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(54) Title: COMPOSITIONS AND METHODS COMPRISING ENDOPHYTIC BACTERIUM FOR APPLICATION TO GRASSES TO INCREASE PLANT GROWTH, SUPPRESS SOIL BORNE FUNGAL DISEASES, AND REDUCE VIGOR OF WEEDY COMPETITORS

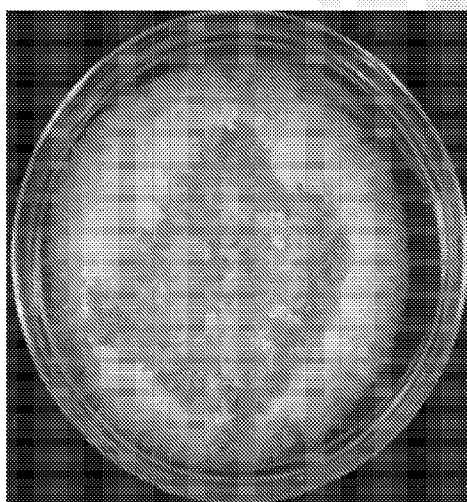


Fig. 1A

(57) Abstract: Endophytic bacteria, compositions comprising the same, and methods of use thereof are disclosed which increase the root and shoot growth of grass hosts, suppress growth of soil borne fungal pathogens of host plants, increase resistance of the grass to diseases and reduce competitiveness of distantly related competitor weeds of the crop.



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5 This application claims priority to US Patent Application 15/657,100 filed July 21, 2017,
the entire contents being incorporated herein by reference as though set forth in full.

Sequence Listing

The instant application contains a Sequence Listing with 16 sequences which has been submitted via EFS-Web and is hereby incorporated by reference in its entirety. Said ASCII
10 copy, created on July 20, 2017 is named SEQLIST and is 15kb in size.

Field of the Invention

This invention relates to the fields of plant biology and endophytic bacteria. More specifically, the invention provides new strains of endophytic bacteria which provide beneficial features to the plant upon colonization of the same.

15 **Background of the Invention**

Several publications and patent documents are cited throughout the specification in order to describe the state of the art to which this invention pertains. Each of these citations is incorporated by reference herein as though set forth in full.

Invasive *Phragmites australis subsp. australis* (Cav.) Trin., native to Eurasian wetlands, is an introduced grass in North America that displaces entire communities of native flora and fauna. This plant is highly competitive and generally out-competes other plants to produce large monospecific stands. It is increasingly becoming clear that microbes that form symbiotic associations with *Phragmites* play roles in increasing its invasive character (Clay et al., 2016). Soares et al. (2016a) recently demonstrated that endophytic bacteria isolated from tiller meristems were capable of increasing nitrogen assimilation into plants in greenhouse experiments. In another study, Soares et al. (2016b) showed that fungal endophytes that enter into roots of *Phragmites* growing in saline soils may enhance salt tolerance in the host, enabling

it to thrive in high saline soils. Ernst, Mendgen and Wirsal (2003) demonstrated that a seed-borne fungal endophyte in genus *Stagonospora* enhanced biomass accumulation in experiments in microcosms. It is thus evident that symbiotic microbes may impact the aggressiveness of *Phragmites*.

Summary of the Invention

In accordance with the present invention, a method of increasing root and shoot growth of a grass plant host, comprising inoculating the host grass or seed with one or more biologically pure *Pseudomonas spp.*, *Microbacterium spp.* or *Enterobacter spp.* endophyte strains isolated from *Phragmites australis*, which are heterologously disposed to said grass plant host, and effective to increase one or more of root growth promotion, shoot growth promotion, resistance to salt stress, competition with undesirable plant species and resistance to fungus in a plant produced from the seed, when compared to a reference plant grown under the same conditions. In certain embodiments, the bacteria comprise specific rDNA sequences selected from the group consisting of SEQ ID NO: 1-16 or sequences having at least 90%, at least 95%, or at least 98% identity therewith. In a particularly preferred embodiment, the endophytic bacteria comprise one or more of Sandy LB4, Sandy LB6, Sandy Y8, RoY12, RoLB13w, WY9y, WY6, West 9, WY14, RiY3, SandyY7w, RiLB4, SY1, SY5, WY9w, and *Microbacterium oxydans* B2. The aforementioned method can also be used to advantage to suppress growth of soil borne fungal pathogens of said host grass relative to host grasses which have not been inoculated with the one or more endophyte. Plants treated with the method may also exhibit increased resistance to disease relative to untreated host plants. Inoculated plants also are effective to reduce competitiveness of distantly related competitor weeds of said grass plant host.

The host grass includes, without limitation, a turf and forage grass selected from *Agrostis spp.*, *Poa spp.*, *Festuca spp.*, *Lolium spp.*, *Cynodon spp.*, *Zoysia spp.*, *Koeleria spp.*, *Paspalum spp.*, *Panicum spp.*, *Stenotaphrum spp.*, *Eremochloa spp.* and *Danthonia spp.*, *Triticum aestivum*, *Oryza sativa*, *Avena sativa*, *Hordeum vulgare*, *Panicum miliaceum*, *Eleusine coracana*, *Sorghum bicolor*, *Pennisetum glaucum*, *Setaria italica*, *Zea mays*. *Agrostis spp.*, *Poa spp.*, *Festuca spp.*, *Lolium spp.*, *Cynodon spp.*, *Zoysia spp.*, *Koeleria spp.*, and *Danthonia spp.* In a preferred embodiment of the method, the host grass is a food grass crop selected from *Oryza sativa*, *Triticum aestivum*, and *Zea mays*. In yet another preferred embodiment, the host grass is a Pooideae grass. In certain aspects of the invention, the bacteria are present in a synthetic seed

ball. Plants may also be sprayed with a liquid formulation comprising the bacteria. In another embodiment, the bacteria are present in a controlled release fertilizer formulation.

The invention also encompasses a synthetic combination comprising at least one purified bacterial population in association with a seed of a turf and forage grass, wherein the purified bacterial population comprises a seed bacterial endophyte that is heterologously disposed to the turf or forage grass seed and is selected from one or more of Sandy LB4, Sandy LB6, Sandy Y8, RoY12, RoLB13w, WY9y, WY6, West 9, WY14, RiY3, SandyY7w, WY9w, RiLB4, SY1, SY5 and *Microbacterium oxydans* B2, the seed bacterial endophyte being present in the synthetic combination at a concentration of at least 10^3 CFU/seed on the surface of a seed, and in an amount effective to increase one or more of root growth promotion, shoot growth promotion, resistance to salt stress, competition with undesirable plant species and increase resistance to fungus in a plant produced from the seed, when compared to an untreated reference plant grown under the same conditions. In this embodiment of the invention, the bacteria may be lyophilized, encapsulated in an alginate bead formulation, or present in a seed ball. Grass seed to be treated, include without limitation, seeds from a grass selected from *Agrostis spp.*, *Poa spp.*, *Festuca spp.*, *Lolium spp.*, *Cynodon spp.*, *Zoysia spp.*, *Koeleria spp.*, and *Danthonia spp.* In other preferred embodiments *Oryza sativa*, *Triticum aestivum*, *Avena sativa*, *Hordeum vulgare*, *Secale cereal*, and *Zea mays* are treated. In a one embodiment, the is *Microbacterium oxydans* B2 and said seed is a rice seed.

In another aspect of the invention, a method of reducing undesirable weed growth in soils comprising turf and forage grasses, comprising drenching or treating the soil with one or more biologically pure *Pseudomonas spp.* endophyte strains wherein said strains reduce weed growth relative to weed growth observed in untreated soil is provided. In one embodiment, the one or more *Pseudomonas spp.* is Sandy LB4 and/or West 9.

In certain aspects, the synthetic combination comprises a combination of bacteria selected from the group consisting of i) protease positive strains consisting of Sandy LB4 or Sandy LB6 in combination with strains SY1 or SY5; ii) protease positive strains WY6 or WY14 in combination with protease negative strain WY9y and iii) phosphatase negative strain *Microbacterium oxydans* B2 in combination with phosphatasepositive strains Sandy LB4 or Sandy LB6 which can act synergistically to increase root and shoot growth of grass plant hosts. An exemplary method comprises artificially inoculating the host grass with one or more

biologically pure *Pseudomonas spp.* endophyte strains wherein said strains confer protection to the host grass. Preferably, said one or more *Pseudomonas sp.* is Sandy LB4 and West 9 used alone or in combination.

5

Brief Description of the Drawings

Figures 1A-1D. Fungus without bacteria (Fig. 1A) *Pseudomonas fluorescens* Sandy LB4, (Fig. 1B) *Pseudomonas sp.* West 9 (Fig. 1C) and *Microbacterium oxydans* B2 (Fig. 1D) were co-cultured on LBA with soil fungi *Sclerotinia homeocarpa*, (the causal agent of Dollar spot disease in turfgrass) for 7 days. Formation of zones of inhibition are shown.

10 **Figures 2A-2B.** Fig. 2A. *Cynodon dactylon* (Bermuda grass) Seedlings growing on 0.7% agarose and stained with DAB for reactive oxygen. Fig. 2B. Bermuda grass (*Cynodon dactylon*) grass seedling root in agarose without bacteria showing absence of root hairs.

Figures 3A-3B. Fig. 3A. Bacterial strain West 9 colonizes the Bermuda grass seedling root-tip meristem and becomes intracellular in meristem cells then transmits to all parts of the seedling root and root hairs as the meristem cells divide. In these seedlings root hairs form very close to the root tip. West 9 restores root hair development in sterile seedlings. Root stained in agarose using DAB (diaminobenzidine) to visualize reactive oxygen (H₂O₂) production around bacteria. Fig. 3B. Root hairs (arrow) behind the root-tip are seen to stain brown internally due to production of reactive oxygen in the vicinity of the intracellular bacteria.

20 **Figures 4A-4E.** Fig. 4A. Early developing root hairs show internal presence of the spherical wall-less L-forms (arrows) of bacterial strain West 9. Roots were stained with DAB then counterstained with aniline blue stain. Fig. 4B. Early developing root parenchyma cells also show chains of the spherical L-form bacteria. The root was stained with DAB then counterstained with aniline blue stain. Fig. 4C. Cells from the root meristem show internal presence of bacteria (arrows). Fig. 4D. The bacteria are degraded internally. They are seen to swell and become clear internally (arrows). Clusters of degraded bacteria are seen in these root parenchyma cells. Fig. 4E. More degrading bacteria (arrows) in root parenchyma cells. Bacterial L-forms swell as their internal contents vanish due to degradation.

Figures 5A-5B. Fig. 5A. As the root continues to develop the bacteria within root hairs and root parenchyma cells are seen to disappear, apparently due to complete degradation by the root cells. Here several root hairs are seen to be free of internal bacteria (arrows). Bacteria that were intercellular are not degraded and remain as endophytes in the plant. Fig. 5B. In older parts of the root all root hairs are seen to be free of internal bacteria and reactive oxygen staining (brown coloration). In the parenchyma cells of the root axis, the brown coloration also lessens due to degradation of the internal bacteria.

Figures 6A- 6C. Fig. 6A. *Poa annua* seedling root in agarose without bacteria showing absence of root hairs. Stained using DAB to visualize reactive oxygen. Fig. 6B. *Poa annua* seedling root bearing strain West 9, showing production of root hairs. Fig. 6C. *Poa annua* root hair showing internal bacteria (arrows).

Figures 7A-7B. Fig. 7A. Strain Sandy LB4 (also from *Phragmites*) is also a meristem colonizer. It is shown here in a Bermuda grass seedling root. Fig. 7B. The bacteria appear to persist longer in Bermuda grass seedling roots than strain West 9. It appears that the effect of Sandy LB4 in Bermuda grass seedlings is of longer duration than that of West 9.

Figures 8A-8C. Fig. 8A. *Rumex crispus* (Curly dock). Fig. 8B. Strain Sandy LB4 from *Phragmites* in root hair of *Rumex crispus*. Root hair blockage may affect the absorptive function of root hairs. This may reduce the competitive ability of seedlings. Fig. 8C. Normal root hair function involves internal cytoplasmic streaming. Cytoplasmic streaming cessation in blocked hairs may reduce nutrient absorption.

Figures 9A-9B. Fig. 9A. Dandelion seedlings (3-wk-old) showing high mortality when colonized by *Phragmites* bacterium Sandy LB4. Fig. 9B. Dandelion seedlings from inhibition experiment in soil.

Figure 10. Exemplary salt stress experiments in magenta boxes demonstrating that Sandy LB4, Sandy LB6 and SY1 are effective to increase salt stress tolerance in rice and Bermuda grass (*Cynodon dactylon*).

Detailed Description of the Invention

Phragmites australis is an invasive grass that decreases biodiversity and produces dense stands in North America. In the present invention, we tested the hypothesis that invasive *Phragmites australis* establishes mutualisms with endophytic microbes that increase its growth, improve its ability to survive stressful environments and resist pathogens, and contribute to its competitive capacity against other plant species. Ten bacterial strains were isolated from the surfaces of *Phragmites* seeds collected from natural stands at four sites in New Jersey. 80% of the isolates were identified to genus *Pseudomonas*. Using a surrogate test grass *Cynodon dactylon* in seedling assays we determined that two of the isolated strains Sandy LB4 (*Pseudomonas fluorescens*; deposited as accession number NRRL-B67308) and West 9 (*Pseudomonas* sp.; deposited as accession number NRRL-B67307) were endophytes that enter into plant meristems and become intracellular within cells of root tissues. These bacterial strains, along with an endophyte (*Microbacterium oxydans* B2; deposited as accession number NRRL-B67470) isolated from *Phragmites* shoot tissues, were found to increase plant growth using the surrogate grass host *Poa annua*. These strains were also found to slightly improve seed germination and increase seedling root branching in assays using *Poa annua*. Strains Sandy LB4 and West 9 were further shown to protect plants from damping off disease caused by fungal pathogen *Fusarium oxysporum* using *Poa annua* as the surrogate host plant in magenta box experiments. To evaluate whether *Phragmites* bacteria might play a role in inhibiting growth of competitor plant species we conducted experiments using dandelion (*Taraxacum officinale*) and curly dock (*Rumex crispus*). In these experiments we found that one of the endophytic strains, Sandy LB4, has the capacity to increase mortality in seedlings of both competitor species. Further, Sandy LB4 reduces the growth rates of both competitor plant species and increases their susceptibility to diseases.

I. DEFINITIONS:

An "endophyte" or "endophytic microbe" is an organism that lives within a plant or is otherwise associated therewith. Endophytes can occupy the intracellular or intercellular spaces of plant tissue, including the leaves, stems, flowers, fruits, seeds, or roots. An endophyte can be either a bacterial or a fungal organism that can confer a beneficial property to a plant such as an

increase in yield, biomass, resistance, or fitness in its host plant. As used herein, the term "microbe" is sometimes used to describe an endophyte.

Several strains of the bacteria described herein can be identified by their distinct ribosomal 16S sequences. 16S ribosomal RNA (or 16S rRNA) is the component of the 30S
5 small subunit of a prokaryotic ribosome that binds to the Shine-Dalgarno sequence. The genes coding for it are referred to as 16S rRNA gene and are used in reconstructing phylogenies, due to the slow rates of evolution of this region of the gene. Ribosomal RNA sequences from the bacteria described herein are provided in SEQ ID NOS: 1- 16

As used herein, "sequence identity" generally refers to the percent identity of nucleotide
10 bases or amino acids comparing a first polynucleotide or polypeptide to a second polynucleotide or polypeptide using algorithms having various weighting parameters. Sequence identity between two polynucleotides or two polypeptides can be determined using sequence alignment by various methods and computer programs (e.g., BLAST, CS-BLAST, FASTA, HMMER, L-ALIGN, and the like) available through the worldwide web at sites including but not limited to
15 GENBANK (on the world wide web at ncbi.nlm.nih.gov/genbank/) and EMBL-EBI (on the world wide web at ebi.ac.uk). Sequence identity between two polynucleotides or two polypeptide sequences is generally calculated using the standard default parameters of the various methods or computer programs. A high degree of sequence identity, as used herein, between two polynucleotides or two polypeptides is typically between about 90% identity and
20 100% identity, for example, about 90% identity or higher, preferably about 95% identity or higher, more preferably about 98% identity or higher. A moderate degree of sequence identity, as used herein, between two polynucleotides or two polypeptides is typically between about 80% identity to about 85% identity, for example, about 80% identity or higher, preferably about 85% identity. A low degree of sequence identity, as used herein, between two polynucleotides or two
25 polypeptides is typically between about 50% identity and 75% identity, for example, about 50% identity, preferably about 60% identity, more preferably about 75% identity.

The terms "promoting plant growth" and "stimulating plant growth" are used interchangeably herein, and refer to the ability to enhance or increase at least one of the plant's height, weight, leaf size, root size, shoot length, stem size, competition with competitor plants,
30 resistance to fungal infection, increased protein yield from the plant or increased grain yield of the plant.

Particular formulations to be applied in spraying forms such as water dispersible concentrates or wettable powders may contain surfactant such as wetting and dispersing agents, e.g., the condensation product of formaldehyde with naphthalene sulphonate, an alkyl-aryl-sulphonate, a lignin sulphonate, a fatty alkyl sulphate an ethoxylated alkylphenol and an ethoxylated fatty alcohol.

As used herein the terms "spray" or "spraying" include the technique of applying to an exterior surface an ejected liquid material.

As used herein, the terms "coat" or "coating" include application, typically of a liquid or flowable solid, to an exterior surface such as a seed.

As used herein, a "stabilizer" includes a chemical compound that can be added to a formulation to prolong the stability and/or viability of components of the formulation, a critical aspect of product shelf-stability. A stabilizer can be one of a variety of compounds, such as a dessicant.

As used herein, a "preservative" includes any chemical compound and/or physical conditions that prevent the decomposition of organic constituents of seeds treated with formulations. Chemical preservatives could include, for example, synthetic or non-synthetic antioxidants and physical preservatives could include, for example, refrigeration, freeze-drying or drying.

According to an embodiment the at least one dispersing agent can be in the range of about 2% to about 60% on a dry weight by weight basis. Various dispersing agents are commercially available for use in agricultural compositions, such as those marketed by Rhone Poulenc, Witco, Westvaco, International Speciality products, Croda chemicals, Borregaard, BASF, Rhodia, etc. According to an embodiment the dispersing agents which can be used in the agricultural composition can be chosen from a group comprising polyvinylpyrrolidone, polyvinylalcohol, lignosulphonates, phenyl naphthalene sulphonates, ethoxylated alkyl phenols, ethoxylated fatty acids, alkoxyated linear alcohols, polyaromatic sulfonates, sodium alkyl aryl sulfonates, glyceryl esters, maleic anhydride copolymers, phosphate esters, condensation products of aryl sulphonic acids and formaldehyde, condensation products of alkylaryl sulphonic acids and formaldehyde, addition products of ethylene oxide and fatty acid esters, salts of addition products. of ethylene oxide and fatty acid esters, sulfonates of condensed naphthalene, addition products of ethylene oxide and fatty acid esters, salts of addition products of ethylene

oxide and fatty acid esters, lignin derivatives, naphthalene formaldehyde condensates, sodium salt of isodecylsulfosuccinic acid half ester, polycarboxylates, sodium alkylbenzenesulfonates, sodium salts of sulfonated naphthalene, ammonium salts of sulfonated naphthalene, salts of polyacrylic acids, salts of phenolsulfonic acids and salts of naphthalene sulfonic acids. However, those skilled in the art will appreciate that it is possible to utilize other dispersing agents known in the art without departing from the scope of the claims of the present invention.

In some embodiments, a bacterial endophyte is a seed-origin bacterial endophyte. As used herein, a "seed-origin bacterial endophyte" refers to a population of bacteria associated with or derived from the seed of a grass plant. For example, a seed-origin bacterial endophyte can be found in mature, dry, undamaged (e.g., no cracks, visible fungal infection, or prematurely germinated) seeds. The bacteria can be associated with or derived from the surface of the seed; alternatively, or in addition, it can be associated with or derived from the interior seed compartment (e.g., of a surface-sterilized seed). In some cases, a seed-origin bacterial endophyte is capable of replicating within the plant tissue, for example, the interior of the seed. Also, in some cases, the seed-origin bacterial endophyte is capable of surviving desiccation.

