

Table S1. Disease compatibility of specific combinations of race-specific secreted effectors (Ave1 and V2) on resistant cultivars of tomatoes containing different combinations of the race 1 (Ve1) and race 2 (V2) resistance phenotypes.

| