

Review Article

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## Defendable Agricultural Boost by Profiteering Endophytes

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### ABSTRACT

Endophytes are the organisms that live within plant tissues and spend at least a half of their life time without causing disease under any circumstances. So, the endophytes are commonly defined as those organisms whose infections are inconspicuous and symptomless in host tissue and internal microbial colonization. Both prokaryote (bacteria) and eukaryotes (fungi) colonize the plant tissues exhibiting a little variation in their colonization mode. The former colonize intracellularly in vascular tissues and the later shows asymptomatic colonization either inter or intracellularly. It has been indicated that endophytes protect plants against pest attack and make them evolve molecular mechanism to face the challenges. The various climatic conditions including global warming, drought stress and deforestation are found to affect the cellular compartments, mutualistic interactions, etc. The successful isolation of endophytes relies on effective surface sterilization. Studies have also been made on the molecular characterization of endophytes including metagenomic studies, molecular markers, molecular cloning, etc. The future studies at molecular levels warranted to ascertain the protective role of PGPB against fungal pathogens.

#### Keywords

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### Introduction

Plants are the major contributors in fixing atmospheric CO<sub>2</sub> on earth. The photo energy from sunlight enables the plants in reducing carbon in CO<sub>2</sub> and produces the wide range of carbonaceous compounds. The diverse microorganisms are normally associated with plants and animals. In animals, the bacterial flora present in the gut region has made a remarkable role in stimulating immunity (Hooper *et al.*, 2011). Similarly in plants, the bacteria play a significant role in plant

defense responses (De Matos Nogueira *et al.*, 2001).

Numerous bacteria and fungi thrive on plant surfaces are called epiphytes. Certain microbes grow intimately with the plant hosts which influence plant metabolism hormonal pathways in addition to regular nutritional and biosynthetic capacities. Many of the phenotypic properties of plants are derived from the plant – microbe interaction. These naturalistic traits of microbes influence plants' interaction with competitors,

mutualists, pathogens and can play a role in growth, productivity and nutrient fluxes (Maren L. Friesen *et al.*, 2011).

### **Endophytes**

Plants harbor diverse communities of microorganisms like fungi and bacteria. These organisms are known to occur on plant surfaces called epiphytes and those present inside the plant tissues are endophytes. A Range of microorganisms including fungi, bacteria, virus and complex interaction are found within the plant tissues. As Hyde and Soy Tong (2008) mentioned, there is a doubtful series of definitions to describe an endophyte. De Bary (1866), defined endophytes as any organisms occurring within plant tissues.

Recent reviews of bacterial and fungal endophytes recommend that the term endophyte should refer to “habitat only and not function” and should include the microbes which colonize internal plant tissues for all or a part of their life cycle (Hallmann *et al.*, 2001, Schulz and Boyle, 2005).

The above said definition found to be similar to that of Hallmann *et al.*, (1997), but considers all contributing microbes. “Endophytes are microbes which occur within plant tissues for at least part of their life time without causing disease under any known circumstances.”

This statement explains that certain microorganisms which are considered endophytes in beginning may be changed to harmful. It is observed that the genomes of fungi considered as endophytes often possess plant pathogenic genes. Endophytes promote the growth and yield of plants, suppress pathogens by removing contaminants, involve in PO<sub>4</sub> solubilization and nitrogen fixation to plants.

### **History**

Endophytes were first described by the German Botanist Heinrich Friedrich Link in 1809. Endophytes are termed as “microzymas” by the French scientist BeChamp. Until 1887, it was believed that plants were healthy under sterile conditions, after which it was discovered that bacteria occur inside the plants tissues (Victor Galippe, 1887)

The word ‘endophyte’ is derived from Greek word (“Endon” =within, “Phyton” =plant) which literally means “within the plant”. The usage of this term is as board as its literal definition and spectrum of potential hosts and inhabitants. Although there are diverse uses for the word endophyte, they are most commonly defined as those organisms whose infections are inconspicuous, symptomless in host tissue and internal microbial colonisation (Stone *et al.*, 2000).