Seed-origin means that the bacterial entity is obtained directly or indirectly from the seed surface or seed interior compartment or is obtainable from a seed surface or seed interior compartment. For example, a seed-origin bacterial entity can be obtained directly or indirectly from a seed surface or seed interior compartment when it is isolated, or isolated and purified, from a seed preparation; in some cases, the seed-origin bacterial entity which has been isolated, or isolated and purified, may be cultured under appropriate conditions to produce a purified bacterial population consisting essentially of a seed-origin bacterial endophyte. A seed-origin bacterial endophyte can be considered to be obtainable from a seed surface or seed interior compartment if the bacteria can be detected on or in, or isolated from, a seed surface or seed interior compartment of a plant.

In some embodiments, the present invention contemplates methods of manually or mechanically combining an endophyte described herein with one or more plant elements, such as a seed, a leaf, or a root, in order to confer an improved agronomic trait or improved agronomic trait potential to said plant element or host plant. In some embodiments, the present invention contemplates methods of manually or mechanically combining a plurality of endophytes described herein with one or more plant elements.

As used herein, an endophyte is “heterologously disposed” when mechanically or manually applied, artificially inoculated or disposed onto or into a plant element, seedling, plant or onto or into a plant growth medium or onto or into a treatment formulation so that the endophyte exists on or in said plant element, seedling, plant, plant growth medium, or treatment formulation in a manner not found in nature prior to the application of the endophyte, e.g., said combination which is not found in nature. In some embodiments, such a manner is contemplated to include: the presence of the endophyte; presence of the endophyte in a different number, concentration, or amount; the presence of the endophyte in or on a different plant element, tissue, cell type, or other physical location in or on the plant; the presence of the endophyte at different time period, e.g. developmental phase of the plant or plant element, time of day, time of season, and combinations thereof. In some embodiments, plant growth medium is soil, a hydroponic apparatus, or artificial growth medium such as commercial potting mix. In some embodiments, the plant growth medium is soil in an agricultural field. In some embodiments, the plant growth medium is commercial potting mix. In some embodiments, the plant growth medium is an artificial growth medium such as germination paper. As a non-limiting example, if the plant element or seedling or plant has an endophyte normally found in the root tissue but not in the leaf tissue, and the endophyte is applied to the leaf, the endophyte would be considered to be heterologously disposed. As a non-limiting example, if the endophyte is naturally found in the mesophyll layer of leaf tissue but is applied to the epithelial layer, the endophyte would be considered to be heterologously disposed. As a non-limiting example, an endophyte is heterologously disposed at a concentration that is at least 1.5 times, between 1.5 and 2 times, 2 times, between 2 and 3 times, 3 times, between 3 and 5 times, 5 times, between 5 and 7 times, 7 times, between 7 and 10 times, 10 times greater, or even greater than 10 times higher number, amount, or concentration than that which is naturally present. As a non-limiting example, an endophyte is heterologously disposed on a seedling if that endophyte is normally found at the flowering stage of a plant and not at a seedling stage.

The compositions provided herein are preferably stable. The seed-origin bacterial endophyte is optionally shelf stable, where at least 10% of the CFUs are viable after storage in desiccated form (i.e., moisture content of 30% or less) for 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or greater than 10 weeks at 4° C. or at room temperature. Optionally, a shelf stable formulation is in a dry formulation, a powder formulation, or a lyophilized formulation. In some embodiments, the

formulation is formulated to provide stability for the population of bacterial endophytes. In one embodiment, the formulation is substantially stable at temperatures between about 0° C and about 50°C for at least about 1, 2, 3, 4, 5, or 6 days, or 1, 2, 3 or 4 weeks, or 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11 or 12 months, or one or more years. In another embodiment, the formulation is

5 substantially stable at temperatures between about 4°C and about 37°C for at least about 5, 10, 15, 20, 25, 30 or greater than 30 days.

An agricultural plant can be a monocotyledonous (i.e., an "agricultural grass plant") or a dicotyledonous plant typically used in agriculture. An agricultural grass plant includes, but is not limited to, maize (*Zea mays*), common wheat (*Triticum aestivum*), spelt (*Triticum spelta*),

10 einkorn wheat (*Triticum monococcum*), emmer wheat (*Triticum dicoccum*), durum wheat (*Triticum durum*), Asian rice (*Oryza sativa*), African rice (*Oryza glaberrima*), wild rice (*Zizania aquatica*, *Zizania latifolia*, *Zizania palustris*, *Zizania texana*), barley (*Hordeum vulgare*), Sorghum (*Sorghum bicolor*), Finger millet (*Eleusine coracana*), Proso millet (*Panicum miliaceum*), Pearl millet (*Pennisetum glaucum*), Foxtail millet (*Setaria italica*), Oat (*Avena*

15 *sativa*), Triticale (*Triticosecale*), rye (*Secale cereal*), Russian wild rye (*Psathyrostachys juncea*), bamboo (*Bambuseae*), grasses, including *Agrostis* spp., *Poa* spp., *Festuca* spp., *Lolium* spp., *Cynodon* spp., *Zoysia* spp., *Koeleria* spp., *Danthonia* spp., or sugarcane (e.g., *Saccharum arundinaceum*, *Saccharum barberi*, *Saccharum bengalense*, *Saccharum edule*, *Saccharum munja*, *Saccharum officinarum*, *Saccharum procerum*, *Saccharum ravennae*, *Saccharum*

20 *robustum*, *Saccharum sinense*, or *Saccharum spontaneum*).

A "host plant" includes any plant, particularly an agricultural plant, which an endophytic microbe such as a seed-origin bacterial endophyte can colonize. As used herein, a microbe is said to "colonize" a plant or seed when it can be stably detected within the plant or seed over a period

25 time, such as one or more days, weeks, months or years; in other words, a colonizing microbe is not transiently associated with the plant or seed. A preferred host plant is a cereal plant.

As used herein, a "reference agricultural plant" is an agricultural plant of the same species, strain, or cultivar to which a treatment, formulation, composition or endophyte preparation as described herein is not administered/contacted. Exemplary reference agricultural plants are described herein. A reference agricultural plant, therefore, is identical to the treated

30 plant with the exception of the presence of the endophyte and can serve as a control for detecting the effects of the endophyte that is conferred to the plant.

"Biomass" means the total mass or weight (fresh or dry), at a given time, of a plant tissue, plant tissues, an entire plant, or population of plants. Biomass is usually given as weight per unit area. The term may also refer to all the plants or species in the community (community biomass).

5 A "bacterial network" means a plurality of endophyte entities (e.g., bacteria, fungi, or combinations thereof) co-localized in an environment, such as on or within a grass agricultural plant. Preferably, a bacterial network includes two or more types of endophyte entities that synergistically interact, such synergistic endophytic populations capable of providing a benefit to the agricultural seed, seedling, or plant derived thereby.

10 An "increased yield" can refer to any increase in biomass or seed or fruit weight, seed size, seed number per plant, seed number per unit area, bushels per acre, tons per acre, kilo per hectare, or carbohydrate yield. Typically, the particular characteristic is designated when referring to increased yield, e.g., increased grain yield or increased seed size.

15 A "transgenic plant" includes a plant or progeny plant of any subsequent generation derived therefrom, wherein the DNA of the plant or progeny thereof contains an exogenous DNA not naturally present in a non-transgenic plant of the same strain. The transgenic plant may additionally contain sequences that are native to the plant being transformed, but wherein the "exogenous" gene has been altered in order to alter the level or pattern of expression of the gene, for example, by use of one or more heterologous regulatory or other elements.

20 The terms "pathogen" and "pathogenic" in reference to a bacterium includes any such organism that is capable of causing or affecting a disease, disorder or condition of a host containing the organism.

25 As used herein, an "agricultural seed" is a seed used to grow a plant typically used in agriculture (an "agricultural plant"). The seed may be of a monocot or dicot plant, and may be planted for the production of an agricultural product, for example grain, food, fiber, etc. As used herein, an agricultural seed is a seed that is prepared for planting, for example, in farms for growing.

30 In some cases, the present invention contemplates the use of microbes that are "compatible" with agricultural chemicals, for example, a fungicide, an anti-bacterial compound, or any other agent widely used in agricultural which has the effect of killing or otherwise interfering with optimal growth of microbes. As used herein, a microbe is "compatible" with an agricultural chemical when the microbe is modified, such as by genetic modification, e.g.,

contains a transgene that confers resistance to an herbicide, or is adapted to grow in, or otherwise survive, the concentration of the agricultural chemical used in agriculture. For example, a microbe disposed on the surface of a seed is compatible with the fungicide metalaxyl if it is able to survive the concentrations that are applied on the seed surface.

5 In some embodiments, an agriculturally compatible carrier can be used to formulate an agricultural formulation or other composition that includes a purified bacterial preparation. As used herein an "agriculturally compatible carrier" refers to any material, other than water, which can be added to a seed or a seedling without causing or having an adverse effect on the seed (e.g., reducing seed germination) or the plant that grows from the seed, or the like.

10 As used herein, a "portion" of a plant refers to any part of the plant, and can include distinct tissues and/or organs, and is used interchangeably with the term "tissue" throughout.

 A "population" of plants, as used herein, can refer to a plurality of plants that were subjected to the same inoculation methods described herein, or a plurality of plants that are progeny of a plant or group of plants that were subjected to the inoculation methods. In addition,
15 a population of plants can be a group of plants that are grown from coated seeds. The plants within a population will typically be of the same species, and will also typically share a common genetic derivation.

 A "reference environment" refers to the environment, treatment or condition of the plant in which a measurement is made. For example, production of a compound in a plant associated
20 with a purified bacterial population (e.g., a seed-origin bacterial endophyte) can be measured in a reference environment of drought stress, and compared with the levels of the compound in a reference agricultural plant under the same conditions of drought stress. Alternatively, the levels of a compound in plant associated with a purified bacterial population (e.g., a seed-origin bacterial endophyte) and reference agricultural plant can be measured under identical conditions
25 of no stress.

 As used herein, a "colony-forming unit" ("CFU") is used as a measure of viable microorganisms in a sample. A CFU is an individual viable cell capable of forming on a solid medium a visible colony whose individual cells are derived by cell division from one parental cell.

30

II. Methods of the Invention

The invention relates to a process and method for the production and use of endophytes as plant inoculants products that provide unique inoculant feature/benefits for the promotion of plant vigor, health, growth and yield comprising bacteria isolated from *Phragmites*, e.g., Strains Sandy LB4, SY1 and strain West 9. Several of these new strains of bacteria have been deposited with the NRRL Agriculture Research Service Culture Collection (herein after "NRRL") under the terms of the Budapest Treaty, under accession numbers provided herein.

The invention also relates to a process and method for producing economically acceptable quantities of preparations of the aforementioned bacteria. The invention further relates to an endophyte product(s) produced by such processes and methods. The endophyte product(s) may comprise a solid substrate of, for example, certain cereals e.g. rye, which contain sufficient natural emulsifiers in the form of various proteins, lignin, to provide the bacteria with excellent natural dispersing/wetting/sticker agents that allows for rapid site occupation on/in plants. The product formulation allows/enables practical use and application of the product(s) to roots, stems, leaves, flowers, bulbs, etc of plants as a water-based sprayable formulations or as dusts for other uses e.g. seed treatment, or as dusts for insect/mite vectors. The product when applied to seeds, roots, stems, leaves, flowers, wounds or cut surfaces of plants enables the endophyte to act as an inoculant within the tissues of plants. The product provides improved roots, leaf, stem and or vegetative bud (flowers) growth to plants and or enhances/improves the germination and emergence of seeds and also causes mortality in neighboring competitor species. The product provides for reduction of environmental or cultural stress to plants e.g. root loss due to trimming, pruning, cutting or other stresses. The product provides improved crop quality and faster development to marketability of the crop. The product provides for a reduction in the dependency on chemical pesticides for pest control e.g. control of *Botrytis*, *Fusarium*, *Pythium* and the like. The product can be used for the production of a variety of greenhouse, horticultural and agronomic field crops. The composition of the invention can be a plant inoculant composition comprising the bacteria described above in admixture with an agrochemically acceptable diluent or carrier. The invention also relates to a method of enhancing growth, health vigor or yield of a plant which method comprises applying the plant inoculant composition of the invention to a plant or plant locus. The invention further relates to a method

of combating a plant fungus which method comprises applying an antifungal effective amount of the composition.

Lyophilization Procedure

5 Freeze drying bacteria (lyophilization) is a very well-established method for the archiving and long-term storage. Initial reports of freeze drying bacteria can be found in the middle of last century. The approaches used vary widely, but they all following the standard process associated with lyophilization, namely the freezing of the sample, application of a high vacuum, warming of the sample while under vacuum which causes water sublimation, driving off excess water
10 through a drying phase, and finally sealing of the sample to prevent water uptake. This general process is used to preserve bacteria, fungi, yeasts, proteins, nucleic acids, and any other molecules which may be degraded due to the presence of water.

 Thus, in one aspect of the invention, one or more of the endophytic bacteria will be applied to a plant or a plant part (such as seeds) as a lyophilized (freeze-dried) powder. In brief,
15 the liquid culture will be: centrifuged, re-suspended in a lyophilization medium which will optionally include cryoprotectants and biological- and/or chemical-oxygen scavengers, transferred to a shelf lyophilizer, lyophilized, and packaged for transport and storage.

 In an alternative approach, one or more bacteria may be encapsulated in alginate beads enriched with humic acid as described by Young CC et al., *Biotechnol Bioeng.* 2006 Sep
20 5;95(1):76-83. Also see “Alginate beads as a storage, delivery and containment system for genetically modified PCB degrader and PCB biosensor derivatives of *Pseudomonas fluorescens* F113 B” by Power et al., *Journal of Applied Microbiology* 110, 1351–1358, 2011.

 Other approaches include coating seeds with preparations comprising the endophytic bacteria of the invention. For example, Sandy LB4 can be incorporated into a carrier, which include without
25 limitation, alginate (micro-bead formation), chitosan, carboxymethylcellulose-starch, clay, finely-ground peat mixed with calcium carbonate, methacrylic acid, bio-char and biogels. In certain embodiments, carrier and Sandy LB4 may be mixed with additives (including: adhesives, nutrients, surfactants and stabilizers).

 The carrier and Sandy LB4, for example, (with additives) can be applied to seeds of grass
30 crops using a commercial seed dressing machine (e.g., MAYJOY High Speed Seeds Dressing Machine/Corn Seed Dresser).

Sandy LB4 may also be used as an additive to create seed balls. In this approach, clay mixed with freeze-dried preparation of Sandy LB4 and seeds (2-3) are added to the center of a small clay ball. The seed balls are then dried and stored for future use.

5 Sandy LB4 can also be applied to host plants in soil drenching approaches. For example, freeze dried preparation of Sandy LB4 can be mixed with a liquid carrier (comprising water, buffers, plant nutrients, and microbial nutrients,). This liquid preparation of bacterium and carrier (with additives) may be applied to the soil around plant or seed or in the alternative be applied to soil and plants using a commercial sprayer.

10 The following materials and methods are provided to facilitate the practice of the present invention.

Seed collection

Seeds of *P. australis* were collected at four sites in New Jersey, including Sandy Hook peninsula, and towns of South River, West Windsor and Robbinsville.

Seed preparation

15 Seeds were cleaned of associated awns, lemmas and paleas by rubbing seeds between a fine mesh screen and wooden block. Seeds were washed with agitation in sterile water then the water was discarded. Seeds were then washed a second time in 1-mL of sterile water after which the water was spread over the surface of agar plates containing trypticase soy agar (TSA), Luria Bertani Agar (LBA) and Yeast Extract Sucrose Agar (YESA) and plates incubated at room
20 temperature for a 24-48 hr period. Colonies were removed and streaked to obtain pure cultures. A total of 16 isolates were obtained from *P. australis* seeds using this method.

Bacterial identification and characterization

Genomic DNA from bacteria was isolated using QIAGEN miniprep kits. Bacterial identifications were made by use of 16S rDNA sequences after methods employed by Lane
25 (1991) (Table1). For Sandy LB4 we used the *recA* gene to identify the bacterium to species. Sequences were compared to GenBank accessions using BLASTn (found on the web at www.ncbi.nlm.nih.gov). Cultures on LB agar were also observed under ultraviolet light using an ultraviolet mineralogical lamp (UVGL-21, long wave UV-366nm, UVP, San Gabriel, California,

USA). Phosphate solubilization was assessed by culturing bacteria on Pikovskayas agar, while protease testing was done by culturing in skim milk agar.

Endophytism determination

Two of the Phragmites seed bacteria (*Pseudomonas fluorescens* Sandy LB4 and *Pseudomonas* sp. West 9) were selected along with bacteria *Microbacterium oxydans* B2 (from Phragmites shoots) and *Bosea thiooxidans* TBN (from Japanese knotweed). These bacteria were screened to determine their capacity to become endophytic in grass roots. Bermuda grass (*Cynodon dactylon*) was used to evaluate endophytism because seeds could be surface sterilized and seedlings would readily grow in sterile agarose medium. Seeds without husks were surface disinfected by agitation in 4% NaOCl solution for 45 mins. Seeds were washed thoroughly by agitation in several washes of sterile water until chlorine odor could not be detected on seeds. Seeds were then placed onto 0.7% agarose in Petri dishes. To inoculate seeds one drop of a water suspension of bacteria (at approximately 0.5 OD, 600nm) was placed onto each seed. For each treatment each plate contained 10-15 seeds, with 5 replicates per treatment. An axenic control was also done to confirm behavior of seedlings without bacteria. Plates were incubated at room temperature for 7-10 days.

Visualization of bacteria in/on seedling roots

To visualize bacteria in/on seedling roots, plates bearing seedlings were flooded and stained with a solution of 3, 3'-diaminobenzidine (DAB) using SIGMAFAST®3, 3'-diaminobenzidine tablets from Sigma (White et al., 2014) and incubated at laboratory ambient temperature overnight for staining. Plates were then washed of excess DAB and the seedling roots examined through the reverse of the Petri plate using a compound light microscope. Intracellular penetration of root cells was indicated by dark red or brown staining within root hairs and/or root parenchyma cells.

Assessment of capacity for plant growth promotion

To assess plant growth promotion, *Pseudomonas fluorescens* Sandy LB4, *Pseudomonas* sp. West 9, *Microbacterium oxydans* B2 and *Bosea thiooxidans* TBN were inoculated onto previously surface disinfected (40 mins agitation in 4% NaOCl) seeds of *Poa annua* by soaking

seeds in a suspension of bacteria (at approximately 0.5 OD, 600nm). Controls were not inoculated with bacteria. Seeds were planted in sterile soil in magenta boxes with approximately 15 plants/treatment and placed under florescent lights in the laboratory for 37 days. Plants were harvested by removal from soil and roots wiped of excess soil. Shoot and root lengths were then measured and recorded. Growth promotion experiments were also conducted using rice (cultivar Rex). In these experiments rice seeds were surface disinfected for 1 hour in 4% NaOCl solution, then washed in sterile water. To inoculate seeds, surface disinfected seeds were soaked for 1 hour with continuous agitation in a suspension of *Pseudomonas fluorescens* Sandy LB4 at approximately 0.5 OD, 600nm. Controls were surface disinfected seeds soaked in sterile water with agitation for 1 hour. 25 seeds for each treatment were planted in sterile soil in magenta boxes and incubated for 7 days under fluorescent lights at laboratory ambient temperature. Seedlings were removed from soil and root and shoot lengths determined.

Seed germination and seedling root architecture

Experiments were conducted to evaluate *Pseudomonas fluorescens* Sandy LB4 effects of seed germination and early seedling root architecture. In these experiments, seeds of *Poa annua* were surface disinfected for 45 mins as above then plated onto agarose media with without nutrients (Murashige and Skoogs basal medium) then monitored daily for 7 days for the parameters: percent germination, root development into agarose, root branching, and root and shoot lengths.

Disease protection (co-culture experiments)

To assess potential for disease protection bacteria *Pseudomonas fluorescens* Sandy LB4, *Pseudomonas sp.* West 9 and *Microbacterium oxydans* B2 were co-cultured on LBA with soil fungi, *Sclerotinia homeocarpa* or *Fusarium oxysporum* for 7 days. Formation of zones of inhibition were noted.

