*., 2001 |
| *Azospirillum sp.* | Wheat (*Triticum aestivum*) | Dobbelaere *et al*., 2001 |
| Banana (*Musa acuminata*) | Mia *et al*., 2007 |
| Rice (*Oryza sativa*) | Ladha *et al*., 1982 |
| Soybean (*Glycine max*) | Hungria *et al*., 2013 |
| Maize (*Zea mays*) | Riggs *et al*., 2001 |
| *Bacillus pumilus* | Rice (*Oryza sativa*) | Bacilio-Jimenez *et al*., 2001 |
| *Bacillus subtilis* | Mulberry (*Morus alba*) | Ji *et al*., 2008 |
| Wheat (*Triticum aestivum*) | Upadhyay *et al*., 2012 |
| Onion (*Allium cepa*) | Weilharter *et al*., 2011; Pan *et al*., 2015; |
| *Bacillus japonicum* | Soybean (*Glycine max*) | Hungria *et al*., 2013 |
| *Bacillus sphaericus* | Banana (*Musa acuminata*) | Mia *et al*., 2007 |
| *Bradyrhizobium elkanii* | Soybean (*Glycine max*) | Hungria *et al*., 2013 |
| *Burkholderia phytofirmans* | Grapevine (*Vitis vinifera*) | Compant *et al*., 2008 |
| Maize (*Zea mays*) | Kost *et al*., 2014; Naveed *et al*., 2014 |
| *Burkholderia vietnamiensis* | Rice (*Oryza sativa*) | Govindarajan *et al*., 2008 |
| Sugarcane  (*Saccharum officinarum*) | Govindarajan *et al*., 2006 |
| *Corynebacterium flavescens* | Rice (*Oryza sativa*) | Bacilio-Jimenez *et al*., 2001 |
| *Enterobacter sp.* | Maize (*Zea mays*) | Riggs *et al*., 2001; Naveed *et al*., 2014; Cun *et al.,* 2022 |
| Wheat (*Triticum aestivum*) | Tian *et al*., 2017 |
| *Gluconacetobacter diazotrophicus* | Sugarcane  (*Saccharum officinarum*) | Suman *et al*., 2005; Luna *et al*., 2010 |
| Sorghum (Sorghum bicolor) | Luna *et al*., 2010 |
| Rice (*Oryza sativa*) | Rouws *et al*., 2010; Meneses *et al*., 2017 |
| *Herbaspirillum sp.* | Sugarcane  (*Saccharum officinarum*) | Weber *et al*., 1999 |
| *Herbaspirillum seropedicae* | Maize (*Zea mays*) | Roncato-Maccari *et al*., 2003; Amaral *et al*., 2014; Balsanelli *et al*., 2014; |
| Rice (*Oryza sativa*) | Elbeltagy *et al*., 2001; James *et al*., 2002 |
| Wheat (*Triticum aestivum*) | Pankievicz *et al*., 2016 |
| *Klebsiella pneumoniae* | Maize (*Zea mays*) | Riggs *et al*., 2001; Dong *et al*., 2003; Weilharter *et al*., 2011 |
| Wheat (*Triticum aestivum*) | Iniguez *et al*., 2004 |
| *Mesorhizobium opportunistum* | Chickpea (*Cicer arietinum*) | Nandasena *et al*., 2009 |
| *Nitrospirillum amazonense* | Sugarcane  (*Saccharum officinarum*) | Schwab *et al*., 2018 |
| *Pantoea agglomerans* | Rice (*Oryza sativa*) | Verma *et al*., 2004 |
| Maize (*Zea mays*) | Rodrigues and Forzani, 2016 |
| Pistachio trees  (*Pistacia vera*) | Etminani and Harighi, 2018 |
| *Pseudomonas fluorescens* | Rape (*Brassica napus*) | Sheng *et al*., 2008b |
| Tomato  (*Solanum lycopersicum*) | Duijff *et al*., 1997; Issa *et al*., 2018 |
| *Pseudomonas putida* | Potato (*Solanum tuberosum*) | Andreote *et al*., 2009 |
| Pea (*Pisum sativum*) | Germaine *et al*., 2006 |
| *Mung bean (Vigna radiata)* | Orozco-Mosqueda *et al.,* 2020 |
| Ginger (*Zingiber officinale*) | Jasim *et al.,* 2014 |
| *Ralstonia sp.* | Wheat (*Triticum aestivum*) | Patel and Archana, 2017 |
| *Rhizobium sp.* | Wheat (*Triticum aestivum*) | Tian *et al*., 2017; Saghafi *et al*., 2018 |
| Faba bean (*Vicia faba*) | Tian *et al*., 2008 |
| Common bean (*Phaseolus vulgaris)* | Lopez-Lopez *et al*., 2010 |
| *Rhizobium pusense* | *Mung bean (Vigna radiata)* | Chaudhary *et al.,* 2021 |
| *Serratia marcescens* | Rice (*Oryza sativa*) | Gyaneshwar *et al*., 2001 |
| *Serratia proteamaculans* | Soybean (*Glycine max*) | Taghavi *et al*., 2009 |
| *Staphylococcus sp.* | Wheat (*Triticum aestivum*) | Tian *et al*., 2017 |