In spite of the fact that bacteria are prokaryotes and fungi are eukaryotes, they share many attributes of their association with plant hosts e.g. both colonize root tissues inter and intra-cellularly and often systemically. However, they show a little variation in their modes of colonization i.e. bacteria primarily colonies intracellularly (HallMann *et al.*, 1997) and are frequently found in the vascular tissues of host plants (Kobayashi and Palumbo 2000). In case of fungi, asymptomatic colonization may be inter and intra-cellular throughout the root.

### **Impact of endophytes on plants**

Generally, it is considered that the bacteria are present within plants tissues which may or may not be pathogenic. The non-pathogenic flore occur in root tissues (Perotti, 1926). Presence of bacteria was observed in leaves, stem and roots of healthy plants (Hennig and

Vill forth, 1940). Beijerinck (1888) described the colonization of rhizobia in legume root tissues. Shekhawat *et al.*, (1984), reported the existence of both pathogenic and non-pathogenic endophytic bacteria within potatoes and triggering of disease development by an external event. Certain bacterial species may become pathogenic under certain conditions (Hayward, 1974).

Bacterial endophytes reside inside plants for at least part of their life cycle (Hallmann *et al.*, 1997). A series of articles indicate the benefits of selective fungal or bacterial endophytes in plant protection against pest attack.

The bacterial invasion in plants could have a major impact on growth and health of plant and the plants are supposed to evolve molecular mechanisms to deal with the challenges posed by invaded bacteria.

Severe environmental constraint to agricultural productivity is caused by drought stress. In the modern agriculture system the fertile nature of soil is often maintained by applying bio fertilizers and usage of agrochemicals has been increased to control pathogens and pests.

### **Effect of climate on endophytes**

The population of endophytes varies from plant to plant, species to species and also the same species from one region to other. The variation also occurs with change in climatic condition of same region. In total endophytic fungi, the temporal changes in relative frequency were studied (Chareprasert *et al.*, 2006).

The additional consequences of global warming may be the effect of drought stress which on AMF has been reviewed thoroughly by Auge (2001). The multidimensional stress

drought affects various sub cellular compartments, cell organs and the whole plant level (Choluj *et al.*, 2004, Rahadari *et al.*, 2012). So the quantity and quality of growth in plants is negatively affected due to drought. The migration of drought stress is necessary to achieve designated goals to increase the food production.

Drought is often responsible for plant growth reduction and it also affects both root and aerial plant parts. This causes changes not only in allocation of photosynthates in the rhizosphere but also in extra mycorrhizal mycelia formation. Drought does not affect the density of extra mycorrhizal hyphae as demonstrated with *Glomus sp.* But under drought condition more hyphae were produced when expressed relative to root length (Staddon *et al.*, 2004). This may be explained by the fact that part of mycelium might be dead (Staddon *et al.*, 2003) and so before drought most hyphae have been produced (Staddon *et al.*, 2004).

Some endophytes are seen to be the latent pathogens and only at certain condition the infection proceeds. This occurs due to the environmental conditions such as CO<sub>2</sub> accumulation or O<sub>2</sub> depletion (Lund and Whyatt, 1972)

The greater number of genera and species with higher colonization frequency were found in matured leaves of teak (*Tectone grandis L*) and rain tree (*Samanea saman*) than those in young leaves. During rainy season their occurrence in leaves increased. The endophytic population and frequency tended to differ among sampling dates for all organs studied, namely young petioles and twigs of *Gingko biloba* (Thongsandee *et al.*, 2012) The temperature stress in plants is alleviated by some rhizosphere bacteria and endophytes and these strains induce the growth promotion of different crops at

different climates, soil, temperature (Bilal *et al.*, 1993, Javed and Arsha, 1997).

Endophytes play a major role in mutualistic interactions, enhancing host plant's nutrient uptake and helping host to counteract adverse effects of biotic and abiotic stress (Idris *et al.*, 2007).

While comparing with rhizospheric bacteria or bacterial pathogens, the endophytic bacteria occur at lower population densities (Hallmann *et al.*, 1997; Rosenblueth and Mastineze Romero, 2004) Endophytic population like rhizospheric population is conditioned by biotic and abiotic factors (Seghers *et al.*, 2004) When compared to rhizospheric bacteria the endophytic bacteria could be better protected from biotic and abiotic stress (Hallmann *et al.*, 1997)

Egamberdiyeva and Hoflich (2003) reported that temperature and soil type may affect the performance of the plant beneficial bacteria. From semi-continental climate the *Mycobacterium* sp. 44 and *Pseudomonas fluorescens* and *Pantoea agglomerans* were isolated and found to increase the root and shoot growth in winter wheat at 16° C compared with that 26°C in loamy sand. Rain forest destruction may lead not only to the loss of valuable tree species it also affects the unknown endophytes, especially fungi (Strobel *et al.*, 2004).

The major role played by drought stress is it makes physico-chemical and biological properties of soil in microbial activity and crop yield. Water availability controls the production and consumption of protein and polysaccharides by the bacteria (Roberson and Firestone, 1992) and thus indirectly influence soil structure. The microbes produce Exopolysaccharide (EPS) which protects them from inhospitable conditions and enables their survival. Capsular material

of *A. brasilense* sp 245 contain high molecular weight carbohydrate complex (Lipopolysaccharide protein (LP) complex and Polysaccharide (PL) complex responsible for the protection under extreme conditions like desiccation. These complex addition to a suspension of decapsulated cells of *A. brasilense* sp 245 significantly enhanced survival under drought stress (Konnova *et al.*, 2001). The drought stress causes changes in plant associated communities. The different subpopulations of endophytic bacteria colonizing sunflower cultivated under drought condition (or) under irrigation management were identified recently (Forchetti *et al.*, 2007) Under drought condition, one particular *Achromobacter* strain was found which indicates a better adaptation to drought stress condition. Interestingly, a higher plant growth promotion potential was found in endophytic bacteria isolated from sunflower cultivated under drought when compared with irrigated plant (Forchetti *et al.*, 2007).

### **Isolation of endophytes**

Both Monocotyledonous and Dicotyledonous plants contain endophytic bacterial species, ranging from woody tree species such as oak, pear to herbaceous crop plant such as sugar beet and maize (Miche and Balandreau, 2001) Totally 2003 fungal endophytes were isolated from 750 surface sterilized seeds. In contrast, only 16 endophytic isolates were obtained from 800 surface sterilized seeds (J. Hallmann 1997).

The microbes such as endophytic bacteria, fungi and actinomycetes whose isolation from the plant tissue has been a challenge since the studies on endophytes begins. Several researchers have reviewed different methods of the separation of bacterial endophytes (Hallmann and Reinhold, 1998). An initial surface sterilization is used to isolate endophytes followed by culturing from

ground tissue extract (Rai, 2007) (or by direct culturing of plant tissues on media suitable for bacteria or fungi or actinomycetes (Hata 2008).

*Lodewyckx et al.*, (2002) review highlights the method used to isolate and characterize endophytic bacteria from various plant species. A complete list of bacterial endophytes isolated from a broad range of plants is provided by Rosenblueth and Martinez – Roncero (2006) and Berg and Hallmann (2006) which update the ground work laid by Hallmann *et al.*, (1997) and Lodewyckx *et al.*, (2002).

The endophytes such as *Pleurostoma*, *Chaetomium*, *Coniochaeta*, *Daldinia*, *Xylaria*, *Hypoxylon*, *Nodulisporum*, *Cazia* and *Phellinus* isolated from *Hyperzia serrata* were confirmed for the first time by rDNA ITS analysis (Chen, 2011).