Damping off disease control experiment

To evaluate whether the *Phragmites* bacteria could protect seedlings from disease an experiment was set up using the damping off pathogen *Fusarium oxysporum*. In this experiment surface disinfected seeds of *Poa annua* were inoculated by soaking in a suspension (at

approximately 0.5 OD, 600nm) of strains Sandy LB4, West 9 or *Microbacterium oxydans* B2, then planted in magenta boxes containing previously sterilized soil that had been mixed with 10-mL of a conidial suspension of *Fusarium oxysporum*. Controls included treatment without bacterial or fungal treatments, and with fungus but without bacteria. Each treatment included 25 replicates. After 8 days and 15 days seedlings evident on the surface of the soil were counted. Also presence of fungal mycelium on the surface of soil was determined. To test for movement of the bacteria into soil a sterile probe was placed into soil between plants in each replicate, then the probe was streaked on a fresh Petri plant containing LB agar and incubated for 24 hours prior to examination.

Effects of *Phragmites* bacteria on growth of competitor plants

Dandelion (*Taraxacum officinalis*) was used to screen *Phragmites* microbes for any antagonism leading to increased mortality in the dandelion seedlings. Here dandelion seeds were rubbed against a nylon screen to remove awns and associated fibers. Seeds were then surface disinfected for 30 mins in 0.4% NaOCl, then rinsed in five changes of sterile water to remove residual chlorine. Seeds were used in a series of experiments.

Competitor inhibition experiment 1 (*Phragmites* mixture 1)

Eleven seeds were placed onto the surface of agarose medium (0.7%) in Petri dishes. One drop of bacterial mixture 1 (including equal parts of *Microbacterium oxydans* B2, *Pseudomonas fluorescens* Sandy LB4, and *Pseudomonas sp.* West 9 at approximately OD 0.5, 600nm) was applied to the surface of each seed (with 9 replica plates). Controls were moistened with sterile water only. Plates were then incubated at approximately 21°C for two weeks in alternating light/dark conditions (10-hours light/14-hours dark). Seedlings were then examined to determine whether they were alive (light green) or dead (brown).

Competitor inhibition experiment 2 (*Phragmites* mixture 1 and 2 in soil)

In a second experiment dandelion seeds were cleaned of external fibers, surface disinfected as above and placed onto sterile soil in magenta boxes (10 seeds/treatment). Treatments included bacterial mixture 1 (as indicated above) and bacterial mixture 2 (including strains: *Pseudomonas sp.* WY6, *Pseudomonas sp.* Sandy LB6 *Pseudomonas sp.* Sandy Y8,

Pseudomonas sp. RoLB13w, *Pantoea* sp. RoY12w). A control was not inoculated with Phragmites bacteria. To inoculate seeds with bacteria 2 mL of a bacterial suspension at approximately 0.5 OD, 600nm was mixed into soil in magenta boxes. Seeds of *Poa annua* were also pre-moistened with bacterial suspension or water (control) then placed into soil and boxes
5 incubated for four weeks under alternating light/dark conditions (10-hours light/14-hours dark) and 21°C. After four weeks seedlings were removed from soil, weighed, and root and shoot lengths determined (Table 4).

Competitor inhibition experiment 3 (examination of individual bacterial strains)

A screen was conducted using individual strains of the bacteria included in mixture 1 (as
10 indicated above). In this preliminary individual screen, 14 disinfected dandelion seeds were placed on agarose media in Petri dishes. Seeds were inoculated as above with each bacterium, and control seeds were not inoculated. Three replica plates were made for each treatment and control. Plates were then incubated at approximately 21°C for two weeks in alternating light/dark conditions (10-hours light/14-hours dark), after which mortality was assessed in each plate.

Competitor inhibition experiment 4 (*P. fluorescens* Sandy LB4)

A larger experiment was conducted using strain Sandy LB4. In this experiment dandelion
15 seeds were cleaned and surface disinfected as above, then placed on 0.7% agarose in 38 Petri dishes (15 seeds/dish). Seeds on nineteen of the plates were inoculated using a suspension of Sandy LB4 as above, and the other nineteen were not inoculated. Plates were incubated for two
20 weeks under the conditions of the previous experiment, after which seedling mortality was assessed.

Competitor inhibition experiment 5 (effects on seedlings of curly dock (*Rumex crispus*))

Seeds of locally collected curly dock were cleaned of bracts and associated wings, then
25 surface disinfected for 40 minutes in 4% NaOCl. Ten seeds were then placed onto each of five Petri plates containing 0.7% agarose. Each seed was inoculated with a drop of bacterial suspension (strain Sandy LB4). Five control replica plates were prepared where seeds were not inoculated. Plates were incubated at approximately 21C for one week in alternating light/dark

conditions (10-hours light/14-hours dark) after which seedlings were stained using DAB and examined microscopically.

Competitor inhibition experiment 6 (curly dock in soil)

A test was conducted in which 2 mL of individual bacteria (Sandy LB4, West 9, M. oxydans B2, or WY6 was mixed into previously sterilized soil in magenta boxes, then ten surface disinfected seeds of curly dock were placed into the soil 2-3 mm beneath surface. A control treatment was not inoculated with bacteria. After two weeks, the number of seedlings emerging from the soil surface was determined.

The following examples are provided to illustrate certain embodiments of the invention. They are not intended to limit the invention in any way.

Example I

Host promotional properties of endophytic *Pseudomonas* spp. from seeds of invasive reed grass (*Phragmites australis*)

This invention involves application of strains (e.g., Sandy LB4) of endophytic bacteria (belonging to genus *Pseudomonas*) for increasing growth of turf and forage grasses (*Agrostis* spp., *Poa* spp., *Festuca* spp., *Lolium* spp., *Cynodon* spp., *Zoysia* spp., *Koeleria* spp., *Danthonia* sp.) and food grass crops (*Triticum aestivum*, *Oryza sativa*, *Zea mays*), and suppress soil borne fungal diseases (e.g., damping off disease), and reduce vigor of seedlings of distantly-related weeds including *Taraxacum officinale* (dandelion) and curly dock (*Rumex crispus*). Each of the endophytes described herein (See Table 1) was obtained from seeds of the invasive grass *Phragmites australis*. For example, Sandy LB4 colonizes root meristems and enters into the cells of roots. This endophyte is unique in entering cells of the plant where it elicits production of reactive oxygen. Elicitation of reactive oxygen likely increases the resistance of the grass host to oxidative stress by stimulating the plant to up-regulate oxidative stress resistance genes. Sandy LB4 increases the growth of grass seedlings resulting in seedlings with larger root systems and larger shoots compared to seedlings without the bacterium. Sandy LB4 also moves from the plant roots out into the soil and colonizes mycelium of soil fungi suppressing growth of fungal pathogens resulting in reduced seedling disease. Sandy LB4 colonizes seedlings of non-grass competitor plants (dandelions and curly dock) and increases mortality of non-grass seedlings in laboratory experiments.

Application of the endophytic microbes obtained from *Phragmites australis* described herein to heterologous grass crops (e.g., turfgrasses, wheat, rice, sorghum, etc.) improves crop growth and reduces soil borne diseases and weeds. The endophytic bacteria listed in Table 1, including Sandy LB4, may be used alone or in combination in soil additives or seed treatments to grow crops with reduced agrochemical inputs. Widespread application of these endophytic bacteria commercially will enable decreased use of fertilizers, fungicides and herbicides on grass crops, resulting in reduced cost for expensive agrochemicals and reduced contamination of soil and water.

Isolates

Sixteen bacterial isolates were obtained from the surfaces of *Phragmites* seeds. Of these, 12 were shown to belong in genus *Pseudomonas* based on 16S rDNA sequence data (Table 1). Two strains examined closely *Pseudomonas* sp. West 9 and *Pseudomonas fluorescens* Sandy LB4 were shown to become endophytic in the Bermuda grass seedlings (Fig. 1 and Fig. 2). These bacteria colonized root tip meristems and became intracellular in root cells, inducing formation of root hairs on seedling roots.

Table 1. Bacteria isolated from *Phragmites australis* seed surfaces

	Collection Site	Strain ID	UV	P ¹	Prot. ²	Genus and Species	NRRL deposit #
20	Sandy Hook	Sandy LB4	BF*	Y	Y	<i>P. fluorescens</i>	NRRL-B67308
	Sandy Hook	Sandy LB6	LF	Y	N	<i>Pseudomonas</i> sp.	NRRL-XXXX
	Sandy Hook	Sandy Y8	LF	Y	N	<i>Pseudomonas</i> sp.	NRRL-XXXX
	Robbinsville	RoY12	NF	Y	N	<i>Pantoea</i> sp.	NRRL-XXXX
	Robbinsville	RoLB13w	LF	Y	N	<i>Pseudomonas</i> sp.	NRRL-XXXX
25	West Windsor	WY9y	NF	Y	N	<i>Enterobacter</i> sp.	NRRL-XXXX
	West Windsor	WY6	LF	Y	N	<i>Pseudomonas</i> sp.	NRRL-XXXX
	West Windsor	West 9	LF	Y	N	<i>Pseudomonas</i> sp.	NRRL-B67307
	West Windsor	WY14	NF	Y	Y	<i>Pseudomonas</i> sp.	NRRL-XXXX
	South River	RiY3	NF	Y	N	<i>Pseudomonas</i> sp.	NRRL-XXXX
30	Sandy Hook	SandyY7w	--	Y	N	<i>Pseudomonas</i> sp.	NRRL-XXXX
	South River	RiLB4	NF	Y	N	<i>Pantoea</i> sp.	NRRL-XXXX
	Sandy Hook	SY1	NF	Y	N	<i>Pseudomonas</i> sp.	NRRL-XXXX
	Sandy Hook	SY5	NF	Y	N	<i>Pseudomonas</i> sp.	NRRL-XXXX
	West Windsor	WY9w	--	-		<i>Pseudomonas</i> sp.	NRRL-XXXX
35	Cook Campus	B2				<i>Microbacterium oxydans</i>	NRRL-B67470

*BF = bright fluorescence; Lf = light fluorescence; NF = no fluorescence observed.

¹Phosphorus solubilization, Y = yes; -- = missing data.

²Protease production based on milk agar clearing; Y = yes; N = no; -- = missing data.

Partial 16S rDNA sequences for these strains from top to bottom are provided in SEQ ID NOS:

5 1, 2, 3, 6, 5, 12, 7, 10, 9, 4, 13, 14, 15, 16, and 8. Sequences are provided in Table 6.

Modes of colonizing Bermuda grass seedling roots

Several strains, including Sandy LB4 and West 9 were shown to enter root cells at the root meristem. Squash preparations of root tips showed presence of bacteria within meristematic
 10 cells. See Fig. 3A- 3B. In older root cells, the spherical bacteria were seen to stain brown (positive indication of reactive oxygen) using the reactive oxygen stain DAB. Spherical bacterial L-forms were evident in root hairs and parenchyma cells (Fig. 4A –Fig. 4E) that comprised the main axis of the root. In older parts of the root the L-forms were seen to swell and the contents lacked protein as evident by failure to stain blue using the aniline blue counterstain (Fig. 5). In
 15 still older root parts, including root hairs and parenchyma, the internal bacterial inclusions were no longer evident. Seedlings derived from seeds that had not been inoculated with either Sandy LB4 or West 9 did not show presence of bacterial L-forms within root cells.

Plant growth promotion capacity

Pseudomonas fluorescens Sandy LB4, *Pseudomonas sp.* West 9 and *Microbacterium*
 20 *oxydans* B2 were found to increase growth of *Poa annua* in the magenta box experiment. Roots and shoots of seedlings bearing these three bacteria were shown to be significantly longer than those without bacteria or inoculated with *Bosea thiooxidans* TBN (from Japanese knotweed; family *Polygonaceae*; Table 2). In the experiment where Sandy LB4 was applied to rice seeds, seedling shoot growth was increased 46% over seedlings without bacteria (Table 3). In general
 25 root and shoot lengths were significantly larger in inoculated seedlings than those of non-inoculated controls. In addition, roots of inoculated seedlings showed more adherent soil particles compared to non-inoculated seedling roots.

Table 2A. Growth promotion of *Poa annua* in soil for 37 days (15 plants in each treatment)

Treatment	Shoot length (mm)*	Root length (mm)
No bacteria	88.87 ± 23.53 ^a	17.27 ± 8.59 ^a
TBN	86.2 ± 32.35 ^a	13.73 ± 5.01 ^a
Sandy LB4	122.33 ± 27.38 ^b	23.8 ± 8.54 ^b
West 9	119.33 ± 20.43 ^b	27.33 ± 14.56 ^b
<i>Microbacterium oxydans</i> B2	113.73 ± 25.41 ^b	20.87 ± 6.79 ^{a, b}

* Data is given as mean ± standard deviation. Means followed by the same letter within a column are not significantly different according to the Duncan multiple range test ($p < 0.05$).

Table 2B. Growth promotion of Bermuda grass (*Cynodon* spp.)

Endophytes (Bacteria)	Geotropic response (%)	Root length (cm)	Root hairs and branching*	Shoot length (cm)	IAA test
Control	35	1.5-2.0	+	1.5-2.0	-
1. West9	35	1.5-2.0	+++	1.5-2.0	NA
2. Sandy LB4	50	2.0-3.0	+++	1.5-2.5	-
3. <i>Microbacterium oxydans</i> B2	40	1.0-1.5	+	1.5-2.0	NA
4. SY1	40	2.0-3.0	+++	1.5-2.5	-
5. Sandy LB6	60	2.0-3.0	+++	1.5-2.5	-

Table 3. Rice seedling development with and without the indicated endophytic bacteria isolate

Endophytes (Bacteria)	Geotropic response (%)	Root length (cm)	Root hairs and branching*	Shoot length (cm)	Phosphate solubilization
Control	10	1.0-2.0	+	1.0-2.0	-
1. West9	20	2.0-3.0	++	2.0-2.5	+
2. Sandy LB4	40	2.0-3.0	+++	2.0-3.0	+

3. <i>Microbacterium oxydans</i> B2	20	2.0-3.0	++	2.0-2.5	+
4. SY1	25	2.0-2.5	+++	2.0-2.5	+
5. SY5	10	1.0-2.0	No	1.0-2.0	NA
6. Sandy LB6	25	2.0-3.0	++	2.0-3.0	+
7. RiY3	0.0	1.0-1.5	No	0.5-1.5	NA
8. ROLB13y	10	1.0-1.5	No	0.5-1.5	NA

Seed germination rate and root architecture

In the germination and root growth study where we used axenic *Poa annua* to determine bacterial effects on seeds and seedlings, we found that the germination rate was only slightly increased in seeds bearing Sandy LB4. However, there was a slightly greater tendency for roots bearing the bacterium to branch compared to roots without the bacterium (Fig. 6). In the agarose medium roots bearing the bacterium were also seen to more frequently grow downward into the agarose medium than roots without the bacterium. It is notable that in seedlings grown on agarose containing Murashige and Skoogs nutrient base, the majority of the roots tended to grow downward into the medium (Table 4).

Table 4. *Poa annua* germination and growth with *Phragmites* bacterium Sandy LB4 over a 1-week period.

Agarose	Total GERM	% GERM	Roots Down	Roots Up	2 nd root	3 rd root	Shoot Avg.	Root Avg.
No bacteria	117	90	30	87	100	58	18	16
Sandy LB4	123	95	54	68	103	71	21	12
MS	Total GERM	% GERM	Roots Down	Roots Up	2 nd root	3 rd root	Shoot Avg.	Root Avg.
No bacteria	118	91	116	0	67	1	26	12
Sandy LB4	122	94	116	4	78	6	24	13

Production of fungal inhibitors

Inhibition zones were evident in co-culture experiments, where West 9 and Sandy LB4 were cultured with *Sclerotinia homeocarpa* and *Fusarium oxysporum* and seen to inhibit them in culture. *Microbacterium oxydans* B2 did not produce inhibitory substances in culture. Another
5 endophytic bacteria, SY1, also produced fungal inhibitors as described further below.

Damping off disease protection

In the experiment evaluating capacity of *Phragmites* bacteria to protect *P. annua* seedlings from damping off caused by *Fusarium oxysporum*, we showed that the least damping
10 off was seen in the pathogen and bacterial free treatment with 23 seedlings surviving after 15 days. After 15 days Sandy LB4 showed the highest level of protection with 20 seedlings surviving; followed by West 9 with 16 seedlings surviving, *Microbacterium oxydans* B2 with 15 seedlings surviving, and the pathogen + no bacterium treatment with 12 seedlings surviving. SYI also effectively inhibited fungal growth as described below.

Effects of *Phragmites* bacteria on growth of seedlings of competitor species

In competitor inhibition experiment 1 we showed that a mixture of *Phragmites* bacteria (including *Microbacterium oxydans* B2, *Pseudomonas sp.* West 9 and *Pseudomonas fluorescens* Sandy LB4) increased mortality in dandelion seedlings compared to the non-inoculated control
20 with the difference being statistically significant at the 5% level of significance in the Student's t-test. The bacterial mixture gave an $83.4 \pm 15.7\%$ mortality compared to the axenic control at $16.4 \pm 22.8\%$ mortality.

In competitor inhibition experiment 2 conducted in soil in magenta boxes we found that inoculation of soil with bacterial mixture 1 (including Sandy LB1, West 9 and *Microbacterium*
25 *oxydans* B2) resulted seedlings that were significantly smaller in terms of weight, and shoot and root lengths than those of non-inoculated seedlings or those grown in soil containing *Phragmites* bacterial mixture 2 containing five other isolates (Table 5).

Table 5. Dandelion seedling growth inhibition in soil by *Phragmites* bacterial mixtures

Treatment	Seedling mass Wet weight (mg)	Shoot length (mm)	Root length (mm)
No bacteria	27.03 ± 9.54 ^{*a}	42.44 ± 12.6 ^a	40 ± 17.11 ^a
Mix 1	14.91 ± 8.71 ^b	23.88 ± 11.86 ^b	14.63 ± 5.21 ^b
Mix 2	26.36 ± 6.78 ^a	40.89 ± 10.71 ^a	43.33 ± 19.55 ^a

*Data given as means ± standard deviation. Within columns means followed by the same letter are not statistically different according to the Duncan multiple range test ($p < 0.05$).

Identification of strains showing dandelion seedling inhibition

In competitor inhibition experiment 3, *Pseudomonas fluorescens* Sandy LB4, *Pseudomonas sp.* West 9 and *Microbacterium oxydans* B2 were tested for their effect on mortality of dandelion seedlings on agarose (Fig. 9). In this test the most potent bacterium for increasing mortality of dandelion seedlings was Sandy LB4 (71%) > *Microbacterium oxydans* B2 (35%) > West 9 (31%) > no bacterium control (23%). On the basis of this test we concluded that Sandy LB4 was the most potent of the bacteria increasing mortality in dandelion seedlings and a larger study (Competitor inhibition experiment 4) was set up to gather statistical data for effect of Sandy LB4 on dandelion seedling mortality. In the larger study a statistically significant ($p < 0.05$ in student's t-test) increased mortality of $71.1 \pm 15.3\%$ was shown for Sandy LB4-inoculated seedlings compared to only $6.8 \pm 7.4\%$ for the non-inoculated control treatment.

Effects of *Pseudomonas fluorescens* Sandy LB4 on curly dock (*Rumex crispus*) seedlings

In competitor inhibition experiment 5 we evaluated the colonization of seedlings of *Rumex crispus* by Sandy LB4. Here we found that Sandy LB4 actively colonized seedlings, inducing root hair development in the inoculated seedlings. Staining the seedlings using the reactive oxygen stain DAB and microscopic examination of roots showed that root hairs appeared to be occluded by large dark clusters of swollen degrading L-form bacteria (Fig. 8). In competitor inhibition test 6 in soil we showed that curly dock seedling emergence was: no bacterium treatment (7 seedlings) > West 9 (3 seedlings) > *Microbacterium oxydans* B2 (2 seedlings) > Sandy LB4 (0 seedlings).

Fungal disease protection in rice (*Oryza sativa*) seedlings by growth promoting seed associated endophytic bacteria from invasive *Phragmites australis*

As discussed at length above, non-cultivated plants carry endophytes that may be used to enhance development and disease resistance of desirable crops. During seedling establishment, rice may be infected by fungal pathogens. Species of *Fusarium* and *Pythium* cause rice seed rot during germination. *Fusarium* blight in rice seedlings is common and affects seedling development.