**Table 2 Some rhizospheric fungal endophytes showing effiecient plant growth promotion in crop plants**

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| **Species** | **Crops** | **References** |
| *Aspergillus nidulans* | Tomato  (*Solanum lycopersicum*) | Xia *et al.,* 2019 |
| *Cladosporium sphaerospermum* | Soybean (*Glycine max*) | Hamayun *et al*., 2009 |
| *Coniothyrium aleuritis* | Tomato  (*Solanum lycopersicum*) | Xia *et al.*, 2019 |
| *Fusarium culmorum* | Wheat (*Triticum aestivum*) | Lr, 2018 |
| *Fusarium oxysporum* | Tomato  (*Solanum lycopersicum*) | Xia *et al*., 2019 |
| *Glomus versiforme* | Chickpea (*Cicer arietinum*) | Alloush *et al*., 2000 |
| *Glomus intraradices* | Pepper (*Capsicum annuum*) | Martin and Stutz, 2004; Beltrano *et al*., 2013 |
| *Glomus caledonium* | Cucumber (*Cucumis sativus*) | Ortas, 2010 |
| *Glomus mosseae* | Garlic (*Allium sativum*) | Sari *et al*., 2002 |
| Maize (*Zea mays*) | Lone *et al*., 2015 |
| Potato (*Solanum tuberosum*) | Lone *et al*., 2015 |
| Peach (*Prunus persica*) | Wu *et al*., 2010 |
| *Metarhizium robertsii* | Maize (*Zea mays*) | Ahmad *et al*., 2020 |
| *Penicillium roqueforti* | Wheat (*Triticum aestivum*) | Ikram *et al*., 2018 |
| *Rhizophagus irregularis* | Wheat (*Triticum aestivum*) | Perez-de-Luque *et al*., 2017 |
| Tomato  (*Solanum lycopersicum*) | Khalloufi *et al*., 2017 |
| *Trichoderma sp.* | Rice (*Oryza sativa*) | Doni *et al*., 2014 |



**Figure 1. Mechanism of plant growth promotion by PGPE**

**3.1 Direct mechanism of plant growth promotion by PGPE**

PGPE promotes plant growth directly through a number of plant-beneficial mechanisms. These can be categorized as biofertilization- plant growth promotion by enabling the bioavailability of plant nutrients and phytostimulation- the synthesis of plant growth regulators like phytohormones, and enzymes (Beneduzi *et al*., 2012). PGPE is an alternative to maximize nutrient acquisition and can be efficiently exploited as an easily accessible microbial source for enhancing agriculture production. Nitrogen fixation, solubilization of insoluble phosphates, sulfur oxidation, and iron chelation are the major pathways through which PGPEs maximize nutrient acquisition (Verma *et al*., 2001; Wakelin *et al*., 2004; Banerjee and Yesmin, 2005; Krause *et al*., 2006; Long *et al*., 2008; Santi *et al*., 2013; Yadav and Yadav, 2019). They also mediate the synthesis of vitamins that are vital for plant growth (Pirttila *et al*., 2004). In addition, rhizospheric endophytes regulate plant growth by facilitating osmotic adjustment, regulating stomatal opening and closure, modifying root morphology, improving photosynthetic efficiency, and enhancing mineral metabolism (Compant *et al*., 2005, Polesi, 2011).

**3.1.1**B**iofertilization- Nutrient Acquisition by PGPE**

Soil is the most essential medium that supplies nutrients for plant growth and development. Most often, the soil may be deficient in one or the other nutrient components required for plant growth. PGPE is well known for enhancing nutrient availability for plant growth stimulation. Endophytes colonizing the root zone facilitate the uptake of limiting plant nutrients, like nitrogen, phosphorus, and iron (Balachandar *et al*., 2006; Glick, 2012).

***3.1.1.1 Nitrogen bioavailability***

Nitrogen is an indispensable nutrient for plant growth and development. Molecular nitrogen is converted to ammonia through a fundamental process known as nitrogen fixation. Rhizospheric PGPE plays a crucial role in enhancing the availability of nitrogen for their host plant by biological nitrogen fixation through the activity of the nitrogenase enzyme. The root-associated endophytes execute efficient nitrogen fixation enabling better utilization of nitrogen in nitrogen-deficient environments (Dobereiner and Pedrosa 1987; Boddey *et al*., 1991; Reis *et al*., 2004). This PGPE includes species of the Rhizobiaceae family which colonize leguminous plant roots and independent endophytes including *Frankia* and cyanobacterial genera like *Azotobacter, Azocarus, Azospirillum*, and *Gluconacetobacter*that establish colonization of non-leguminous plant roots (Zahran, 2001; Ahemad and Khan, 2012; Bhattacharyya and Jha, 2012)*.* A number of nitrogen-fixing endophytes including *Alcaligenes* sp., *Azospirillum* sp., *Bacillus* sp., *Enterobacter* sp., *Herbaspirillum* sp., *Klebsiella* sp., *Pseudomonas* sp. and *Rhizobium* sp. were identified from rice and maize root (Patriquin *et al*., 1983; Boddey *et al*., 1995; Triplett, 1996; James *et al*., 2000). Furthermore, endophytic species of the genus *Gluconacetobacter*are excellent nitrogen fixers like the genus *Rhizobium*andare identified as symbiotic endophytes from the internal root tissues of sugarcane and pine trees (Carrell and Frank, 2014). Inoculating the crops with PGPE like *Gluconacetobacter diazotrophicus, Herbaspirillum seropedicae, H. rubrisubalbicans, Azospirillum amazonense,* and *Burkholderia* sp., is effective in enhancing nitrogen content (Oliveira *et al.,* 2002).