### **Surface sterilization**

To examine the efficiency of Sodium hypochlorite, Ethanol and Mercuric chloride as effective sterilizing agent and survival percentage of treated explants, and to detect the effect on explants treated individually and by the different combining of chemical disinfectant. Sterilization procedures that were at first selected and used were not found efficient agent individually.

However various combination and duration of 70% of Ethanol, 2% of Sodium hypochlorite and 0.1% of Mercuric chloride were applied to the explants to achieve a satisfactory result (Nawed Anjum *et al.*, 2015)

The surface sterilization of plant parts with 70% of Ethanol for 1 minute, 2% of Sodium hypochlorite for 4 minutes and 0.1% of Mercuric chloride for 4 minutes, ethanol for 30 seconds finally 3 rinses in distilled water (Nawed Anjum *et al.*, 2015).

### **Isolation**

Many varieties of dicotyledonous plants were collected and transported to refrigerated box at 4°C to the laboratory. After surface sterilization, the sample extracts from plants made upto 10 ml using sterile distilled water. From that, 1 ml was taken and serially diluted using test tubes containing 9 ml of sterilized distilled water. 0.1 ml from the dilutions  $10^{-5}$ ,  $10^{-6}$  and  $10^{-7}$  were transferred to petriplates containing Nitrogen free malate medium (NFB) (Dobereiner, 1992) and Kings' B medium (Kaare Johnsen and Preben Nielson 2006) for the isolation of *Azospirillum* and *Pseudomonas* species respectively.

Meanwhile, 1 ml of extract was dissolved in semi solid NFB to observe the sub – surface pellicle formation by *Azospirillum* species and kept at room temperature. The separated colonies in petridish were brought to pure culture by several sub culture. Pellicle of *Azospirillum* species in test tubes containing semi solid NFB was streaked in petri plates containing semi solid NFB. The purified strains were maintained in Nutrient agar slant and stored at 4°C for future use.

### **Molecular studies on endophytes**

The molecular characterization of endophytes was subjected to the purpose of identification at the molecular level which includes the metagenomic studies, use of molecular markers, molecular cloning and gene expression studies. Denaturing gradient gel electrophoresis (DGGE) profiles of 16s rRNA gene fragments amplified from total plant DNA were used to detect some non culturable endophytic bacteria by comparing the profile with the bands obtained from the culturable endophytes from citrus plants (Araujo, 2002).

Bacterial endophyte community of potato (*Solanum tuberosum*) was examined by using bacterial automated ribosomal intergenic

spacer analysis (B-ARISA) technique (Marter, 2010). To determine the richness of bacterial operational taxonomic units (OTUS), the pyro sequencing was used. The author included that metagenomic analysis can complement PCR based analysis yield information on whole gene operons (Nikolic, 2011).

Molecular studies in endophytes have gone to the extent of complete genome in *Enterobacter sp.* An endophytic plant growth promoting gamma-proteo bacterium isolated from the stem of polar, a potentially important biofuel feed stock plant was sequenced (Taghavi, 2010).

In contrast to the extensive information on the molecular mechanism of bacteria – plant interactions (Lugtenberg *et al.*, 2002) (Oldryd and Dounie 2004). There is only limited data on the endophyte - host molecule interaction. By the presence of bacteria, plant genes may be modulated and the genes so expressed provide clues to the effects of endophytes in plants.

The gene expression in response to the endophytic colonization of *Gluconacetobacter diazotrophicus* and *Herbaspirillum rubrisubalbicans* are being studied in sugarcane (De Matos Nogueira *et al.*, 2001).

### **Isolation of DNA**

Around 0.5 ml of sample was taken and placed in a mortar and homogenized with 2 ml of extraction buffer.