We found that three bacteria, including *Pseudomonas fluorescens* (Sandy LB4), *Pseudomonas sp.* (Sandy LB6) and *Pseudomonas sp.* (SY1) promoted seedling development, including enhancement of root and shoot growth, and modulation of root hair formation in rice and Bermuda grass. The results with Sandy LB4 and rice are provided above in Table 3. Bacteria were positive for phosphate solubilization. Another bacteria, *Pseudomonas sp.* (SY1) protected rice and Bermuda grass seedlings from *Fusarium* infection. In co-culture experiments, strain SY1 inhibited fungal pathogens with 85.71 % growth inhibition of *F. oxysporum*, 86.33 % growth inhibition of *Curvularia sp.* and 82.14 % growth inhibition of *Alternaria sp.* Seedlings of Bermuda grass and rice treated with bacteria were found much less infected by *F. oxysporum* in comparison to nontreated controls. Bacteria appeared to actively degrade fungal mycelia. Metabolite products of strain SY1 were also found to inhibit fungal growth on nutrient media. *Pseudomonas sp.* (SY1) was found to produce inhibitory volatiles. MALDI-TOF analysis revealed that SY1 produces antifungal lipopeptides of the fengycin family which appear to be responsible for the antifungal activity of SY1.

Table 6. Exemplary partial 16S rDNA and SEQ ID NOS: for strains described herein

Sandy LB4 (*Pseudomonas fluorescens* deposited as NRRL-B67308)

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TAGATACCCTGGTAGTCCACGCCGTAACGATGTCAACTAGCCGTTGGGAGCCTTGAG
CTCTTAGTGGCGCAGCTAACGCATTAAGTTGACCGCCTGGGGAGTACGGCCGCAAG
GTTAAACTCAAATGAATTGACGGGGGCCCGCACAAAGCGGTGGAGCATGTGGTTTA
ATTCGAAGCAACGCGAAGAACCTTACCAGGCCTTGACATCCAATGAACCTTCTAGAG
ATAGATTGGTGCCTTCGGGAACATTGAGACAGGTGCTGCATGGCTGTCGTCAGCTCG
TGTCGTGAGATGTTGGGTTAAGTCCCGTAACGAGCGCAACCCTTGTCCTTAGTTACC
AGCACGTTATGGTGGGCACTCTAAGGAGACTGCCGGTGACAAACCGGAGGAAGGTG
GGGATGACGTCAAGTCATCATGGCCCTTACGGCCTGGGCTACACACGTGCTACAATG
GTCGGTACAAAGGGTTGCCAAGCCGCGAGGTGGAGCTAATCCCATAAAACCGATCG
TAGTCCGGATCGCAGTCTGCAACTCGACTGCGTGAAGTCGGAATCGCTAGTAATCGC

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GAATCAGAATGTCGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACAC CATGGGAGTGGGTTGCACCAGAAGTAGCTAGTTTAACCTTCGGG (SEQ ID NO: 1)
Sandy LB6 (<i>Pseudomonas sp.</i> deposited as NRRL-_____) GTCCTTAGTTACCAGCACGTTATGGTGGGCACTCTAAGGAGACTGCCGGTGACAAAC CGGAGGAAGGTGGGGATGACGTCAAGTCATCATGGCCCTTACGGCCTGGGCTACAC ACGTGCTACAATGGTCGGTACAGAGGGTTGCCAAGCCGCGAGGTGGAGCTAATCTC ACAAAACCGATCGTAGTCCGGATCGCAGTCTGCAACTCGACTGCGTGAAGTCGGAA TCGCTAGTAATCGCGAATCAGAATGTCGCGGTGAATACGTTCCCGGGCCTTGTACAC ACCGCCCGTCACACCATGGGAGTGGGTTGCACCAGAAGTA (SEQ ID NO: 2)
Sandy Y8 (<i>Pseudomonas sp.</i> deposited as NRRL-_____) CTACTTCTGGTGCAACCCACTCCCATGGTGTGACGGGCGGTGTGTACAAGGCCCGGG AACGTATTCACCGCGACATTCTGATTCGCGATTACTAGCGATTCCGACTTCACGCAG TCGAGTTGCAGACTGCGATCCGGACTACGATCGGTTTTGTGAGATTAGCTCCACCTC GCGGCTTGGAACCCCTCTGTACCGACCATTGTAGCACGTGTGTAGCCCAGGCCGTAA GGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCGGTTTGTACCGGCAGTC (SEQ ID NO: 3)
RiY3 (<i>Pseudomonas sp.</i> deposited as NRRL-_____) TTTGACGTTACCGACAGAATAAGCACCGGCTAACTCTGTGCCAGCAGCCGCGGTAAAT ACAGAGGGTGCAAGCGTTAATCGGAATTACTGGGCGTAAAGCGCGCGTAGGTGGTT TGTTAAGTTGGATGTGAAAGCCCCGGGCTCAACCTGGGAACTGCATCCAAAACCTGG CAAGCTAGAGTACGGTAGAGGGTGGTGGAAATTCCTGTGTAGCGGTGAAATGCGTA GATATAGGAAGGAACACCAGTGCGGAAGGCGACCACCTGGACTGATACTGACACTG AGGTGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCACGCCGTA ACGATGTCAACTAGCCGTTGGAATCCTTGAGATTTTAGTGGCGCAGCTAACGCATTA AGTTGACCGCCTGGGGAGTACGGCCGCAAGGTTAAACTCAAATGAATTGACGGGG GCCCCGACAAGCGGTGGAGCATGTGGTTTAATTCGAAGCAACGCGAAGAACCTTAC CGGCCTTGACATGCAGAGAACTTTCAGAGATGGATGG (SEQ ID NO: 4)
RoLB13w (<i>Pseudomonas sp.</i> deposited as NRRL-_____) TGGGAACTGCATCCAAAACCTGGCAAGCTAGAGTACGGTAGAGGGTGGTGGAAATTC CTGTGTAGCGGTGAAATGCGTAGATATAGGAAGGAACACCAGTGCGGAAGGCGACC ACCTGGACTGATACTGACACTGAGGTGCGAAAGCGTGGGGAGCAAACAGGATTAGA TACCCTGGTAGTCCACGCCGTAAACGATGTCAACTAGCCGTTGGAATCCTTGAGATT TTAGTGGCGCAGCTAACGCATTAAGTTGACCGCCTGGGGAGTACGGCCGCAAGGTT AAAACCTCAAATGAATTGACGGGGGCCCCGCACAAGCGGTGGAGCATGTGGTTTAATT CGAAGCAACGCGAAGAACCTTACCAGGCCTTGACATGCAGAGAACTTTCAGAGAT GGATTGGTGCCTTCGGGAACTCTGACACAGGTGCTGCATGGCTGTCGTCAGCTCGTG TCGTGAGATGTTGGGTAAAGTCCCGTAACGAGCGCAACCCTTGTCCTTAGTTACCAG CACGTTATGGTGGGCACTCTAAGGAGACTGCCGGTGACAAACCGGAGGAAGGTGGG GATGACGTCAAGTCATCATGGCCCTTACGGCCTGGGCTACACACGTGCTACAATGGT CGGTACAGAGGGTTGCCAAGCCGCGAGGTGGAGCTAATCTCAGAAAACCGATCGTA GTCCGGATCGCAGTCTGCAACTCGACTGCGTGAAGTCGGAATCGCTAGTAATCGCGA ATCAGAATGTCGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACACCA TGGGAGTGGGTTGCACCAGAAG (SEQ ID NO: 5)
RoY12 (<i>Pantoea sp.</i> deposited as NRRL-_____)

AAGCTCGTGTGTGAAATGTTGGGTAAAGTCCCGCAACGAGCGCAACCCTTATCCTT
 TGTGTCAGCGATTCCGGTCGGGAACCTCAAAGGAGACTGCCGGTGATAAACCGGAGG
 AAGGTGGGGATGACGTCAAGTCATCATGGCCCTTACGAGTAGGGCTACACACGTGC
 TACAATGGCGCATACAAAGAGAAGCGACCTCGCGAGAGCAAGCGGACCTCACAAA
 GTGCGTCGTAGTCCGGATCGGAGTCTGCAACTCGACTCCGTGAAGTCGGAATCGCTA
 GTAATCGTGGATCAGAATGCCACGGTGAATACGTTCCCGGGCCTTGTAACACACCGCC
 CGTCACACCATGGGAGTGGGTTGCAAAAGAAGTA (SEQ ID NO: 6)

WY6 (*Pseudomonas* sp. deposited as NRRL-_____)

GAAGGTTAGACTAGCTACTTCTGGTGCAACCCACTCCCATGGTGTGACGGGCGGTGT
 GTACAAGGCCCGGGAACGTATTCACCGCGACATTCTGATTCGCGATTACTAGCGATT
 CCGACTTCACGCAGTCGAGTTGCAGACTGCGATCCGGACTACGATCGGTTTTGTGGG
 ATTAGCTCCACCTCGCGGCTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTGT
 AGCCCAGGCCGTAAAGGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCGGTTTG
 TCACCGGCAGTCTCCTTAGAGTGCCCAACATCACGTGCTGGTAACTAAGGACAAGG
 GTTGCGCTCGTTACGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCC
 ATGCAGCACCTGTCTCAATGTTCCCGAAGGCACCAATCCATCTCTGGAAAGTTCATT
 GGATGTCAAGGCCTGGTAAGGTTCTTCGCGTTGCTTCGAATTAAACCACATGCTCCA
 CCGCTTGTGCGGGCCCCCGTCAATTCATTTGAGTTTTAACCTTGCGGCCGTACTCCCC
 AGGCGGTCAACTTAATGCGTTAGCTGCGCCACTAAGAGCTCAAGGCTCCCAACGGC
 TAGTTGACATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCAC
 GCTTTCGCACCTCA (SEQ ID NO: 7)

WY9w (*Pseudomonas* sp. deposited as NRRL-_____)

CTTCTGGTGCAACCCACTCCCATGGTGTGACGGGCGGTGTGTACAAGGCCCGGGAA
 CGTATTCACCGCGACATTCTGATTCGCGATTACTAGCGATTCCGACTTCACGCAGTC
 GAGTTGCAGACTGCGATCCGGACTACGATCGGTTTTGTGGGATTAGCTCCACCTCGC
 GGCTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTGTAGCCCAGGCCGTAAAG
 GCCATGATGACTTGACGTCATCCCCACCTTCCTCCGGTTTGTCACCGGCAGTCTCCTT
 AG (SEQ ID NO: 8)

WY14 (*Pseudomonas* sp. deposited as NRRL-_____)

TACCGTCCCCCGAAGGTTAGACTAGCTACTTCTGGTGCAACCCACTCCCATGGTGT
 GACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCGCGACATTCTGATTCGCG
 ATTACTAGCGATTCCGACTTCACGCAGTCGAGTTGCAGACTGCGATCCGGACTACGA
 TCGGTTTTATGGGATTAGCTCCACCTCGCGGCTTGGCAACCCTCTGTACCGACCATTG
 TAGCACGTGTGTAGCCCAGGCCGTAAAGGGCCATGATGACTTGACGTCATCCCCACCT
 TCCTCCGGTTTGTACCGGCAGTCTCCTTAGAGTGCCCAACATAACGTGCTGGTAAC
 TAAGGACAAGGGTTGCGCTCGTTACGGGACTTAACCCAACATCTCACGACACGAGC
 TGACGACAGCCATGCAGCACCTGTCTCAGTGTTCGCGAAGGCACCAAACCATCTCTG
 GTAAGTTCACTGGATGTCAAGGCCTGGTAAGGTTCTTCGCGTTGCTTCGAATTAAAC
 CACATGCTCCACCGCTTGTGCGGGCCCCCGTCAATTCATTTGAGTTTTAACCTTGCGG
 CCGTACTCCCCAGGCGGTCAACTTAATGCGTTAGCTGCGCCACTAAAATCTCAAGGA
 TTCCAACGGCTAGTTGACATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCTGT
 TTGCTCCCCACGCTTTCGCACCTCAGTGTGAGTATGAGCCCAGGTGGTCGCCTTCGC
 CACTGGTGTTCCTTCTATATCTACGCATTTACCGCTACACAGGAAATTCCACCACC
 CTCTGCCCTACTCTA (SEQ ID NO: 9)

West 9 (*Pseudomonas* sp. deposited as NRRL-B67307)

GTACCGTCCTCCCGAAGGTTAGACTAGCTACTTCTGGTGCAACCCACTCCCATGGTG
 TGACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCGCGACATTCTGATTCGCG
 ATTACTAGCGATTCCGACTTCACGCAGTCGAGTTGCAGACTGCGATCCGGACTACGA
 TCGGTTTTGTGGGATTAGCTCCACCTCGCGGCTTGGCAACCCTCTGTACCGACCATTG
 TAGCACGTGTGTAGCCCAGGCCGTAAGGGCCATGATGACTTGACGTCATCCCCACCT
 TCCTCCGGTTTGTACCGGCAGTCTCCTTAGAGTGCCCAACATAACGTGCTGGTAAC
 TAAGGACAAGGGTTGCGCTCGTTACGGGACTTAACCCAACATCTCACGACACGAGC
 TGACGACAGCCATGCAGCACCTGTCTCAATGCTCCCGAAGGCACCGATCCATCTCTG
 GAAAGTTCATTGGATGTCAAGGCCTGGTAAGGTTCTTCGCGTTGCTTCGAATTAAAC
 CACATGCTCCACCGCTTGTGCGGGCCCCCGTCAATTCAATTTGAGTTTTAACCTTGCGG
 CCGTACTCCCCAGGCGGTCAACTTAATGCGTTAGCTGCGCCACTAAAATCTCAAGGA
 TTCCAACGGCTAGTTGACATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCTGT
 TTGCTCCCCACGCTTTCGCACCTCAGTGTCAGTATGAGCCCAGGTGGTCGCCTTCGC
 CACTGGTGTTCTTCTATATCTACGCATTTACCGCTACACAGGAAATTCCACCACC
 CTCTGCCCTACTCTAGCTTGCCAGTTTTTGGATGCAGTTCCAGGTTGAGCCCGGGGA
 TTTACATCCAACCTTAACAAACCACCTACGCGCGCTTTACGCCAGTAATTCCGATT
 AACGCTTGACCCCTCTGTATTACCGCGGCTGCTGGCACAGAGTTAGCCGGTGCTTAT
 TCTGTGCGTAACGTCAAAACACTAACGTATTAGGTTAATGCCCTTCTCCCAACTTA
 AAGTGCTTTACAATCCGAAGACCTTCTTACACACGCGGCATGGCTGGATCAGGCTT
 TCGCCCATTTGTCCAATATTCCTCCACTGCTGCCTCCCGTAGGAGTCTGGACCGTGTCTC
 AGTTCCAGTGTGACTGATCATCCTCTCAGACCAGTTACGGATCGTCGCCTTGGTGAG
 CCATTACCTACCAACTAGCTAATCCGACCTAGGCTCATCTGATAGCGTGAGGTCCG
 AAGATCCCCCACTTTCTCCCGTAGGACGTATGCGGTATTAGCGTCCCTTTTCGAGACG
 TTGTCCCCCACTACCAGGCAGATTCTAGGCATTACTACCCGTCGCGCGCTGAATC
 AGGGAGCAAGCTCCCTCATCCGCTCGACTGCA (SEQ ID NO: 10)

Microbacterium B2 (*Microbacterium oxydans* deposited as NRRL-B67470)

AGTCGAACGAGTGAACACGGAGCTTGCTCTGTGAGGATCAGTGGCGAACGGGTGAG
 TAACACGTGAGCAACCTGCCCCTGACTCTGGGATAAGCGCTGGAAACGGCGTCTAA
 TACTGGATATGTGACGTGACCGCATGGTCTGCGTCTGGAAAGAATTTTCGGTTGGGGA
 TGGGCTCGCGGCTATCAGCTTGTGTTGGTGAGGTAATGGCTCACCAAGGCGTCGACGG
 GTAGCCGGCCTGAGAGGGTGACCGGCCACACTGGGACTGAGACACGGGCCAGACTC
 CTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGCAAGCCTGATGCAGCAA
 CGCCGCGTGAGGGACGACGGCCTTCGGGTTGTAAACCTCTTTTAGCAGGGAAGAAG
 CGAAAGTGACGGTACCTGCAGAAAAAGCGCCGGCTAACTACGTGCCAGCAGCCGCG
 GTAATACGTAGGGCGCAAGCGTTATCCGGAATTATTGGGCGTAAAGAGCTCGTAGG
 CGGTTTGTGCGTCTGCTGTGAAATCCGGAGGCTCAACCTCCGGCCTGCAGTGGGTA
 CGGGCAGACTAGAGTGCGGTAGGGGAGATTGGAATTCCTGGTGTAGCGGTGGAATG
 CGCAGATATCA
 GGAGGAACACCGATGGCGAAGGCAGATCTCTGGGCGGTAACCTGACGCTGAGGAGCG
 AAAGGTGGGGAGCAACAGGCTTAGATACCCTGGTAGTCCACCCCGTAACGTTGGGA
 ACTAGTTGTGGGGTCCATTCCACGGATTCCGTGACGCAGCTAACGCATTAAAGTTCCC
 CGCCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGACCCGCA
 CAAGCGGCGGAGCATGCGGATTAATTCGATGCAACGCGAAGAACCTTACCAAGGCT
 TGACATATACGAGAACGGGCCAGAAATGGTCAACTCTTTGGACACTCGTAAACAGG

TGGTGCATGGTTGTCGTCAGCTCGTGTGTCGTGAGATGTTGGGTAAAGTCCCGCAACGA
GCGCAACCCTCGTTCTATGTTGCCAGCACGTAATGGTGGGAACCTCATGGGATACTGC
CGGGGTCAACTCGGAGGAAGGTGGGGATGACGTCAAATCATCATGCCCTTATGTCT
TGGGCTTCACGCATGCTACAATGGCCGGTACAAAGGGCTGCAATACCGCGAGGTGG
AGCGAATCCCAAAAAGCCGGTCCCAGTTCGGATTGAGGTCTGCAACTCGACCTCAT
GAAGTCGGAG
TCGCTAGTAATCGCAGATCAGCAACGCTGCGGTGAATACGTTCCCGGGTCTTGTACA
CACCGCCCGTCAAGTCATGAAAGTCGGTAACACCTGAAGCCGGTGGCCTAACCTTTT
TGGAGGGAGCCGTCGAAGGTGGG (SEQ ID NO: 11)

WY9y (*Enterobacter* sp. deposited as NRRL-_____)

GCCAGCGGTAGGCCGGGAACCTCAAAGGAGACTGCCAGTGATAAACTGGAGGAAG
GTGGGGATGACGTCAAGTCATCATGGCCCTTACGACCAGGGCTACACACGTGCTAC
AATGGCGCATACAAAGAGAAGCGATCTCGCGAGAGCCAGCGGACCTCATAAAGTGC
GTCGTAGTCCGGATTGGAGTCTGCAACTCGACTCCATGAAGTCGGAATCGCTAGTAA
TCGTGAATCAGAATGTCACGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTC
ACACCATGGGAGTGGGTTGCAAAAGAAGT (SEQ ID NO: 12)

Sandy Y7w (*Pseudomonas* sp. deposited as NRRL-_____)

TGCCAGCGGGTAATGCCGGGAACCTCAAAGGAGACTGCCGGTGATAAACCGGAGGAA
GGTGGGGATGACGTCAAGTCATCATGGCCCTTACGGCCAGGGCTACACACGTGCTA
CAATGGCGCATACAAAGAGAAGCGACCTCGCGAGAGCAAGCGGACCTCATAAAGT
GCGTCGTAGTCCGGATCGGAGTCTGCAACTCGACTCCGTGAAGTCGGAATCGCTAGT
AATCGTAGATCAGAATGCTACGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCG
TCACACCATGGGAGTGGGTTGCAAAAGAAG
(SEQ ID NO: 13)

RiLB4 (*Pantoea* sp. deposited as NRRL-_____)

CTCGTGTTGTGAAATGTTGGGTAAAGTCCCGCAACGAGCGCAACCCTTATCCTTTGTT
GCCAGCGATTTCGGTCGGGAACCTCAAAGGAGACTGCCGGTGATAAACCGGAGGAAGG
TGGGGATGACGTCAAGTCATCATGGCCCTTACGAGTAGGGCTACACACGTGCTACA
ATGGCGCATACAAAGAGAAGCGACCTCGCGAGAGCAAGCGGACCTCACAAAGTGC
GTCGTAGTCCGGATCGGAGTCTGCAACTCGACTCCGTGAAGTCGGAATCGCTAGTAA
TCGTGGATCAGAATGCCACGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTC
ACACCATGGGAGTGGGTTGCAAAAGAAGTAGGTAGCTTAACCT
(SEQ ID NO: 14)