***3.1.1.2 Phosphate solubilization***

Phosphorus is another essential and limiting macronutrient following nitrogen. It is a significant nutrient involved in the activation of enzymes that are involved in major physiological processes governing plant growth and development (Gyaneshwar *et al*., 2002, Ahemad, 2015).  It is typically found as mineral salt in the soil and is sporadically soluble in nature. Thus, the majority of soil P is still unavailable to plants (Miller *et al*., 2010). Rhizospheric endophytes can act as the major source of phosphorus to their host plant as and when needed. Phosphate solubilization is a key feature of endophytic genera including *Bacillus, Azotobacter, Erwinia, Burkholderia, Microbacterium, Enterobacter, Flavobacterium, Pseudomonas, Serratia, Rhizobium*, etc. (Bhattacharyya and Jha, 2012). Around half the population of rhizospheric endophytes colonizing strawberries, sunflower, soybean, and other legumes are efficient phosphate solubilizers (Kuklinsky-Sobral *et al*., 2004; Forchetti *et al*., 2007; Dias *et al*., 2009; Palaniappan *et al*., 2010).

Phosphate mobilization by PGPE is achieved through precipitating the available phosphate by synthesizing organic acids. Processes like chelation and ion exchange are also modes through which phosphate solubilization is achieved by the PGPEs (Nautiyal *et al*., 2000). They can even enhance the availability of phosphorus by producing enzymes like acid phosphatase that can digest organic phosphates (Van Der Heijden *et al*., 2008).

***3.1.1.3 Potassium Solubilization***

Potassium is also an important macronutrient other than nitrogen and phosphorus. Potassium deficiency leads to abnormal development in plants causing improper root development, low viability of seeds, reduced growth, and decreased yield (Kumar and Dubey, 2012). In its original state Potassium is insoluble as it exists as a silicate mineral. Thus, the low availability of potassium is due to its insoluble nature, and plants can absorb only the dissolved form of potassium from the soil (Parmar and Sindhu, 2013). PGPE like *Pseudomonas* sp., *Bacillus edaphicus, B. mucilaginosus*, *Ewingella* sp., *Pantoea* sp., *Ferrooxidans* sp., *Paenibacillus* sp., and *Acidithiobacillus* sp., are reliable potassium solubilizers (Liu *et al*., 2012; Cun *et al.,*2022; Watts *et al.*, 2023). These bacteria efficiently solubilize minerals of potassium into plant-accessible form by secreting organic acid (Liu *et al*., 2012).

***3.1.1.4 Zinc solubilization***

Zinc (Zn) - a vital micronutrient playing a key role in plant growth and development, is essential for regulating various physiological and biochemical processes. It plays a significant role in regulating metabolic processes, enzyme activation, and redox reactions.  Zn is crucial for nitrogen assimilation, protein synthesis, carbohydrate metabolism, auxin biosynthesis, regulation of chlorophyll and other plant pigments, and modulation of cytochrome (Hafeez *et al*., 2013; Surabhi *et al*., 2022).

Persisting Zn deficiency in plants is caused by the reduced Zn availability in the soil owing to its low solubility (Singh *et al.,* 2005; Iqbal *et al*., 2010). Several factors like the presence of bicarbonate ions, excess organic matter content, increased magnesium-calcium ratio, persistent phosphate and iron content, and pH of the soil highly influence Zn availability. An increase in soil pH decreases the Zn solubility (Ahemad, 2011; Prasad *et al*., 2019). To mitigate the Zn deficiency inorganic fertilizers containing Zn serve the plants. However, a major percentage of Zn gets readily transformed into its insoluble state within a few days of application. Mere application of Zn fertilizers to the fields may not alleviate Zn deficiencies in crop plants as it gets rapidly converted into its insoluble form (Saravanan *et al*., 2004). Zinc-solubilizing microbes serve the plants by ensuring the bioavailability of Zn in the soil. PGPE actively solubilizes the insoluble form and facilitates enhanced Zn absorption by plants, subsequently promoting plant growth and yield (Prasad *et al*., 2019).