The extraction buffer (PH 8.0) contains 100 m M Tris, 20 m M EDTA, 0.5 M NaCl, 7M urea, 0.1%  $\beta$  – mercapto ethanol and 25 of SDS.

Take an equal volume of Phenol: Chloroform: Isoamyl alcohol (25:24:1) was added to the

tubes and mixed well by gently shaking the tubes.

The tubes are allowed to centrifuge at room temperature for 15 minutes at 15000 rpm. The upper aqueous phase (supernatant) was collected in a new tube and an equal volume of Chloroform: Isoamyl alcohol (24:1) was added and mixed.

The supernatant obtained after centrifuging at room temperature for 10 minutes at 15000 rpm was transferred to new tubes.

By adding 0.1 volume of 3M sodium acetate, PH – 7.0 and 0.7 volume of Isopropanol for the purpose of DNA precipitation. After the tubes were incubated for 15 minutes at room temperature, the tubes were centrifuged at 4° C for 15 minutes at 15000 rpm.

The DNA pellet was washed with 70% ethanol for twice, then very briefly with 100% ethanol and allowed to air dry. Then the DNA was dissolved in TE (Tris – cl 10m M PH = 8.00 and EDTA 1m M) to remove RNA 5 micro liter of DNase from RNase A (10 mg / ml) was added to the DNA (52).

### **PCR analysis**

AP – PCR analysis: AP – PCR amplification was carried out in a volume of 25 micro liter containing 100ng of template DNA, 2 m M MgCl<sub>2</sub>, 5 $\mu$ m 27 f primer, 2.5  $\mu$ l of 10 X assay buffer which includes (10 m M Tris PH 9), 50 m M Kcl, 1.5 m M Mgcl<sub>2</sub> and 0.01% Gelatin) 10 m M each of dNTPS and 5 units/  $\mu$ l of Taq DNA polymerase.

1 cycle of 5 minutes at 94°C for denaturation, 5 minutes at 45°C for annealing and 5 minutes at 72°C for extension and 35 cycles of 1 minute at 94°C, 1 minute at 45°C and 1 minute at 72°C followed by a final 10 minutes extension at 72°C.