SY1 (*Pseudomonas* sp. deposited as NRRL-_____)

CCTTGCGCTATCAGATGAGCCTAGGTCGGATTAGCTAGTTGGTGAGGTAATGGCTCA
CCAAGGCGACGATCCGTAACCTGGTCTGAGAGGATGATCAGTCACACTGGAACCTGAG
ACACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGGCGAA
AGCCTGATCCAGCCATGCCGCGTGTGTGAAGAAGGTCTTCGGATTGTAAAGCACTTT
AAGGTGGGAGGAAGGGTTGTAGATTAATACTCTGCAATTTTGACGTTACCGCCAGA
ATAGCACCGGCTAACTCTGTGCCAGCAGCCGCGGTAATACAGAGGGTGCAAGCGTT
AATCGGAATTACTGGGCGTAAAGCGCGCGTAGGTGGTTTGTAAAGTCGGATGTGAA
ATCCCCGGGCTCAACCTGGGAACCTGCATCCGAACTGGCAAGCTAGAGTATGGTAG
AGGGTAGTGGAATTTCTGTGTAGCGGTGAAATGCGTAGATATAGGAAGGAACACC
AGTGGCGAAGGCGACTACCTGGACTGATACTGACACTGAGGTGCCAAAGCGTGGGG

AGCAAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGATGTCAACTAGCCG
 TTGGGAGCCTTGAGCTCTTAGTGGCGCAGCTAACGCATTAAGTTGACCGCCTGGGGA
 GTACGGCCGCAAGGTTAAACTCAAATGAATTGACGGGGGCCCCGCACAAGCGGTGG
 AGCATGTGGTTTAATTCGAAGCAACGCGAAGAACCTTACCAGGCCTTGACATCCAAT
 GAACTTTCCAGAGATGGATTGGTGCCTTCGGGAACATTGAGACAGGTGCTGCATGGC
 TGTCGTCAGCTCGTGTCTGAGATGTTGGGTAAAGTCCCGTAACGAGCGCAACCCTT
 GTCCTTAGTTACCAGCACGTTATGGTGGGCACTCTAAGGAGACTGCCGGTGACAAAC
 CGGAGGAAGGTGGGGATGACGTCAAGTCATCATGGCCCTTACGGCCTGGGCTACAC
 ACGTGCTACAATGGTCGGTACAGAGGGTTGCCAAGCCGCGAGGTGGAGCTAATCTC
 ACAAACCGATCGTAGTCCGGATCGCAGTCTGCAACTCGACTGCGTGAAGTCGGAA
 TCGCTAGTAATCGCGAATCAGAATGTCGCGGTGAATACGTTCCCGGGCCTTGTACAC
 ACCGCCCGTCACACCATGGGAGTGGGTTGCACCAGAAGTAGCTAGTCTAACCTTCG
 GGAGGAC (SEQ ID NO: 15)

SY5 (*Pseudomonas* sp. deposited as NRRL-_____)

AGCTCCACCTCGCGGCTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTGTAGC
 CCAGGCCGTAAGGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCGGTTTGTCA
 CCGGCAGTCTCCTTAGAGTGCCCAACATAACGTGCTGGTAACTAAGGACAAGGGTTG
 CGCTCGTTACGGGACTTAACCAACATCTCACGACACGAGCTGACGACAGCCATGC
 AGCACCTGTCTCAATGTTCCCGAAGGCACCAATCCATCTCTGGAAAGTTCATTGGAT
 GTCAAGGCCTGGTAAGGTTCTTCGCGTTGCTTCGAATTAAACCACATGCTCCACCGC
 TTGTGCGGGCCCCCGTCAATTCATTTGAGTTTTAACCTTGCGGCCGTACTCCCCAGGC
 GGTCAACTTAATGCGTTAGCTGCGCCACTAAGAGCTCAAGGCTCCCAACGGCTAGTT
 GACATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTT
 CGCACCTCAGTGTGAGTATCAGTCCAGGTAGTCGCCTTCGCCACTGGTGTTCCTTCCT
 ATATCTACGCATTTACCCGCTACACAGGAAATTCCTACTACCCTCTACCATACTCTAG
 CTTGCCAGTTTCGGATGCAGTTCCAGGTTGAGCCCGGGGA (SEQ ID NO: 16)

Endophytic nature of seed-associated bacteria

We closely examined the properties of several endophytic bacterial species, including *Pseudomonas fluorescens* Sandy LB4, *Pseudomonas* sp. West 9 and SY1 strains isolated from
 5 invasive *Phragmites*. In our in vitro assay using axenic Bermuda grass seedlings, we showed that on germination of seeds both bacteria colonize the root meristem and enter into meristematic cells. In the process of becoming intracellular the bacteria lose their cell walls and their rod shapes to become spherical L-form bacteria. As the root meristem proliferates the intracellular bacteria become distributed in most or all cells that make up the root. Intracellular bacteria are
 10 visible early on in young differentiating root cells as small spherical aniline blue staining structures. The use of DAB to stain for H₂O₂ shows that the intracellular bacteria are generally surrounded by H₂O₂ evident as red or brown staining. H₂O₂ is likely produced by the host plant

as a mechanism to control the intracellular bacterial endophytes. As tissues of the root, including root hairs become older the intracellular bacteria swell and their cytoplasmic contents disappear. Presence of H₂O₂ around intracellular bacteria suggests that the disappearance of bacteria is due to their degradation by the host cells through oxidation. In more mature root tissues all evidence of the intracellular bacteria completely disappears from plant cells. Bacterial cells that do not enter the root tip cells do not appear to elicit a strong reactive oxygen response and may not be degraded; instead they likely remain as intercellular endophytes in the mature root tissues.

Degradation of endophytic microbes as a nutritional source

Degradation of microbes in or on roots was proposed to be a mechanism of transfer of nutrients from microbes to plant (Paungfoo-Lonhienne et al., 2010; White et al., 2012; Beltran et al., 2015). Paungfoo-Lonhienne et al. (2014) denominated the microbe consumption process ‘rhizophagy’ because roots consumed microbes. For a mechanism like rhizophagy to provide substantial nutrients to plants the microbes must obtain their nutrients from outside roots, then enter roots and release their nutrients to the plant tissues. We have experimental evidence that in the case of the *Pseudomonas* endophytes of *Phragmites* the bacteria move from plant out into the soil and perhaps may return to recolonize the plant root meristem. If colonization of roots by the soil bacteria is occurring with any regularity a constant flow of nutrients from soil reserves to the plant may be occurring. In previous research we hypothesized that microbes may aid plants by scavenging organic or inorganic nitrogen (White et al., 2015; Soares et al., 2016a). However, all of the bacteria we isolated from *Phragmites* seeds showed the capacity to solubilize phosphate when grown on Pikovskayas agar. Further, *Pseudomonas fluorescens* and other fluorescent pseudomonads are known to secrete fluorescent siderophores called pyoverdines that actively bind iron (Visca, Imperi and Lamont, 2006). We confirmed that Sandy LB4 secreted a fluorescent pigment that fluoresced brightly in culture and that several other strains from *Phragmites* seeds (Table 1) fluoresced lightly in culture when viewed under an ultraviolet light, suggesting that several of our isolates possess the siderophores. Sandy LB4 was also shown to produce a secreted protease (Table 1) that might enable it to degrade microbial proteins present around roots and thus acquire organic nitrogen (White et al., 2015). The capacity of the motile pseudomonads to actively move out into the soil and obtain iron and solubilize and absorb phosphorus and nitrogenous nutrients could enable the pseudomonads to provide critical

nutrients to *Phragmites* to fuel its rapid growth. The endophytic pseudomonads could thus function as nutrient scavengers for their *Phragmites* seedlings (Zolg and Ottow, 1975).

Intracellular endophytes as elicitors of reactive oxygen

Bacterial endophytes that internally colonize root meristems and increase occurrence of reactive oxygen in the young differentiating tissues of the root may trigger the host to express oxidative stress resistance features to make the host more resistant to the internal oxidations. In a study of fungal endophytes in grasses it has been shown that endophytes increase the oxidative stress resistance in the host (White and Torres, 2009; Hamilton and Bauerle, 2012; Hamilton et al., 2012).

Growth promotion

Using several species to test for growth promotion capacity of the *Phragmites* endophytes described herein, (e.g., *P. fluorescens* Sandy LB4, *Pseudomonas* sp. West 9, *Microbacterium oxydans* B2 and SY1) and another bacterium, *Bosea thiooxidans* TBN from the distant family *Polygonaceae*, we found that several *Phragmites* isolates increased growth of grass compared to the non-inoculated treatment (Table 2A and Table 2B); while the bacterium *Bosea thiooxidans* TBN partially colonized grass seedlings and triggered root hair development but did not increase grass shoot or root growth in the soil experiment. We interpret that the *Phragmites* endophytes are compatible with the test grasses we used, but *Bosea thiooxidans* TBN is only partially compatible with the grasses, incompletely colonizing root hairs and failing to increase growth of the grass host in experiments.

In an experiment where we inoculated rice seeds with Sandy LB4, seedlings bearing the bacterium showed significantly more shoot growth compared to seedlings without the bacteria. Roots were also slightly longer in bacterial treated seedlings. It was also notable that roots bearing bacteria showed greater adherence of soil particles to roots after extraction from the soil. This indicates that the presence of the bacterium may affect the physical relationship of the plant root to the soil matrix around roots. The bacterium could be increasing extraction of nutrients from soil, which could account for the increased growth of seedlings when the bacterium is present.

Effects on seedling development

The effects of Sandy LB4 on *Poa annua* seedling growth were seen very early in development, including a very slightly increased germination rate and greater root branching (Table 4).

5 Suppression of soil borne disease

In co-culture experiments we demonstrated that Sandy LB4 and West 9 produced unknown antifungal substances that inhibited *Sclerotinia homeocarpa* and *Fusarium oxysporum*, while *M. oxydans* B2 did not appear to produce antifungal substances. In an experiment to assess protection from damping off disease caused by *Fusarium oxysporum* (Table 4) we showed that
 10 Sandy LB4, West 9 and *M. oxydans* B2 reduced damping off in seedlings when compared to controls inoculate with the pathogen; however Sandy LB4 was the most effective at reducing damping off. *Microbacterium oxydans* B2 was the least effective at reducing damping off disease.

In this same experiment we showed that the two pseudomonads originally inoculated
 15 onto seeds moved into the soil for some distance away from seedlings, suppressing mycelial growth in and on the surface of the soil. While *M. oxydans* B2 also moved out into the soil, it did not have any mycelial growth suppression effects and fungi grew readily over the surface of the soil in these treatments.

Some other *Pseudomonas spp.* that have been shown to be effective in biological control
 20 of diseases have been found to produce antifungal metabolites, including pyrrolnitrin, phenazines, pyoluteorin, hydrogen cyanide, etc. (Ligon et al., 2000).

Endophyte inhibition of competitor species

Through a series of experiments using *Phragmites* bacteria to inoculate dandelion (*Taraxacum officinale*) and curly dock (*Rumex crispus*) we showed that the strain Sandy LB4,
 25 whether alone or in mixtures of bacteria, substantially increased mortality rates of dandelion seedlings (Table 5) and in general reduced growth and vigor in both weed species. One of the frequent features of *Pseudomonas fluorescens* is the production of hydrogen cyanide (HCN). HCN combines with cytochrome oxidase and prevents aerobic respiration. Perhaps differential

susceptibility of the weeds and the grasses to HCN or other pseudomonad metabolite could be responsible for the inhibition of weed seedling growth and increased mortality. Differing sensitivity to HCN was previously suggested by Zeller, Brandle and Schmid (2007) to explain differing reactions of host and non-host species to *Pseudomonas* rhizobacteria. In this respect it is notable that many grasses secrete significant amounts of reactive oxygen (H_2O_2) from roots (White et al., 2012). H_2O_2 rapidly degrades HCN to form carbonate and ammonium ions (Yeddou et al., 2010). High production of hydrogen peroxide in grass roots could detoxify any HCN but also produce ammonia that could be absorbed and used to support plant growth. We did not see abundant H_2O_2 production in the roots of seedlings of either dandelion or curly dock.

Another possible mechanism to explain differential responses of host and non-host seedlings was evident when seedling roots were stained using DAB to visualize bacteria in roots. In grass roots bacteria remained as small discrete spherical bodies, likely flowing in the cytoplasm with the normal cyclosis. However, the bacteria in the non-host species appeared to form large clusters that appeared to occlude the root hairs, likely blocking normal cyclosis in the root hairs. Differential sensitivity to HCN or physical occlusion of absorptive root hairs or other structural issues may be responsible for the different responses of host grasses and non-host dandelion and curly dock seedlings to Sandy LB4.

Regardless of the mechanism of inhibition the presence of some endophytes in *Phragmites* that show capability to inhibit competitor plants could provide a mechanism by which *Phragmites* excludes competitor species from its stands. The allelopathic exclusion of plants from *Phragmites* stands has been observed and previously attributed to secreted chemical inhibitors such as gallic acid (Weidenhamer et al., 2013) and phytotoxicity of *Phragmites* residues (Uddin et al., 2014). It is possible that some of the allelopathic exclusion of competitors is a function of the growth promotional pseudomonads that are endophytic in plants and likely move out into soils colonizing and inhibiting competitor plant species that are not adapted to them.

Endophytes increase tolerance to salt stress

As shown in Figure 10, each of Sandy LB4, Sandy LB6 and SY1 demonstrably increase resistance to salt stress in rice in magenta box experiments.

Conclusions

It is clear from our studies that *Phragmites* seeds carry pseudomonads that may stimulate growth and increase nutrients available to the grass, but may also inhibit soil borne fungi, suppress diseases and inhibit competitor weed species in heterologous species. The pseudomonad symbionts of *Phragmites* appear to be diverse, multifunctional and appear to increase the competitive capacity of the host.

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20 **Example II**
Application of combinations of heterologous endophytic bacteria
to improve agronomic traits

As described in the previous example, the endophytic bacteria listed in Table 1 can be used alone or in combination to confer beneficial phenotypic changes to host plants. These
25 changes include one or more of increases in: root growth promotion, shoot growth promotion, resistance to salt stress, competition with undesirable plant species and resistance to fungus in a plant produced from the seed, as compared to a reference plant which is not treated with the inventive bacterial combination. The bacteria can act synergistically in combination or their effects may be additive in achieving the advantages set forth above. Such combinations can
30 include 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or all of the strains listed in Table 1. The strains may be present in differing amounts or ratios, e.g., 1:1, 1:2, 1:3, 1:4, 1:2:1, 1:5:1, etc. Exemplary

combinations include, without limitation: protease positive strains such as Sandy LB4 or Sandy LB6 in combination of antifungal (protease negative) strains SY1 or SY5. This combination provides increased functionality to the plant microbe combination. Protease positive strains WY6 or WY14 in combination with protease negative strain WY9y should also act to enhance
5 plant growth. The phosphatase negative strain *Microbacterium oxydans* B2 combined with phosphatase positive strains Sandy LB4 or Sandy LB6 should enhance phosphate scavenging capability as well as other processes effective to increase of plant growth.

The combined bacteria may be formulated to facilitate administration to target plants of interest, using methods described herein above. They may be lyophilized, and optionally
10 formulated into synthetic alginate beads for distribution into soil. Alternatively, they can be formulated as an aerosol for spraying on areas to be treated. Such methods are known to those of skill in the art of plant and crop propagation.

The formulations described above can also be added to certain fertilizer compositions, such as the controlled release fertilizer composition described in US Patent 9,266,787. Such
15 fertilizer compositions can optionally comprise one or more reagents selected from urea, ammonia, ammonium nitrate, ammonium sulfate, calcium nitrate, diammonium phosphate, monoammonium phosphate, potassium nitrate and sodium nitrate. monopotassium phosphate, dipotassium phosphate, tetrapotassium pyrophosphate, and potassium metaphosphate and optionally a macronutrient selected from the group consisting of sulfur, calcium and magnesium
20 and/or micronutrients including boron, copper, iron, manganese, molybdenum and zinc provided that such reagent does not interfere with the growth promoting action of the endophytic bacteria described herein.

These aforementioned formulations and compositions can further comprise a dispersing agent, such as those disclosed in US Patent 8,241,387.

25 While certain of the preferred embodiments of the present invention have been described and specifically exemplified above, it is not intended that the invention be limited to such embodiments. Various modifications may be made thereto without departing from the scope and
30 spirit of the present invention, as set forth in the following claims.

What is claimed is:

1. A method of increasing root and shoot growth of a grass plant host, comprising inoculating the host grass or seed with one or more biologically pure *Pseudomonas spp.*, *Microbacterium spp.* or *Enterobacter spp.* endophyte strains, which are heterologously disposed
5 to said grass plant host, said strains being effective to increase one or more of root growth promotion, shoot growth promotion, resistance to salt stress, competition with undesirable plant species and resistance to fungus in a plant produced from the seed, as compared to a reference plant grown under the same conditions.
2. The method of claim 1, wherein said bacteria comprise specific rDNA sequences selected
10 from the group consisting of SEQ ID NO: 1-16 or sequences having at least 97% identity therewith.
3. The method of claim 2, wherein said endophytic bacteria comprise three or more of the bacteria selected from Sandy LB4, Sandy LB6, Sandy Y8, RoY12, RoLB13w, WY9y, WY6,
15 West 9, WY14, RiY3, SandyY7w, RiLB4, SY1, SY5, WY9w, and *Microbacterium oxydans* B2 and at least one of said bacteria are SY1 or SY5.
4. The method of claim 3, wherein said bacteria are selected from the group consisting of Sandy LB4, Sandy LB6, West 9, and SY1.
- 20 5. The method of claim 3, wherein said bacteria comprise Sandy LB4.
6. The method of claim 3, wherein said bacteria comprise Sandy LB6.
7. The method of claim 3, wherein said bacteria comprise SY1.
8. The method of claim 3, wherein said bacteria comprise *Microbacterium oxydans* B2
9. The method of claim 4, wherein the inoculated host grass exhibits suppressed growth of soil
25 borne fungal pathogens of said host grass relative to host grasses which have not been inoculated with said endophyte.

10. The method of claim 2 wherein said inoculated plant host exhibits increased resistance to disease relative to untreated host plants.

11. The method of claim 2, wherein said inoculated plants reduce competitiveness of distantly related competitor weeds of said grass plant host.

5 12. The method of claim 2, wherein the host grass is a turf and forage grass selected from *Agrostis* spp., *Poa* spp., *Festuca* spp., *Lolium* spp., *Cynodon* spp., *Zoysia* spp., *Koeleria* spp., *Paspalum* spp., *Panicum* spp., *Stenotaphrum* spp., *Eremochloa* spp. and *Danthonia* spp., *Triticum aestivum*, *Oryza sativa*, *Avena sativa*, *Hordeum vulgare*, *Panicum miliaceum*, *Eleusine coracana*, *Sorghum bicolor*, *Pennisetum glaucum*, *Setaria italica*, *Secale cereal*, *Zea mays*,
10 *Agrostis* spp., *Poa* spp., *Festuca* spp., *Lolium* spp., *Cynodon* spp., *Zoysia* spp., *Koeleria* spp., and *Danthonia* spp.

13. The method of claim 2, wherein the host grass is a food grass crop selected from *Oryza sativa*, *Triticum aestivum*, *Avena sativa*, *Hordeum vulgare*, *Secale cereal*, and *Zea mays*.

14. The method of claim 2, wherein the host grass is a Pooideae grass.

15 15. The method of claim 1, wherein said bacteria are present in a synthetic seed ball.

16. The method of claim 1, wherein said plant host is sprayed with a liquid formulation comprising said bacteria.

17. The method of claim 3, wherein said bacteria are present in a controlled release fertilizer formulation.

20 18. A synthetic combination comprising at least one purified bacterial population in association with a seed of a grass plant host or seed wherein the purified bacterial population comprises a seed bacterial endophyte that is heterologously disposed to the grass plant host or seed and is selected from one or more of Sandy LB4, Sandy LB6, Sandy Y8, RoY12, RoLB13w, WY9y, WY6, West 9, WY14, RiY3, SandyY7w, WY9w, RiLB4, SY1, SY5 and *Microbacterium*
25 *oxydans* B2, wherein the seed bacterial endophyte is present in the synthetic combination at a concentration of at least 10^3 CFU/seed on the surface of a seed, and in an amount effective to increase one or more of root growth promotion, shoot growth promotion, resistance to salt stress,

competition with undesirable plant species and increase resistance to fungus in a plant produced from the seed, as compared to a reference plant grown under the same conditions.