***3.1.1.5 Iron availability***

Iron is the most essential micronutrient for every life form. Major physiological processes like photosynthesis, respiration, and transpiration are regulated chiefly by enzymes and proteins with iron as a core element (Ma *et al*., 2016). Iron in its natural form - Fe3+ (ferric), retained as insoluble hydroxides or oxyhydroxides, is unavailable to both plants and microorganisms. Microbial siderophores are the main available source of iron for plants. Siderophores are low molecular weight iron-chelating molecules that bind to the insoluble ferric ions due to their high binding affinity for complexing iron (Rajkumar *et al*., 2009; Ma *et al*., 2016). They are water-soluble molecules categorized as intracellular and extracellular siderophores. Microbial siderophores are the key compounds that solubilize iron from different mineral sources under an iron deficit environment. They convert the insoluble form of iron into an easily accessible source of iron.  Rhizobacteria bear an iron-siderophore complex on the cell membrane that releases Fe2+ into the cell by the reduction of Fe3+ in the Fe3+-Siderophore complex. The siderophore molecule may be either degraded or recycled (Neilands, 1995; Rajkumar *et al*., 2010; Priyanka, 2018).

PGPE shows a strong affinity for iron complexes by secreting iron-chelating siderophores. Major siderophores produced by PGPEs include rhizobactin, enterobactin, and ferrioxamine. *Strenotrophomonas maltophilia* genome sequence revealed the ability to produce enterobactin, a catechol-type siderophore (Ryan *et al*., 2009). Endophytes that produce siderophores were identified in the root tissues of major crops (Loaces *et al*. 2011). Siderophores produced by PGPE are assimilated by plants via various methods, such as ligand-signalling reactions, chelating and degrading siderophore in order to absorb the iron component, or through direct absorption of the iron-siderophore complex. Therefore, bio-inoculation of siderophore-producing PGPE can facilitate iron absorption and mediate plant growth.

PGPEs vary in their ability to metabolize siderophores. While some are competent in siderophore utilization, others are poor in cross-linking siderophores. The second category consumes the iron-rich molecules released by other rhizomicrobes (Khan *et al*., 2009). Rhizomicrobial strains that can efficiently invade the root tissues were recognized as efficient siderophore producers. Rhizobacteria like *Escherichia coli, Pseudomonas aeruginosa, P. fluorescens, Bacillus subtilis, Brevibacillus brevis, Azospirillum brasilense*, *Azotobacter sp., Rhizobium sp., Serratia* *sp*. and *Enterobacter sp.*are active siderophore producers (Kloepper *et al*., 1980; Arora *et al*., 2001; Gupta and Gopal, 2008; Rajkumar *et al*., 2010; Saha *et al*., 2012).

Plant growth promoting endophytic siderophores have much important rhizospheric significance. They are directly involved in plant growth promotion by solubilizing iron and indirectly act as biocontrol agents by inhibiting the growth of plant pathogens (Loper and Henkels, 1999; Priyanka, 2018).  By limiting the iron availability in the plant tissues and in the surrounding soil, siderophore-producing endophytic bacteria inhibit the growth of phytopathogens including pathogenic moulds (Calvente *et al*., 2001; Ahmad *et al*., 2008). *Bacillus subtilis and Pseudomonas* *putida* efficiently inhibit the growth of wilt and dry root rot-causing fungi - *Fusarium oxysporium* (Kloepper *et al*., 1980; Patil *et al*., 2014).

Soil fungi like *Penicillium chrysogenum* and *P. aeruginosa* are actively involved in siderophore production. They aid nodule formation along with nitrogen fixation, stimulation of plant growth promotion and phytopathogen control, secretes organic acids and direct abiotic stress management (Mahmod and Allah, 2001). In addition, endophytic actinomycetes are efficiently involved in the biocontrol of the phytopathogen *Pythium*in crops like cucumber and wheat (Franco, 2006; El-Tarabily, 2009). *Trichoderma*is known to release bio-chelators that inhibit the growth of pathogenic fungi.

Siderophores are notably one of the most heterogeneous categories of microbial biomolecules, that execute a variety of tasks that come under both direct and indirect plant growth promotion and health management. They promote heavy metal stress tolerance by forming complexes with the metal ions in the soil (Rajkumar *et al*., 2010). Heavy metal-resistant rhizobacterial endophytic strains like *Pseudomonas fluorescens* G10 and *Microbacterium sp*. G16 exhibits plant growth promotion and lead accumulation in *Brassica napus* (Sheng *et al*., 2008b). PGPE like *Methylobacterium mesophilicum* and *Sphimgomonas sp*. were identified to be tolerant to heavy metals like nickel. Their endophytic habit enables them to survive under lower iron concentrations in the plant tissues (Idris *et al*., 2004). Consequently, microbial siderophore assimilation offers additional benefits to plants by inhibiting the growth of phytopathogens by limiting their nutrient and mineral availability in the soil, thereby conferring abiotic and biotic stress tolerance in addition to plant growth promotion.