**Table.1** Possible endophytes

Plant type	Endophytes present in plants	Reference
Sugarcane ( <i>Saccharum officinarum</i> )	<i>Herbaspirillum seropedicae</i>	Dong <i>et al.</i> , 1994
Corn ( <i>Zea mays</i> )	<i>Enterobacter sp.</i> , <i>Pseudomonas sp.</i> , <i>Klebsiella sp.</i> , <i>Vibrio sp.</i> ,	Fisher <i>et al.</i> , 1992
Rice ( <i>Oryza sativa</i> )	<i>Azorhizobium caulinodans</i>	Engelhard <i>et al.</i> , 2000
Banana ( <i>Musa spp.</i> )	<i>Citrobacter</i> , <i>Azospirillum brasilense</i>	Martinez <i>et al.</i> , 2003
Cotton ( <i>Gossypium spp.</i> )	<i>Agrobacterium sp.</i> , <i>Serratia sp.</i> , <i>Burkholderia sp.</i> , <i>Bacillus sp.</i> , <i>Staphylococcus</i> , <i>Rhizobium sp.</i> , <i>Variovorax sp.</i> , <i>Pseudomonas sp.</i> , <i>Acenitobacter sp.</i> , <i>Arthrobacter.</i> , <i>Enterobacter</i>	MC Inroy and Kleopfer, 1995
Sweet potato ( <i>Ipomoea batatas</i> )	<i>Enterobacter arburiae</i>	Asis and Adachi 2003
Tomato ( <i>Solanum lycopersicum</i> )	Pseudomonadaceae	Samish <i>et al.</i> , 1963
Carrot ( <i>Daucus carota</i> )	<i>Pseudomonas fluorescens</i>	Surette <i>et al.</i> , 2003
Citrus plant	<i>Alcaligenes sp.</i> , <i>Bacillus cereus</i> , <i>B. lentus</i> , <i>B. magaterium</i> , <i>B. pumilus</i> , <i>B. subtilis</i> , <i>Burkholderia cepacia</i> , <i>Curtobacterium flaccumfaciens</i> , <i>Enterobacter cloacae</i> , <i>Methylobacterium extorquens</i> , <i>Pantoea agglomerans</i> .	Araujo <i>et al.</i> , 2001
Soy bean ( <i>Glycine max L.</i> )	<i>Klebsiella oxytoca</i>	Kuklinsky – sobral <i>et al.</i> , 2004
Cucumber ( <i>Cucumis sativus</i> )	<i>Pseudomonas sp.</i> , <i>Bacillus sp.</i> , <i>Enterobacter sp.</i> , <i>Arthrobacter sp.</i> , <i>Agrobacterium sp.</i> , <i>Burkholderia sp.</i> , <i>Stenotrophomonas sp.</i> , <i>Chryseobacterium sp.</i> ,	Kloepfer, 1983
Lettuce ( <i>Lactuca sativa</i> )	<i>Escherichia coli</i>	Ingham <i>et al.</i> , 2005
Yellow lupine, citrus plant	<i>Burkholderia cepacia</i>	Araujo <i>et al.</i> , 2001, Barac <i>et al.</i> , 2004
Alfalfa, carrot, raddish, tomato	<i>Salmonella enterica</i>	Cooley <i>et al.</i> , 2003, Guo <i>et al.</i> , 2002, Islam <i>et al.</i> , 2004
Rice	<i>Serratia marcescens</i>	Gyneshwar <i>et al.</i> , 2001
Wheat ( <i>Triticum aestivum</i> )	<i>Azospirillum sp.</i> ,	Webster <i>et al.</i> , 1997
Rice ( <i>Oryza sativa</i> )	<i>Azoascus sp.</i> , <i>Herbaspirillum sp.</i> ,	Reinhold-Hurek and Hurek, 1997, 1998
Sugarcane, coffee	<i>Gluconoacetobacter diazotrophicus</i>	Cavalcante and Dobereiner 1988, Jimenez Salgado <i>et al.</i> , 1997
Banana, rice, maize, sugarcane	<i>Klebsiella variicola</i>	Rosenblueth <i>et al.</i> , 2004
Sugarcane ( <i>Saccharum officinarum</i> )	<i>Herbaspirillum rubrisulbabicans</i>	Olivares <i>et al.</i> , 1996 (80)
Marigold ( <i>Calendula officinalis</i> )	<i>Microbacterium esteraromaticum</i>	Sturz and Kimpinski 2004 (81)
Sweet potato ( <i>Ipomoea batatas</i> )	<i>Paenibacillus odosifer</i>	Reiter <i>et al.</i> , 2003 (82)

### 16s rRNA sequence determination

The amplified products of approximately 1461 bp of *Azospirillum sp* and 1341 bp of *Pseudomonas sp* were detected and sequenced by 16s primer containing 27 f (forward primer) and 1492 (reverse primer). The sequencing products were purified and the results were observed.

Finally the sequence can be compared with sequence in the NCBI data bank using the BLAST program.

### Future research

The most challenging sectors to climatic change are done by agriculture. Future research is to be focused on developing microbial formulation to boost plant performance under drought stress which reduces the chemical fertilizers and pesticides usage (Kaushal and Wani, 2015). Further studies at the molecular and biochemical levels are warranted to ascertain the protective role of these PGPB isolates against fungal pathogens.

Though some root endophytes are latent pathogens, many others (root endophytes) provide benefits to the plant growth by helping in phosphorus uptake and protecting against pests and diseases (Sieber, 2002).

### Possible endophytes

The possible endophytes in various plants have been reported by scientists at different time period and a few such organisms have been listed in the table 1.

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