19. The synthetic combination of claim 18, wherein said bacteria are Sandy LB4, Sandy LB6,
5 West 9 and SY1, which are lyophilized.

20. The synthetic combination of claim 18 or 19, wherein said bacteria are encapsulated in an alginate bead formulation.

21. The synthetic combination of claim 18, wherein said bacteria are present in a seed ball.

22. The synthetic combination of claim 18, wherein said bacteria is a combination of Sandy
10 LB4, Sandy LB6, West 9 and SY1 and said seed is a rice seed.

23. The synthetic combination of claim 18, wherein said bacteria is a combination of Sandy LB4, Sandy LB6, West 9 and SY1 and said seed is a grass seed from a grass selected from *Agrostis spp.*, *Poa spp.*, *Festuca spp.*, *Lolium spp.*, *Cynodon spp.*, *Zoysia spp.*, *Koleria spp.*, and *Danthonia spp.*

15 24. The synthetic combination of claim 18, wherein said bacteria is a combination of Sandy LB4, Sandy LB6, West 9 and SY1 and said seed is a grass seed from a grass selected from *Oryza sativa*, *Triticum aestivum*, *Avena sativa*, *Hordeum vulgare*, *Secale cereal*, and *Zea may.s*

25. The synthetic combination of claim 18, wherein said bacteria is *Microbacterium oxydans* B2 and said seed is a rice seed.

20 26. A method of reducing undesirable weed growth in soils comprising turf, forage or food grasses, comprising drenching or treating the soil with one or more biologically pure *Pseudomonas spp.* endophyte strains wherein said strains reduce weed growth relative to weed growth observed in untreated soil, wherein said one or more *Pseudomonas spp.* is Sandy LB4 and/or West 9.

25 27. The method of claim 26, wherein said at least one biologically pure *Pseudomonas spp.* endophyte strain is obtained from Phragmites and said bacteria are lyophilized.

28. The synthetic combination of claim 18, comprising a combination of bacteria selected from the group consisting of

i) protease positive strains consisting of Sandy LB4 and Sandy LB6 in combination with strains SY1 or SY5;

5 ii) protease positive strains WY6 and/or WY14 in combination with protease negative strain WY9y; and

iii) phosphatase negative strain *Microbacterium oxydans* B2 in combination with phosphatase positive strains Sandy LB4 and/or Sandy LB6.

29. A method of reducing undesirable weed growth in soils comprising turf, forage or food

10 grasses, comprising drenching or treating or coating seeds from said turf, forage or food grass with a combination of Sandy LB4, Sandy LB6, West 9 and SY1 bacterial strains and said seed is a grass seed selected from *Oryza sativa*, *Triticum aestivum*, *Avena sativa*, *Hordeum vulgare*, *Secale cereal*, and *Zea mays*.