**3.1.2 Phytostimulation**

Phytostimulation is the synthesis of growth-regulating compounds by plants as a result of microbial activities (Kumari *et al*., 2021). Phytostimulation primes the secretion of phytohormones and other regulatory molecules involved in plant growth promotion (Santos *et al*., 2018). Endophytic colonization plays a crucial role in the secretion of phytohormones (Gravel *et al*., 2007; Shi *et al*., 2014).  PGPE mediates direct plant growth promotion by synthesizing phytohormones like auxin, cytokinin, gibberellin (GA), and abscisic acid (Maheshwari *et al*., 2019). Indole-3-acetic acid (IAA) and ethylene biosynthesis is influenced by plant-microbial interaction (Afzal *et al.,* 2019). Auxin, cytokinins, and GA enhance root morphogenesis and improve nutrient and water uptake (Santos *et al*., 2018).

***3.1.2.1 Regulation of Indole-3-acetic acid level in plants***

IAA is the most common and naturally occurring phytohormone of the class auxin. IAA signalling functions in every aspect of plant growth and development. It induces plant cell division, differentiation, and cell elongation. IAA enhances seed and tuber germination. It plays a significant role in pigment formation and initiation of the lateral and adventitious root, besides increasing xylem development. Regulation of photosynthesis, triggering the stimuli-mediated response, activation of plant defense mechanism and modulation of secondary metabolite synthesis too comes under the purview of IAA. In addition, IAA controls cell communication and stress response in plants (Navarro *et al*., 2006; Gravel *et al*., 2007; Spaepen *et al*., 2007; Glick, 2012).

PGPE of various plants actively synthesize and secrete IAA as a secondary metabolite that enhances plant growth (Patten and Glick, 1996). In plants, IAA synthesized by root endophytes induced lateral root formation and increased root biomass (Tsavkelova *et al*., 2007, Dias *et al*., 2009). *Azotobacter, Arthrobacter, Pseudomonas, Bacillus, Acinetobacter, Agrobacterium, Rhizobium, Burkholderia*, and *Xanthomonas* are the common rhizobacterial strains that can colonize the root tissues and induce IAA production (Joo *et al*., 2005; Tsakelova *et al*., 2006; Gravel *et al*., 2007; Miter *et al*., 2021). Endophytic bacteria of the genus *Pseudomonas* are efficient plant growth promoters that are known for enhancing the host IAA level (Kang *et al*., 2014). Rhizobial strains colonizing leguminous plant roots are equally known for their IAA-mediated plant growth regulation (Ahemad and Khan, 2012). Cellular accumulation of IAA varies according to endophytic activation, which sequentially influences many of the plant’s metabolic processes. IAA mediates root nodule formation. Genetically modified *Rhizobium leguminosarum* bv. viciae with enhanced IAA biosynthetic pathway showed increased nitrogen fixation and nodule formation than the wild variety (Camerini *et al*., 2008).

PGPE not only mediates IAA synthesis but also is involved in the degeneration of IAA, correspondingly enhancing plant growth and development. *Pseudomonas putida* Strain 1290 inhibited the effect of exogenous IAA that induced root elongation in radish by degrading the derived IAA and helped to stabilize the IAA concentration benefiting their host plant (Leveau and Lindow, 2005).

**3.2 Indirect mechanism of plant growth promotion by PGPE**

An environment-friendly approach to plant disease management employed using antagonistic microorganisms is one of the indirect mechanisms of plant growth promotion which is a form of biological control against phytopathogens (Lugtenberg and Kamilova, 2009). PGPE are potential biocontrol agents that mediate indirect plant growth promotion without inducing any environmental toxicity in contrast to synthetic pesticides (MacGregor, 2006; Mishra *et al*., 2015). Endophytes survive in the protective microenvironment of the host, ensuring safer survival benefits and many additional advantages above rhizosphere microorganisms. The occurrence of a suppressive rhizomicrobiome by the plant-root endophyte interaction sustains plant growth and maintains plant health even in the presence of phytopathogen. Rhizospheric endophytes inhibit the incursion of phytopathogens through different mechanisms like the secretion of antimicrobial compounds, competitive exclusion of the phytopathogen by efficiently utilizing the microbial niche and the essential nutrients, and initiating Induced Systemic Resistance- ISR (Compant *et al*., 2005; Beneduzi *et al*., 2012; Teixeira *et al*., 2019).

The production of antimicrobial compounds like antibiotics, antifungal toxins, siderophores, hydrolytic enzymes, and volatile organic compounds is the most common microbial defense mechanism involved in the eradication of plant pathogens (Sheoran *et al*., 2015). Plant-beneficial root endophytes also facilitate ISR to protect plants by suppressing pathogens. ISR-mediated responses protect plants from bacterial, fungal, and viral diseases (Alvin *et al*., 2014). ISR is the principal plant defense mechanism that guides the host plants by inducing resistance against the pest and pathogens, even in the uninfected plant parts and ensures future protection from further pathogenic infections (Afzal *et al*., 2019). These mechanisms are beneficial against phytopathogens as they can reduce or prevent the proliferation of these pathogens through antagonistic interactions.

**3.2.1 Antibiosis**

Endophytes regulate the internal environment of the plants. They suppress pathogen growth and maintain overall plant health. Endophytes effectively reduce the intensity of virulence by vying for nutrients in the same ecological niche as that of phytopathogens and secrete chemicals toxic to plant pathogens. Root endophytes secrete antibiotics onto the endosphere, inhibiting the rhizosphere-invading phytopathogens (Gao *et al*., 2010; Bara *et al*., 2013).

Rhizospheric endophytes produce a wide range of antimicrobial compounds during the stationary phase where the cells are liable to produce secondary metabolites not involved in the active growth of the microbes. Major ones being antibiotics like coronamycin, ecomycins, kakadumycins, munumbicins, pseudomycins and xiamycins (Castillo *et al*., 2003; Ezra *et al*., 2004; Christina *et al*., 2013) and antifungal toxins like hydrogen cyanide, 2,4-diacetylphloroglucinol, tensin, phenazines, pyrrolnitrin, pyoluteorin, and viscosinamide (Bhattacharyya and Jha, 2012). These antimicrobial substances act antagonistically towards pathogens.

With their vast array of antimicrobials, endophytic bacterial communities suppress the different phytopathogens. They effectively target nematodes, viruses, bacteria, and fungal pathogens (Raza *et al.,* 2016). The bacterial endophytic population is more prominent in the root than in the aerial portions. Root colonizing endophytic bacteria from the genera *Arthrobacter*, *Actinobacteria*, *Bacillus, Enterobacter, Pantoea, Paenibacillus, Pseudomonas, Rhizobium*, and *Serratia* are known for their antimicrobial activity (Lodewyckx *et al*., 2002; Joseph *et al*., 2007; Aktuganov *et al*., 2008; Liu *et al*., 2010). *Pseudomonas* strains are recognized as direct inhibitors of phytopathogens owing to the unique range of antimicrobial compounds they produce (Weller, 2007; Khare and Arora, 2011; Mishra and Arora, 2012). *Bacillus subtilis* antimicrobials are antagonists for phytopathogen *Erwinia carotovora* (Deng *et al*., 2011). Rhizospheric endophytic bacterial strains *Bacillus megaterium* BP17 and *Curtobacterium luteum* TC10 show nematicidal activity against the phytopathogenic nematode *Radopholus similis*(Aravind *et al*., 2009).

Host-specific endophytes are more efficient biocontrol agents due to their favourable microenvironment. Rhizospheric endophytic genera including *Bradyrhizobium, Enterobacter, and Pseudomonas* were isolated from rice (*Oryza sativa*). Root colonizing PGPEs proficiently eliminate diseases in plants like maize, soybean, sorghum, millets, cowpea, sugarcane, cotton, wheat, rice, black pepper, and potatoes (Zarei *et al*., 2011; Zhang *et al*., 2012; Adeleke and Babalola*,*2020). Species of *Bacillus* are efficient endophytic biocontrol agents and reported to suppress pathogens in millets (*Pennisetum glaucum*), and *Citrus* varieties. *Enterobacter, Pantoea*, and *Pseudomonas* promote the biocontrol of pathogens in Sugarcane (*Saccharum officinarum*) (De Silva *et al*., 2019; Shen *et al*., 2019; Adeleke and Babalola*,*2020).  Additionally, PGPE that are effective against plant pests has been genetically engineered by developing *Pseudomonas fluorescens* that express Cry toxin of *Bacillus thuringiensis* and chitinase activity of *Serratia marcescens* that successfully target *Eldana saccharina* larvae (Sugarcane Borer worm) (Downing *et al*., 2000).

Endorhizal microbes act antagonistically towards phytopathogenic fungi. *Fusarium* root wilt of *Musa sp., Lycopersicon esculentum*, and *Capsicum* has been efficiently controlled by rhizospheric endophytes (Benhamou *et al*., 1998; Chen *et al*., 2011; Sundaramoorthy *et al*., 2012). *Bacillus subtilis* strains efficiently eliminated the root pathogenic fungi *Aspergillus niger* and *Fusarium oxysporum* in groundnut (Ziedan, 2006). Pseudobactin, a siderophore metabolized by pseudomonads, is effective in suppressing the growth of potato soft-rot-causing pathogenic fungi *Erwinia cartovora* (Kloepper *et al*., 1980). Similarly, pyroverdine is another class of pseudomonad-based siderophore with potential biocontrol properties.  Endorhizhal strain *Pseudomonas fluorescens* PICF7 actively curtailed the growth of the fungi *Verticillium dahlia* in olive (Prieto *et al*., 2009). Infection and proliferation of *Phytophthora infestans*late blight disease-causing pathogen, is inhibited by the antimicrobials of *Pseudomonas putida* P9 (Andreote *et al*., 2009).