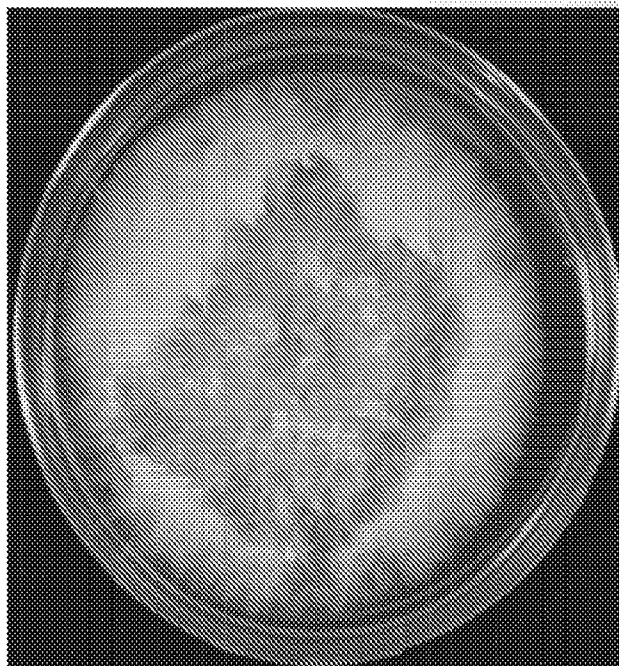


Fig. 1A

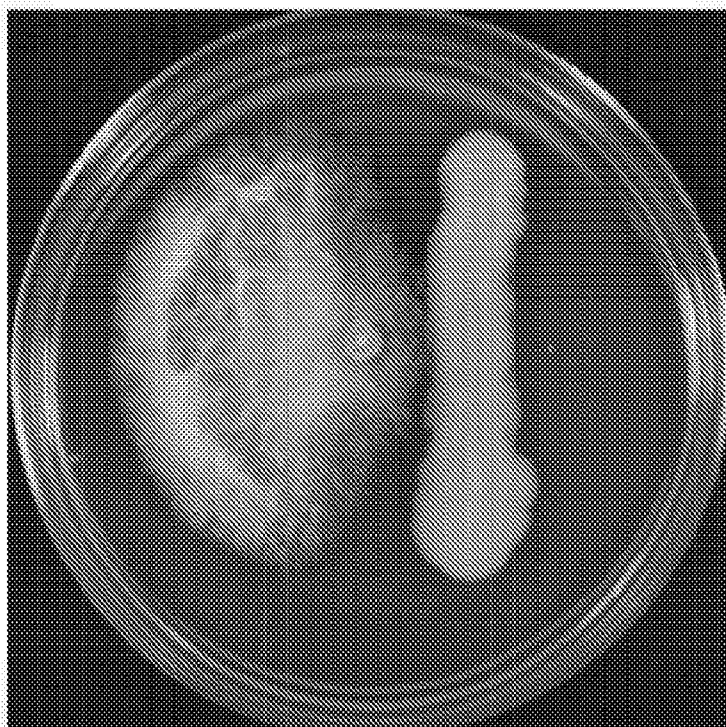


Fig. 1B

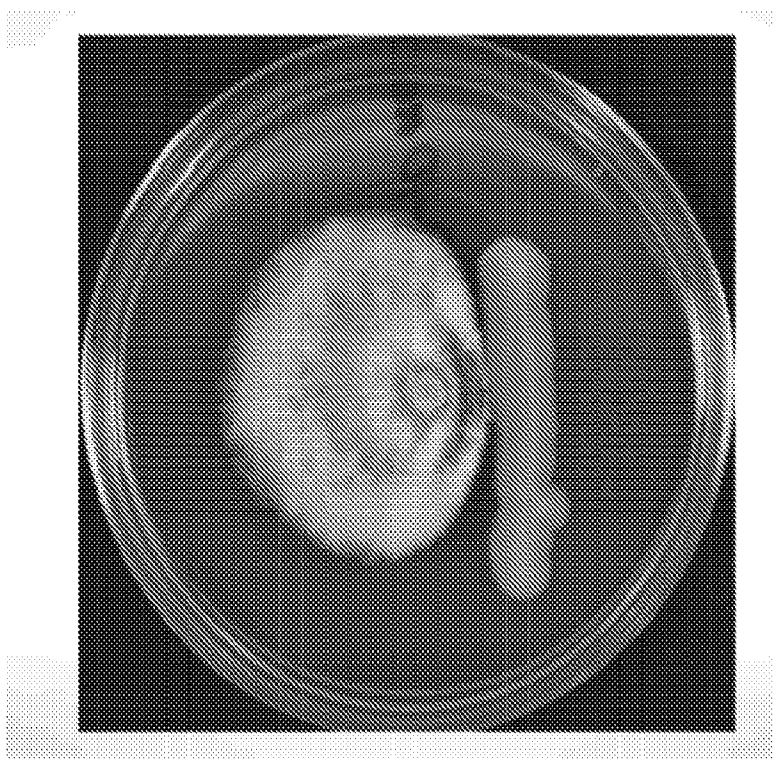


Fig. 1C

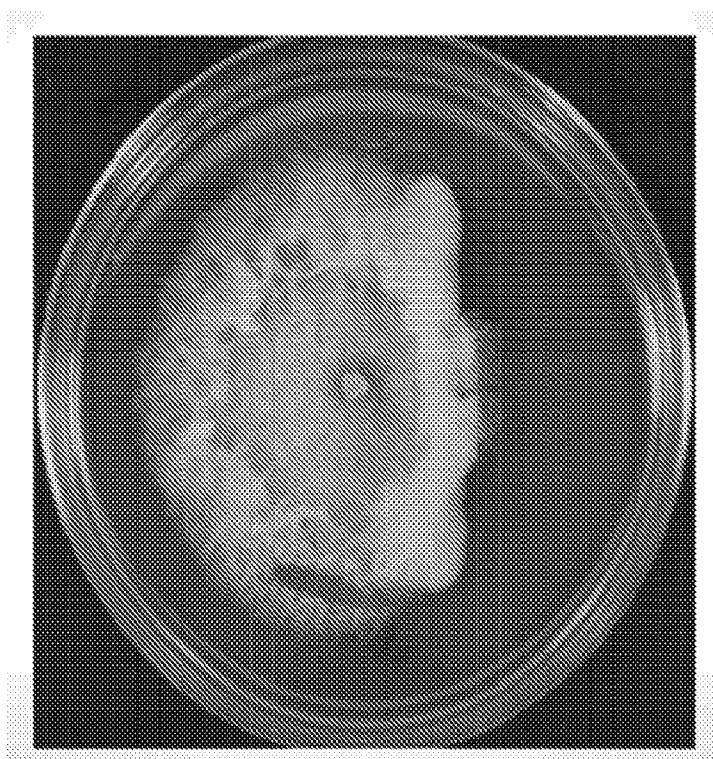


Fig. 1D

Seedlings growing on 0.7% agarose and stained with
DAB for reactive oxygen

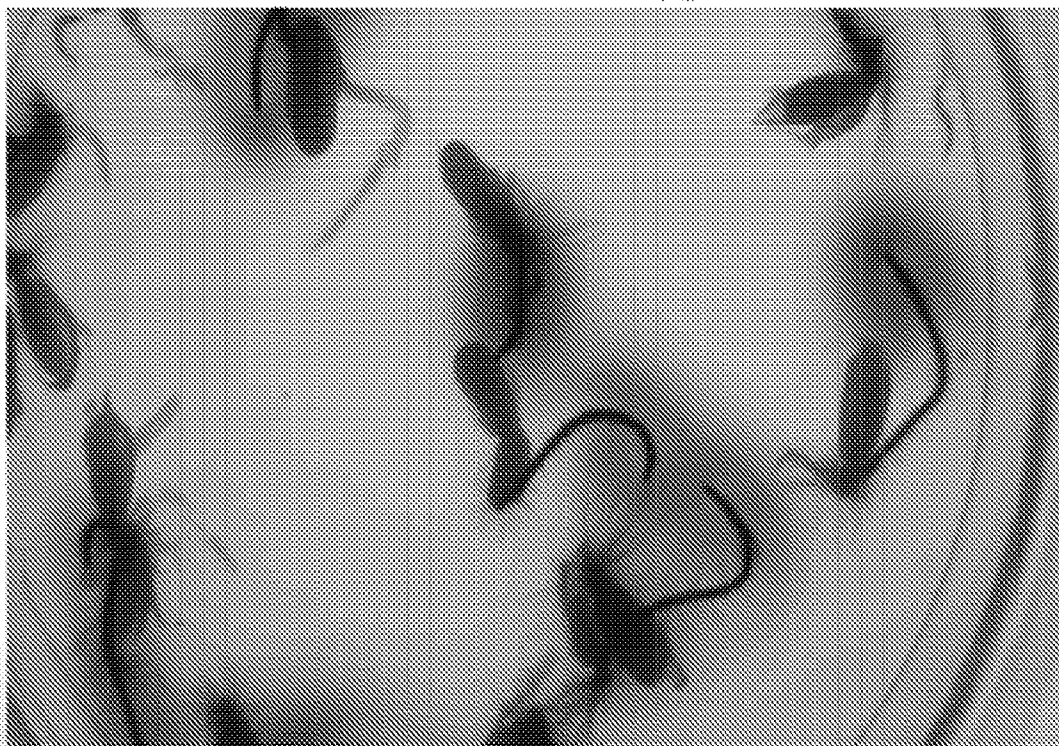


Fig. 2A

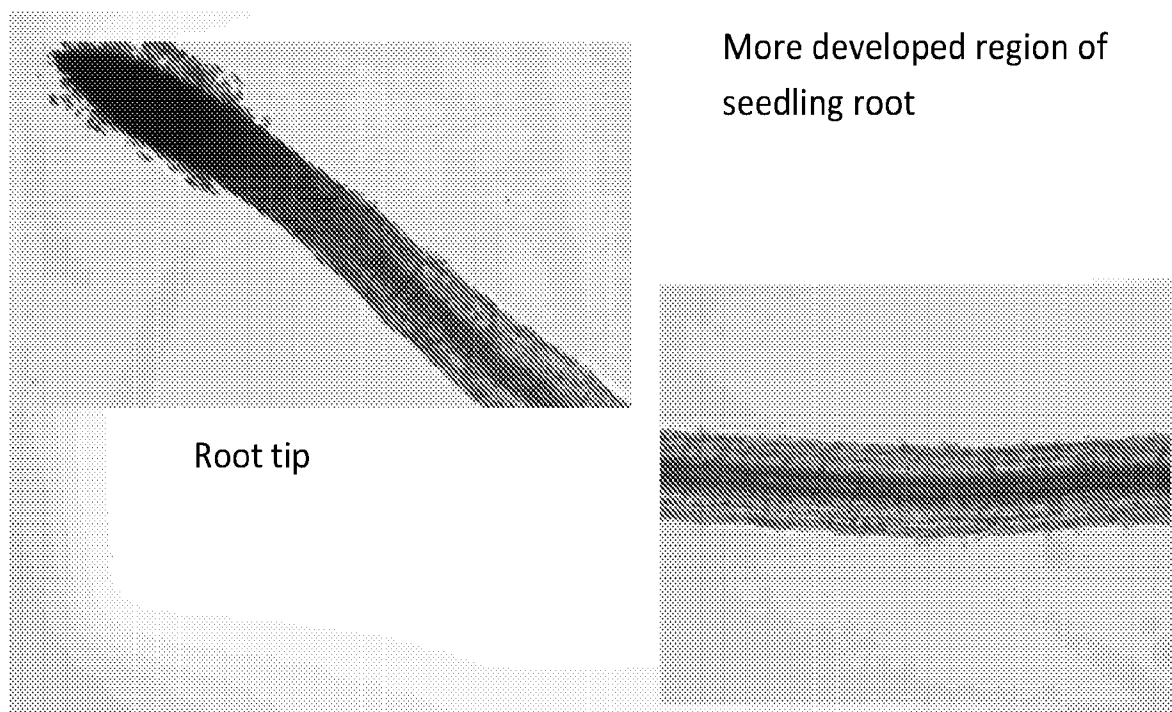


Fig. 2B

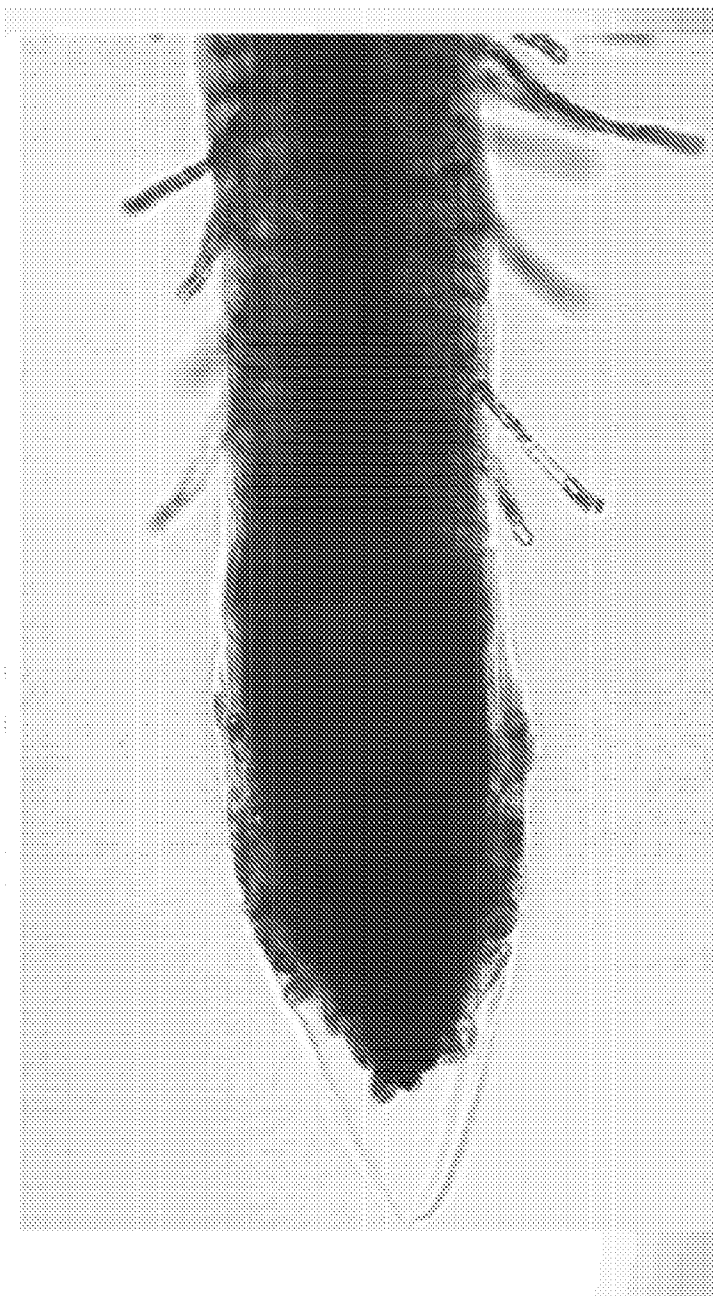


Fig. 3A

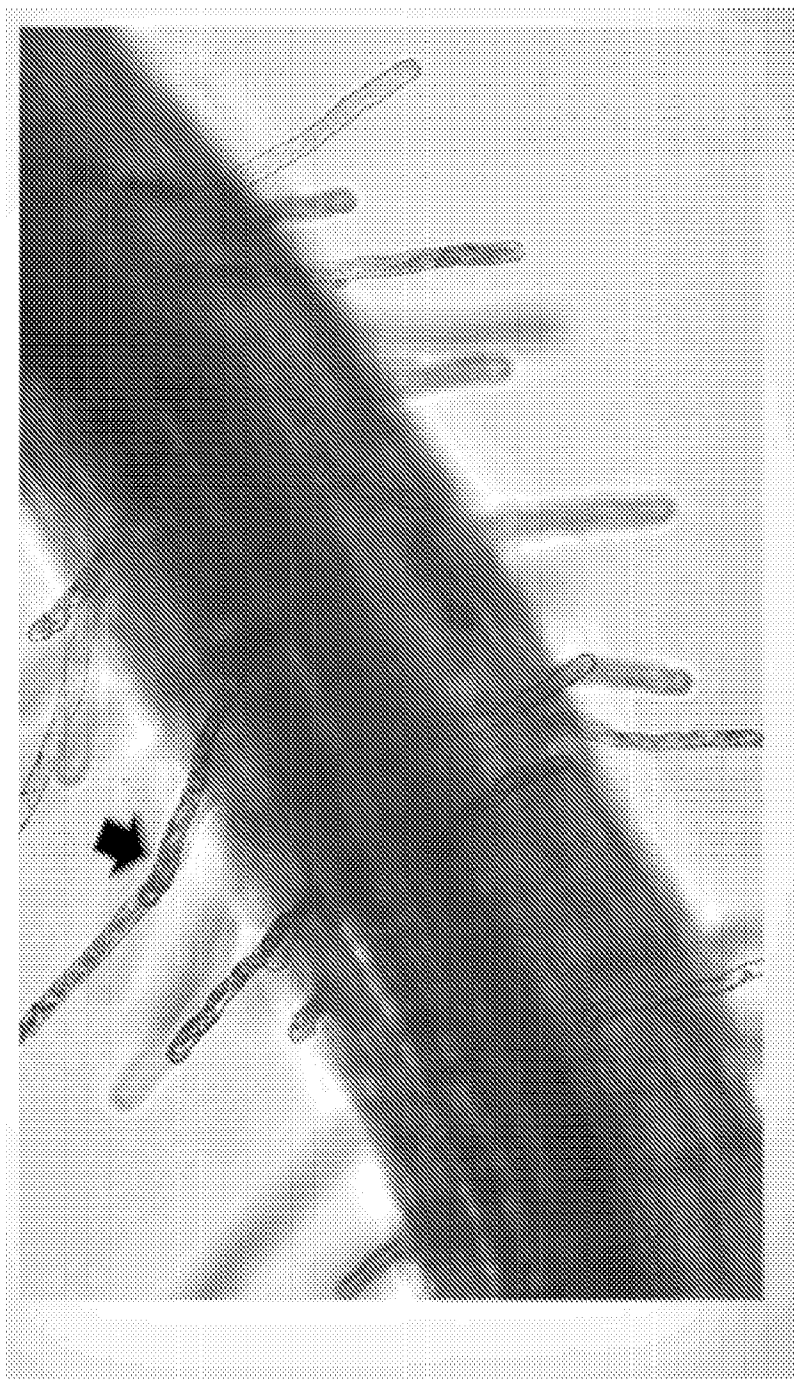


Fig. 3B

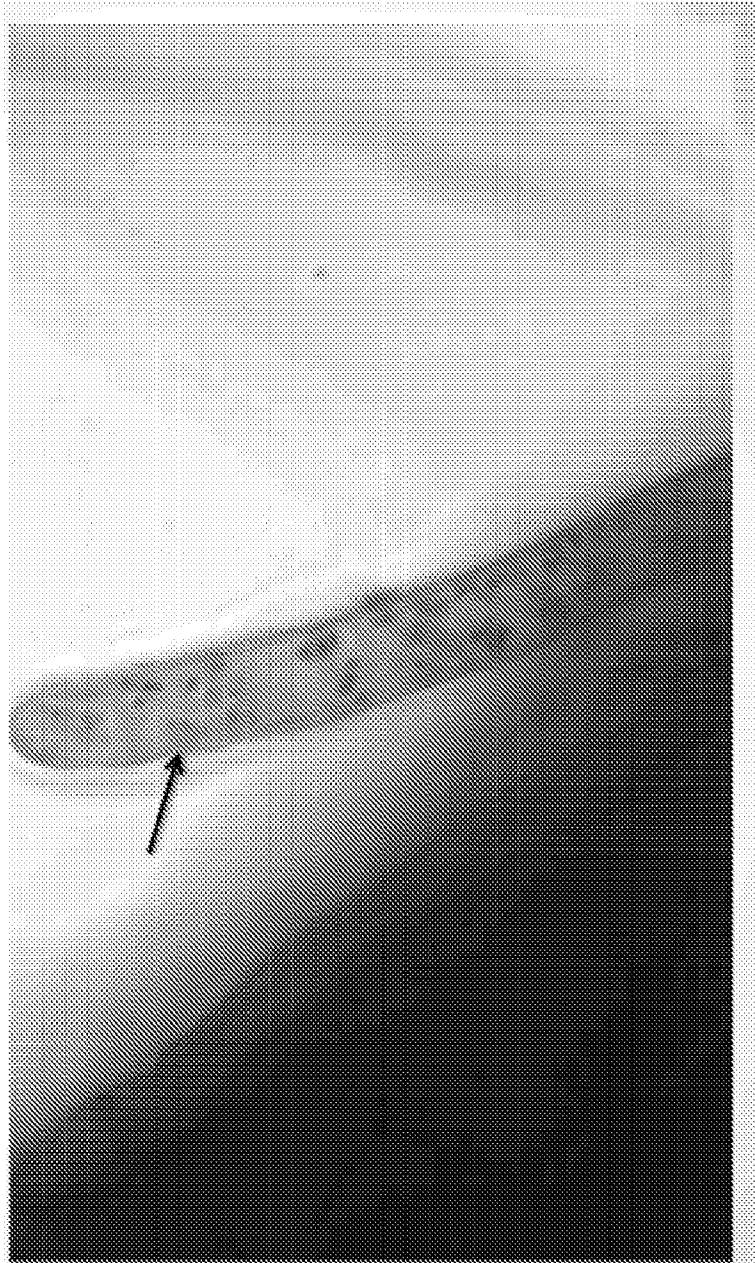


Fig. 4A



Fig. 4B

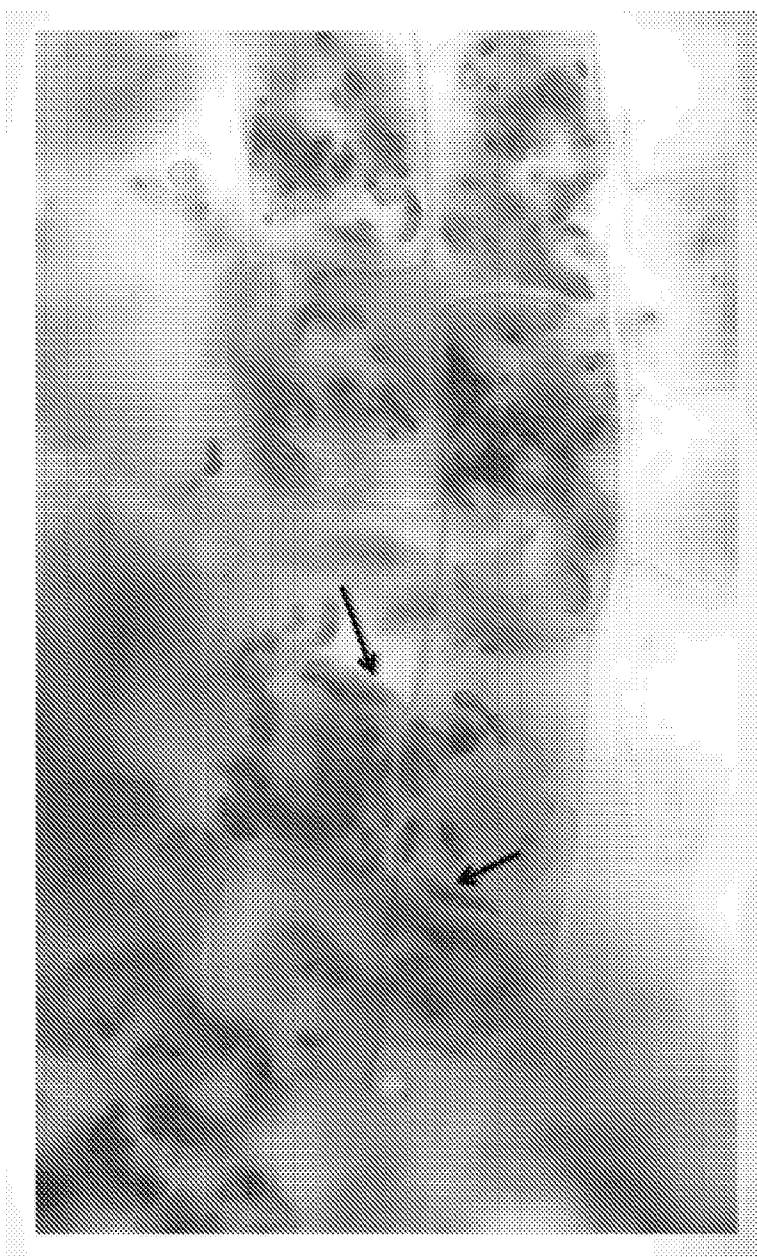


Fig. 4C

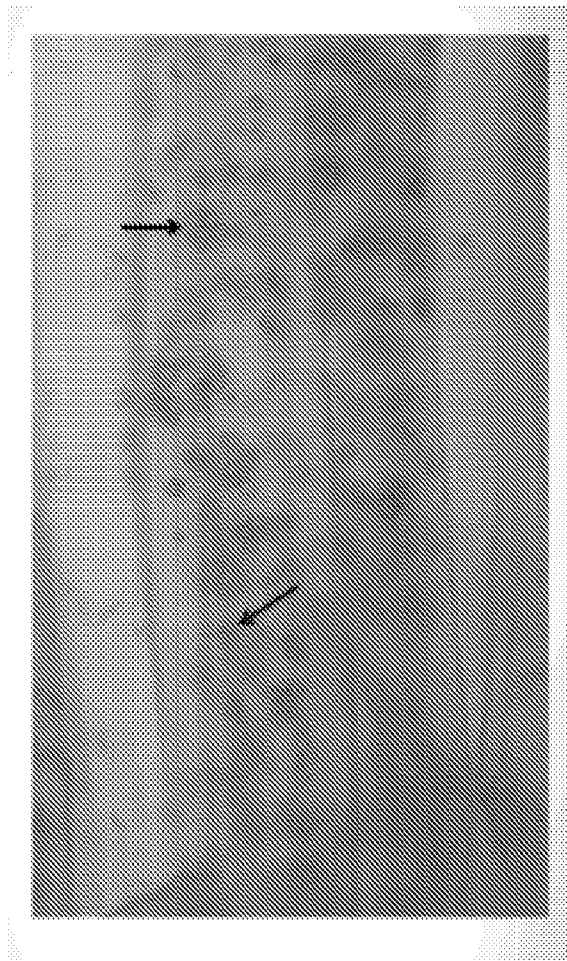


Fig. 4D

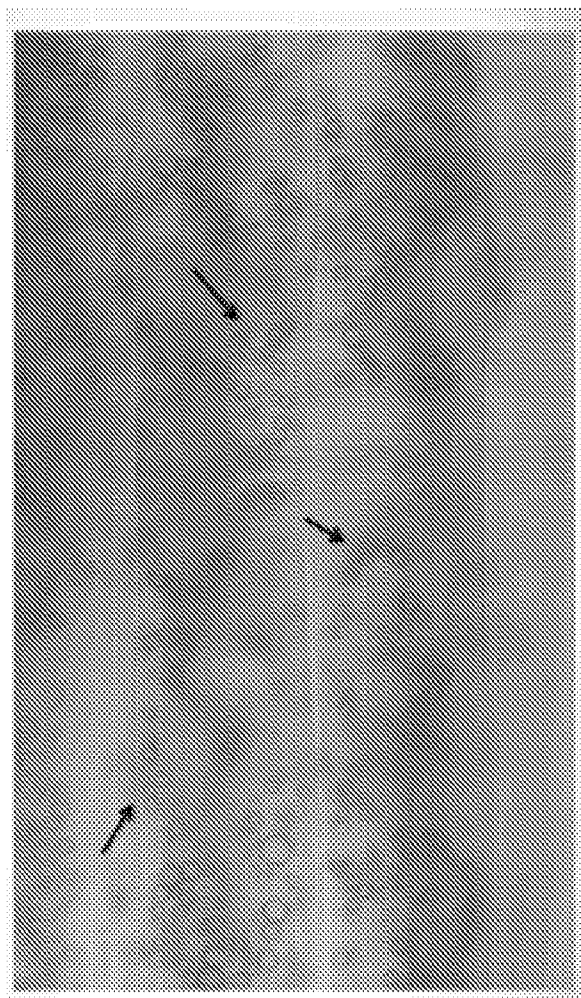


Fig. 4E



Fig. 5A

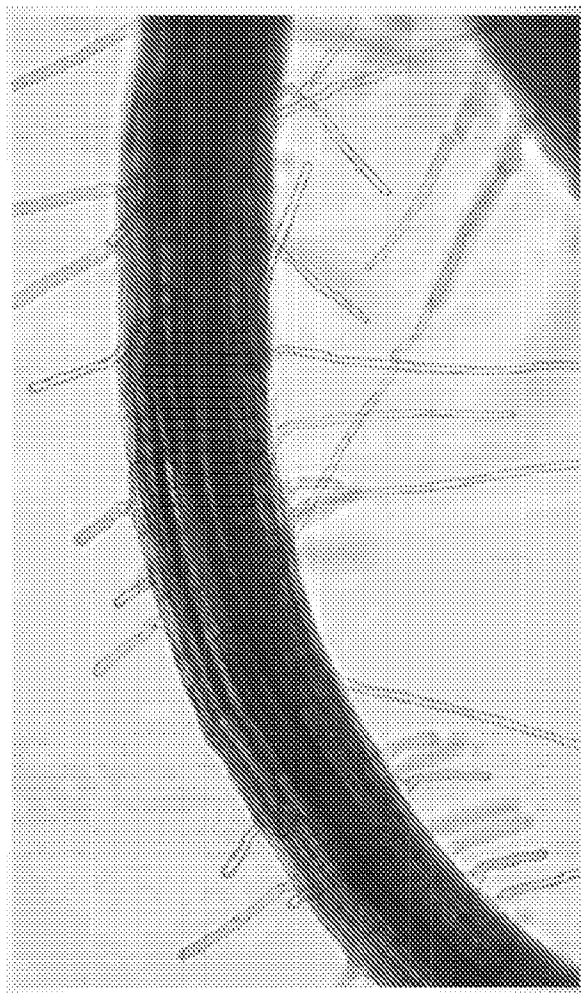


Fig. 5B

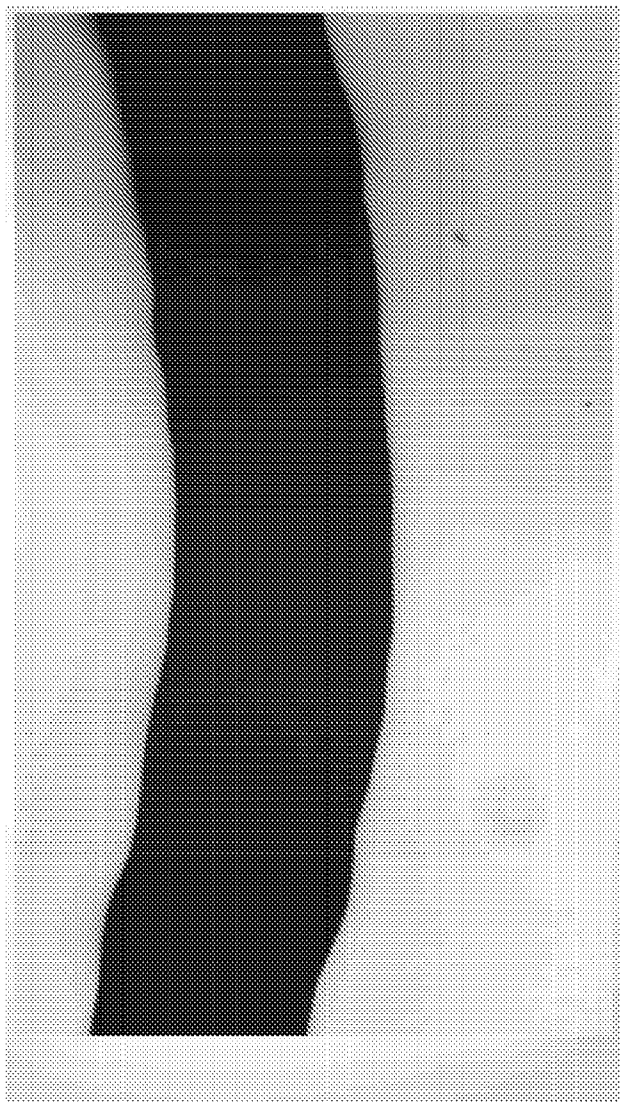


Fig. 6A

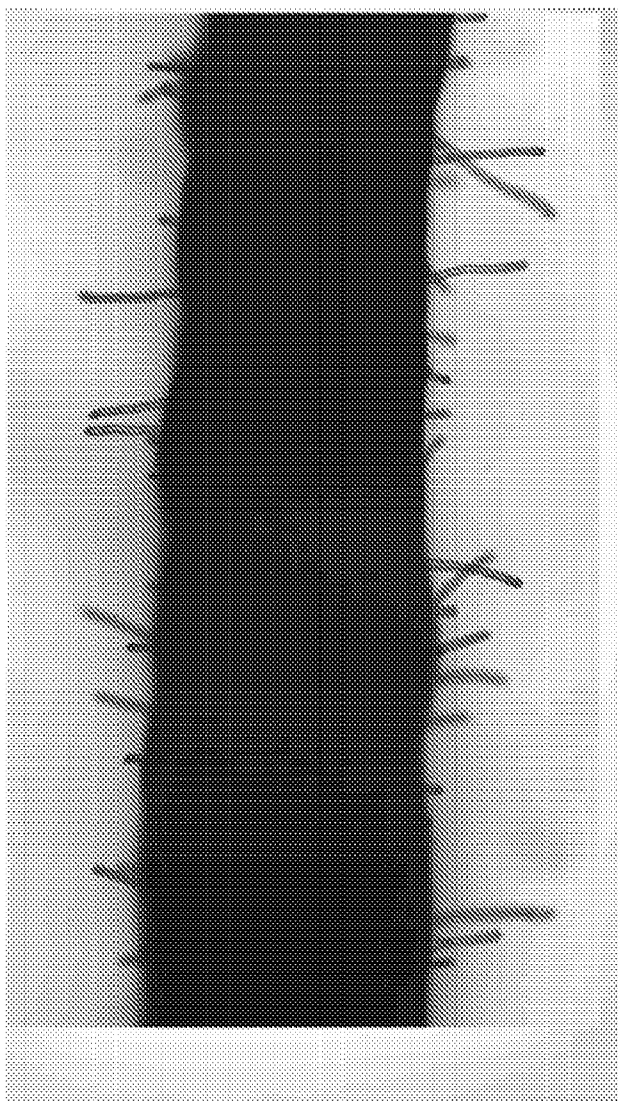


Fig. 6B

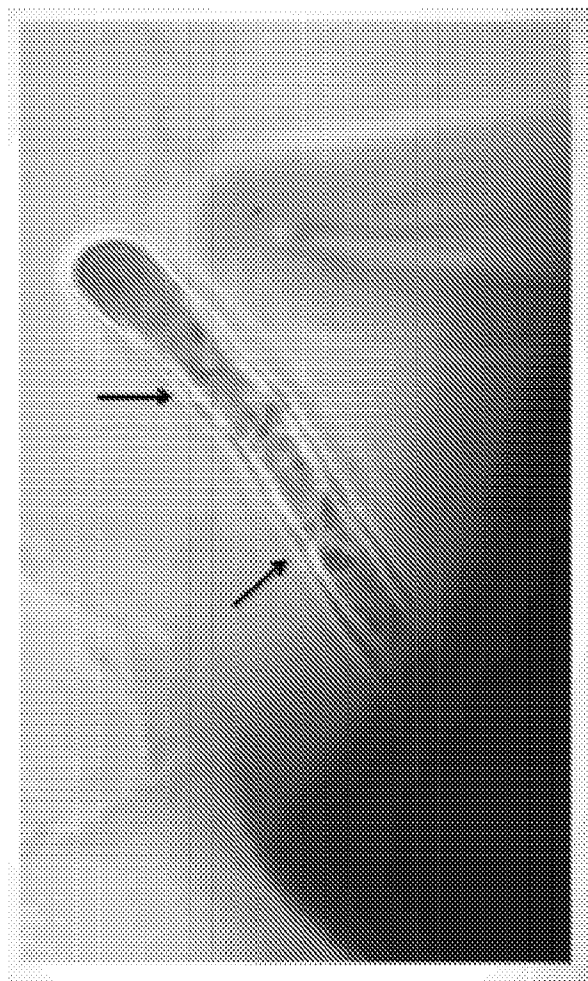


Fig. 6C

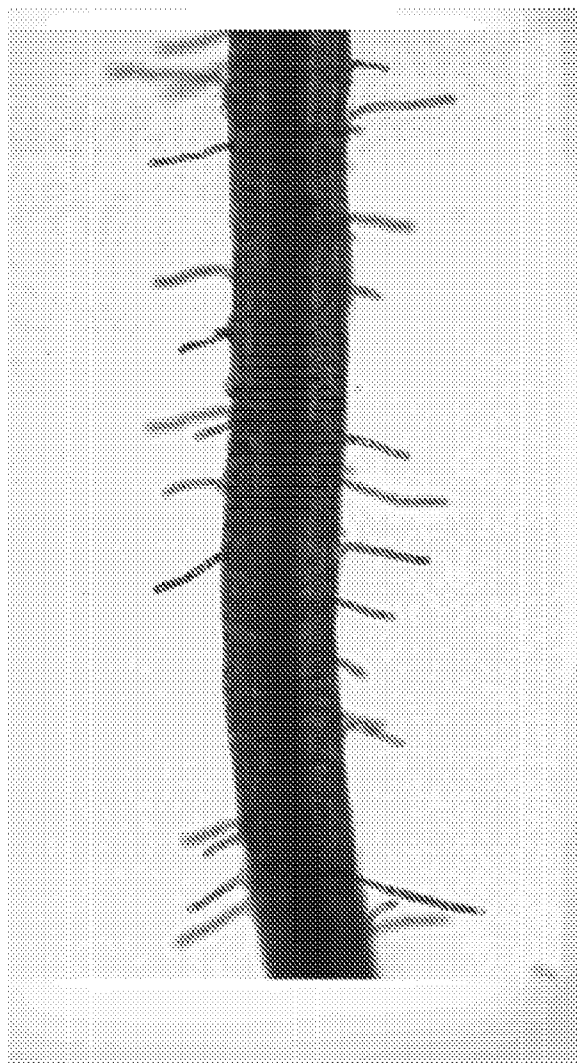


Fig. 7A

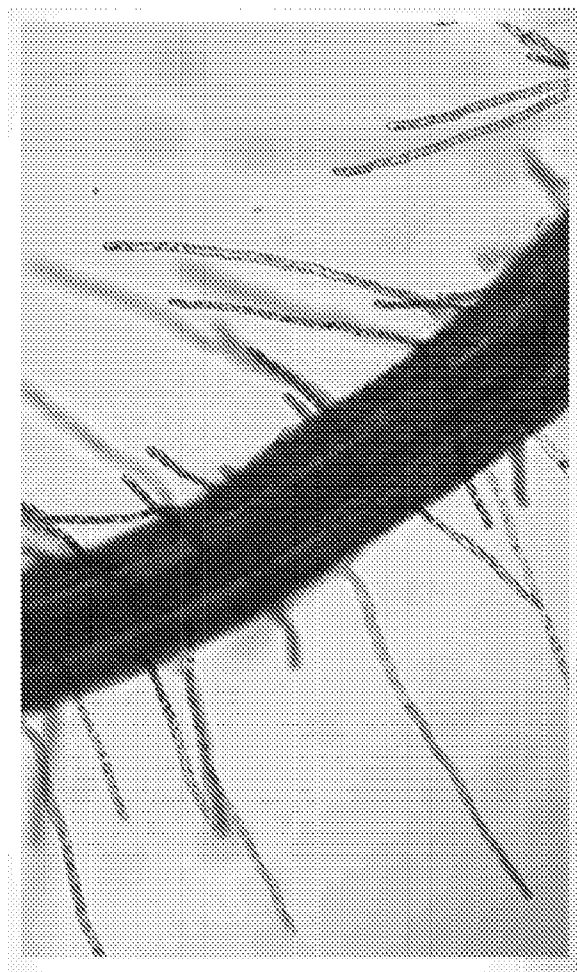


Fig. 7B

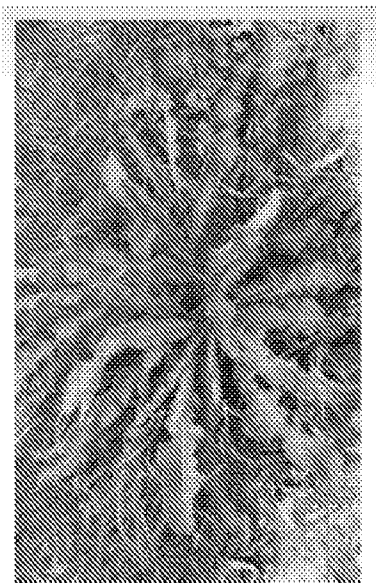


Fig. 8A

Strain Sandy LB4 from Phragmites in root hair
of *Rumex crispus*

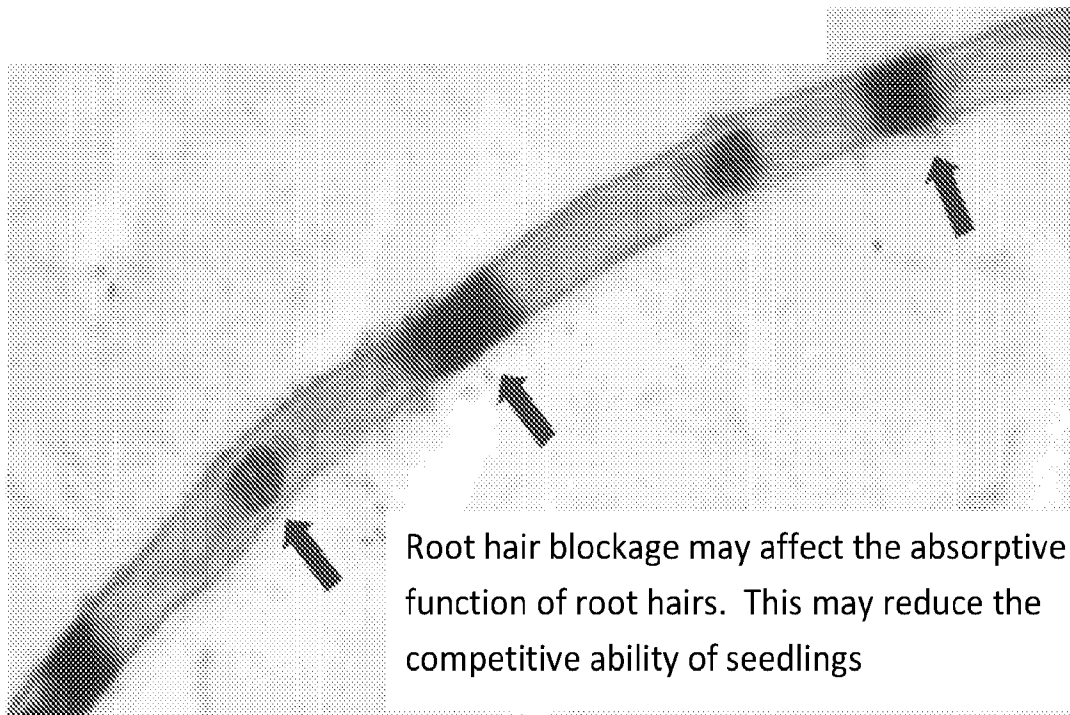


Fig. 8B

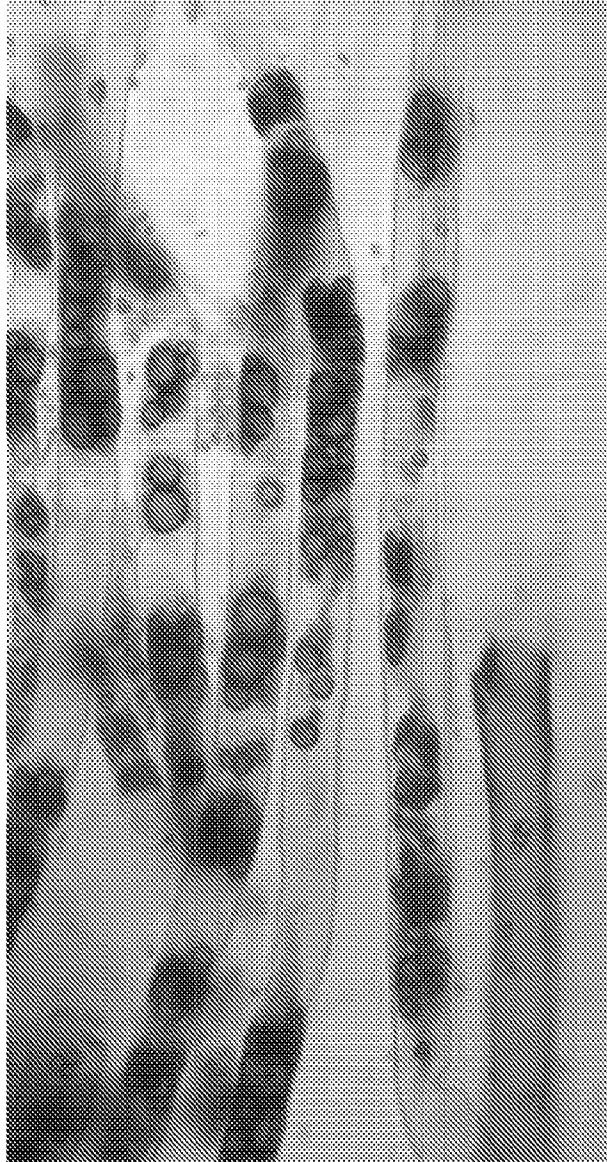


Fig. 8C

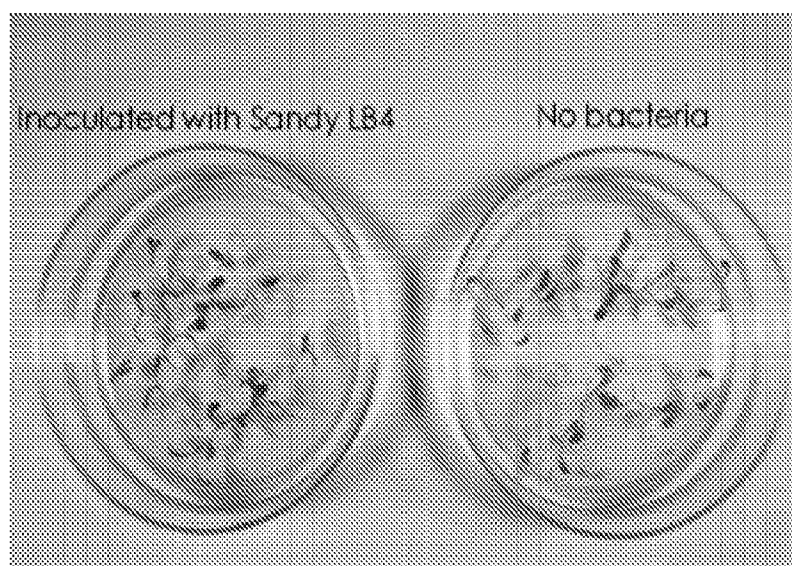


Fig. 9A

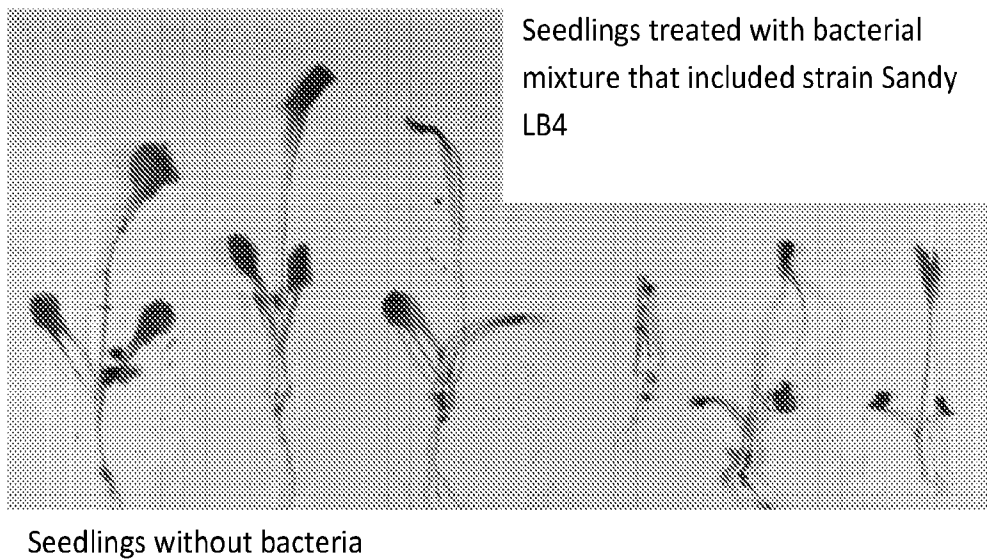


Fig. 9B

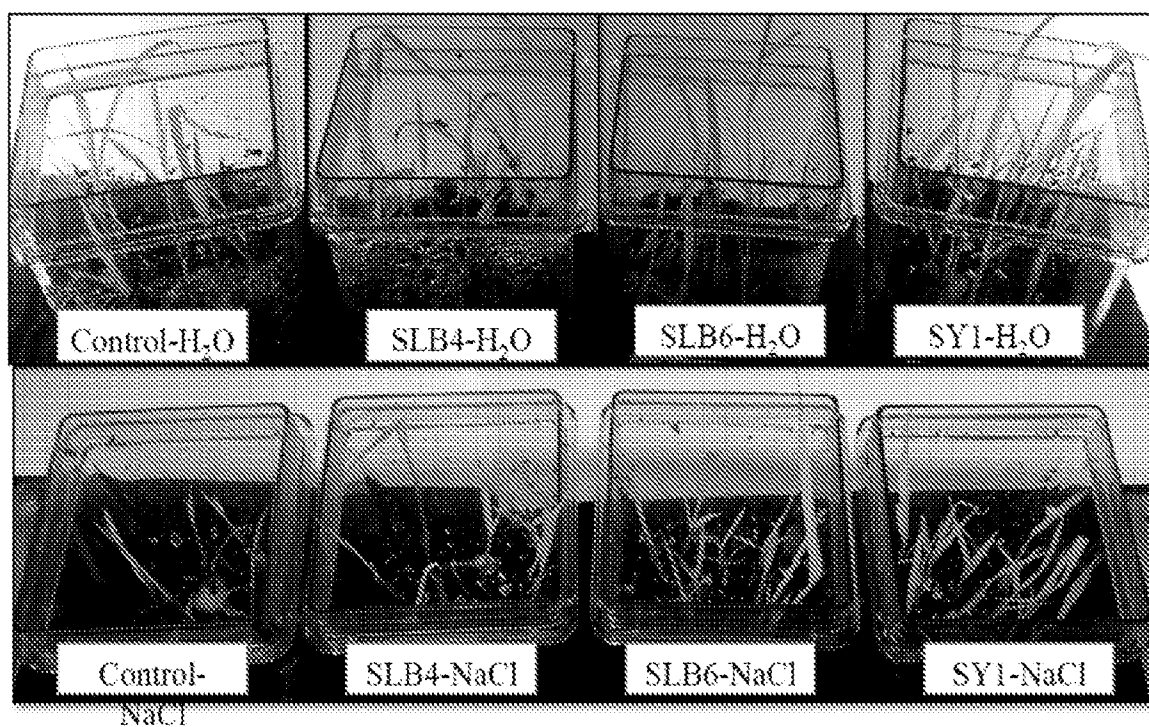


Figure 10

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US2018/043259

A. CLASSIFICATION OF SUBJECT MATTER

IPC(8) - A01C 1/06; A01G 1/00; A01G 7/00; A01G 7/06; A01H 5/00; A01H 17/00; A01N 63/00 (2018.01)
 CPC - A01G 7/00; A01G 22/00; A01N 63/00; A01N 63/02; A01N 63/04; C12N 1/20; C12R 1/01; C12R 1/38; C12R 1/39 (2018.08)

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

See Search History document

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

See Search History document

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

See Search History document

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	US 2016/0316760 A1 (INDIGO AGRICULTURE, INC.) 03 November 2016 (03.11.2016) entire document	1, 2, 10, 12-16, 18, 20, 21
Y		3-9, 11, 17, 19, 22-25, 28, 29
Y	US 2009/0324564 A1 (MAEDA) 31 December 2009 (31.12.2009) entire document	3-9, 17, 19, 22-24, 26, 27, 29
Y	"KY074468: Pseudomonas syringae strain BAV3981 16S ribosomal RNA gene, partial sequence," NCBI GenBank, Accession No. KY074468, 20 December 2016 (20.12.2016), Pgs. 1 of 1. Retrieved from the Internet: <www.ncbi.nlm.nih.gov/nuccore/KY074468> on 21 September 2018 (21.09.2018). entire document	3-9, 17, 19, 22-24, 29
Y	SOARES et al. "Functional Role of Bacteria from Invasive Phragmites australis in Promotion of Host Growth," Microbial Ecology, 01 August 2016 (01.08.2016), Vol. 72, Iss. 2, Pgs. 407-417. entire document	8, 11, 25, 27, 28
Y	"AB109886: Pseudomonas graminis Gene for 16S rRNA, Partial Sequence, Strain: KF701," NCBI Genbank, Accession No. AB109886, 29 July 2009 (29.07.2009), Pgs. 1-2. Retrieved from the Internet: <www.ncbi.nlm.nih.gov/nuccore/AB109886> on 21 September 2018 (21.09.2018). entire document	19, 22-24, 29
Y	US 2017/0020138 A1 (SYMBIOTA, INC.) 26 January 2017 (26.01.2017) entire document	23



Further documents are listed in the continuation of Box C.



See patent family annex.

Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier application or patent but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T"

later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X"

document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y"

document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&"

document member of the same patent family

Date of the actual completion of the international search

21 September 2018

Date of mailing of the international search report

04 OCT 2018

Name and mailing address of the ISA/US

Mail Stop PCT, Attn: ISA/US, Commissioner for Patents
 P.O. Box 1450, Alexandria, VA 22313-1450
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Authorized officer

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INTERNATIONAL SEARCH REPORT

International application No.

PCT/US2018/043259

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	US 2013/0031673 A1 (GRANDLIC et al) 31 January 2013 (31.01.2013) entire document	26, 27, 29
P,X	US 2018/0020672 A1 (RUTGERS, THE STATE UNIVERSITY OF NEW JERSEY) 25 January 2018 (25.01.2018) entire document	1-29
P,X	US 2017/0238552 A1 (UNIVERSITY OF GUELPH) 24 August 2017 (24.08.2017) entire document	1-29
A	"JF901350: Endophytic bacterium 41P-2 16S ribosomal RNA gene, partial sequence," NCBI GenBank, Accession No. JF901350, 07 December 2012 (07.12.2012), Pgs. 1-2. Retrieved from the Internet: <www.ncbi.nlm.nih.gov/nuccore/JF901350> on 21 September 2018 (21.09.2018). entire document	1-29
A	"AY753173: Enterobacter endosymbiont of Metaseiulus occidentalis clone pAJ240 16S ribosomal RNA gene, partial sequence," NCBI GenBank, Accession No. AY753173, Pgs. 1-2. Retrieved from the Internet: <www.ncbi.nlm.nih.gov/nuccore/AY753173> on 21 September 2018 (21.09.2018). entire document	1-29
A	PROENCA et al. "Diversity of Bacteria Associated with Bursaphelenchus xylophilus and Other Nematodes Isolated from Pinus pinaster Trees with Pine Wilt Disease," PLoS One, 09 December 2010 (09.12.2010), Vol. 5, Iss. 1, e15191, Pgs. 1-9. entire document	1-29
A	RODRIGUEZ et al. "Fungal endophytes: diversity and functional roles," New Phytologist, 27 March 2009 (27.03.2009), Vol. 182, No. 2, Pgs. 314-330. entire document	1-29
A	"HQ538790: Pseudomonas sp. M47tronco1 16S ribosomal RNA gene, partial sequence," NCBI GenBank, Accession No. HQ538790, 02 May 2012 (02.05.2012), Pgs. 1-2. Retrieved from the Internet: <www.ncbi.nlm.nih.gov/nuccore/HQ538790> on 21 September 2018 (21.09.2018). entire document	1-29
A	"KP860310: Microbacterium oxydans strain B2 16S ribosomal RNA gene, partial sequence," NCBI GenBank, Accession No. KP860310, 08 June 2016 (08.06.2016), Pgs. 1-2. Retrieved from the Internet: <www.ncbi.nlm.nih.gov/nuccore/KP860310> on 21 September 2018 (21.09.2018). entire document	1-29

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US2018/043259

Box No. I Nucleotide and/or amino acid sequence(s) (Continuation of item 1.c of the first sheet)

1. With regard to any nucleotide and/or amino acid sequence disclosed in the international application, the international search was carried out on the basis of a sequence listing:

a. ☒ forming part of the international application as filed:

☐ in the form of an Annex C/ST.25 text file.

☒ on paper or in the form of an image file.

b. ☒ furnished together with the international application under PCT Rule 13ter.1(a) for the purposes of international search only in the form of an Annex C/ST.25 text file.

c. ☐ furnished subsequent to the international filing date for the purposes of international search only:

☐ in the form of an Annex C/ST.25 text file (Rule 13ter.1(a)).

☐ on paper or in the form of an image file (Rule 13ter.1(b) and Administrative Instructions, Section 713).

2. ☐ In addition, in the case that more than one version or copy of a sequence listing has been filed or furnished, the required statements that the information in the subsequent or additional copies is identical to that forming part of the application as filed or does not go beyond the application as filed, as appropriate, were furnished.

3. Additional comments:

SEQ ID NOs: 1-16 were searched.