Similar to the endophytic rhizospheric bacteria, endophytic fungi also maintain mutualistic relations with the host plant and negative associations with plant pathogens. Mycorrhiza-forming fungi colonize the root internal tissues and sustain strong interactions that in turn protect the plants from disease-causing agents (Verzeaux *et al*., 2017). Endophytic fungi secrete a variety of antimicrobial substances that play a significant role in controlling pest and pathogen invasion (Segaran and Sathiavelu, 2019). Entomopathogenic fungi efficiently control plant pests (Jaber and Ownley, 2018). These entomopathogenic fungi are effective biocontrol agents and are developed into commercial products. The species of *Trichoderma* exert antimicrobial activities against many pathogenic bacteria and fungi (Vizcaino *et al*., 2005). Root endophytic fungi suppress the growth of nematodes like *Meloidogyne incognita*by secretingantimicrobial compounds.  *Verticillium chlamydosporium* synthesizes phomalactone, which slows down the mobility of the nematodes within a few minutes of exposure (Khambay *et al*., 2000).

Plants release low molecular weight secondary metabolites known as phytoalexins to attract endorhizal microbes. PGPE in turn produces certain plant-signalling polyphenolic metabolites like flavonoids with enhanced antimicrobial activities. The antimicrobial activity of endophytic *Trichoderma* sp. from the *Vinca* plant is attributed to the many secondary metabolites like peptaibols, gliotoxin, gliotoxin, polyketides, pyrones, and terpenes produced by them (Vinale *et al*., 2008). Thus, the plant-endophyte interaction mediated through secondary metabolite secretion determines the rhizospheric microbial diversity. Consequently, endorhizal microbial metabolites are promising resources that can be efficiently exploited as a substitute for agrochemicals.

**3.2.2 Induced systemic resistance (ISR)**

Root colonizing PGPE efficiently inhibits the proliferation of phytopathogens causing plant diseases and helps prevent yield loss. The application of endorhizal microflora in agriculture considerably protects plants against phytopathogens. Induced Systemic Resistance (ISR) is triggered as a defense response to pathogen infection or endophytes. This in turn leads to plant-inherent resistance against pathogens. ISR facilitates plant disease resistance via multiple mechanisms. ISR stimulation alters the structural integrity of the plant cell wall by changing the composition and increasing the wall density, which inhibits pathogen invasion into the cortical tissues (Benhamou *et al*., 1996). In plants, ISR accelerates the expression of plant defensive compounds such as hydrogen peroxide, antioxidant enzymes like peroxidase, polyphenol oxidase, glutathione peroxidase, glutathione reductase, superoxide dismutase, etc. Pathogenesis-related proteins like phytoalexins, chitinase, and phenylalanine ammonia-lyase are also elevated by ISR (Harish *et al*. 2009; Gamalero and Glick, 2011).

PGPE elicits phytohormone biosynthesis as a result of ISR. ISR activates salicylic acid (SA), jasmonic acid (JA), and ethylene signalling pathways, which are usually linked together and get modulated according to ISR responses (Pieterse *et al*., 2012). In order to colonize the host plant, endophytes must also need to bypass the ISR-mediated plant defense mechanism (Ma *et al*., 2016). Root colonizing PGPE are greater contestants that induce ISR and are found to impart resistance against a broad spectrum of phytopathogens.

Endophytic strains such as *Bacillus, Pseudomonas*, and *Serratia* trigger plant defence by ISR and induce protection against different pathogens (Kloepper and Ryu, 2006; Pieterse *et al*., 2012). In *Arabidopsis thaliana,* endo bacterial strains of *Bacillus* mediate ISR by activating SA and JA-ethylene pathways against plant pathogens and inducing resistance (Niu *et al*., 2011). Endophyte inoculation in *A. thaliana* enhances ISR response conferring resistance against *P. syringae*varieties (Kavino *et al*., 2007; Kumar *et al*., 2007; Harish *et al*., 2009).

Actinomycete inhabitants of *A. thaliana*also modulate ISR-mediated defense pathway. This provides protection to the plant from the soil-borne pathogens *Erwinia carotovora* and *Fusarium oxysporum*. Nevertheless, the defense pathways differed for both pathogens.   
*E. carotovora* pathogen resistance was activated through the JA- ethylene pathway and SA mediated pathway provided resistance against *F. oxysporum*(Conn *et al*., 2008)*.*PGPE also stimulates ISR response against viral pathogens. Endophytic strains of *Bacillus subtilis*, *B. pumilus* and *B. amyloliquefaciens* facilitated ISR response against the Cucumber Mosaic Virus (CMV) (Zehnder *et al*., 2000). In bananas, endophytes evoked ISR towards Banana Bunchy Top Virus (BBTV) (Kavino *et al*., 2007; Harish *et al*., 2008).

By secreting compounds like chitinase, cellulase, lipopolysaccharides, siderophores, 2,4-diacetyl phloroglucinol, homoserine lactones, volatiles compounds like, monoterpene β-ocimene, limonene, linalool, myrcene, nonanal acetoin, 2,3-butanediol and other compounds like methyl salicylate and indole, PGPE successfully induces ISR (Yue *et al*., 2001; Lugtenberg and Kamilova, 2009; Russo *et al*., 2012; Rybakova *et al*., 2015). Mycorrhizal fungi colonizing the roots of higher plants elicit ISR via the accumulation of lignin in the cell wall of host root tissues, which act as a structural barrier against pathogens (Dehne and Schoenbeck, 1979). Endorhizal *Trichoderma sp.* induces chitinase and cellulase secretion when it comes in contact with the pathogen (Russo *et al*., 2012).

**4.**   **Application of microbial endophytes in agriculture**

Endophytes originate from the plant-associated microbial population in the rhizosphere, the region of soil around the roots (Hardoim *et al*., 2008). The diversity of microorganisms in the host plant is determined by the overall composition of the soil microbiome and influenced by the plant-specific microbial interactions (Sorensen and Sessitsch, 2006). Endophytic microorganisms are highly potent tools capable of improving plantgrowth, reducing susceptibility to plant pathogens and pests, and enhancing tolerance to plant stress post-endophytic inoculation (Watts *et al.*, 2023). Endophytes often significantly benefit associated plants by doubling their fresh mass and dry weight (Hoffmann *et al*., 2004; Govindaraj *et al*., 2017; Sheng *et al*., 2008a; Su, 2017). Endophyte-associated plants increased fresh and dry weight twofold by improved nutrient uptake during zinc and iron-deficient conditions. The presence of endophytes also increases IAA and phytosiderophore synthesis thereby improving the nutrient availability in the soil (Singh *et al*., 2018). Biofortification of crops for essential micronutrients by endophyte application is one of the practical approaches towards high-quality crop production.The most effective strategy for the application of endophytes in crop production is still being studied. The most reliable approach is to use them as soil inoculum or by seed priming (Cocq *et al*., 2017).

Owing to the complex relationship between the existing soil microorganisms and plants, recruiting plant growth-promoting rhizospheric endophytes into field crops is more challenging (Khaksar *et al*., 2016). Priming of seeds or seedlings with PGPE is an intricate process. It is highly competitive with subsequent colonizing microorganisms and is often affected by the biochemical inhibitors of the seeds **(**Gaiero *et al*., 2013**).** Introducing crop-specific plant growth-promoting rhizospheric endophytes can be a potent solution for the specific concern. On the whole, PGPE application within the initial hours of planting is more likely to establish long-term colonization and sustained presence in matured plants (Watts *et al*., 2023). Intensive research for exploiting the crop-specific rhizospheric endophytes would lead to the discovery of potential compatible sources of PGPE.

Rhizospheric endophytes perform a key role in biological nitrogen fixation, solubilization of nutrients like phosphate, potassium, zinc etc., phytohormone synthesis, plant stress tolerance and disease resistance. These beneficial microbial endophytes can be exploited to improve soil health and to enhance plant growth and productivity.

**5. Conclusion and Future Perspectives**

Root endophytes can have a significant implication on plant growth and development.  Knowledge regarding the rhizosphere-plant-microbial interactions can be exploited to improve crop productivity and maintain ecosystem sustainability. Rhizospheric microbes exhibit competitive, mutualistic as well and neutral relationships with the plants. Mutualistic plant-microbe interaction is significant as it has been greatly shown to form endophytic relationships with plants. Plant-rhizospheric endophyte relationships are influenced by the quality of plant-microbe interactions which significantly improve plant growth. The agroecosystem can be improved by employing microbial bio inoculums or engineered bioformulations or their metabolites to increase agricultural productivity. This can be executed by elucidating the mechanism underlying rhizospheric endophyte-mediated plant growth promotion that favours an eco-friendly mode of plant cultivation.

Recent advancements in agrobiotechnology and agromicrobiolgy have made it possible to modify PGPE in an efficient way that could be beneficial for agriculture. Next-generation sequencing techniques have transfigured the analysis of endophytic communities (Adeleke *et al.,* 2021).  In addition, elucidation of the relative diversity of the microbiota, genomic and proteomic techniques including metagenomics and metaproteomics, have thrown light towards the taxonomic and functional profiling of endophytic microbial communities (Adeleke and Babalola, 2022). Metabolomic and proteomics techniques also reveal the functional aspects of protein-coding genes and gene-specific metabolite production (Fadiji *et al.,* 2021). A number of efficient PGPEs have been identified but still, the appropriate technique to enhance the plant growth-promoting potential through genetic engineering remains a challenge. Moreover, the efficacy of genetically engineered PGPE may be subjected to few challenges but can exhibit vibrant agricultural significance as PGPE has the potential to revolutionize agriculture by making crops more productive and resilient.

To exploit the biological, ecological and functional approaches of endophytic rhizomicrobiome, a concerted effort with a multidisciplinary approach integrating the different advanced areas of research like agro microbiology, agrobiotechnology, nanotechnology, chemical engineering, and material science would be required. This would offer novel microbial formulations which could be employed as potential biofertilizers and biocontrol inoculants at a large scale.  These emerging techniques can also be employed to develop potential microbial consortia that would find prospective application in larger agricultural fields. Sustainable agriculture towards eco-friendly and safer food production will be a fruition in the immediate future.

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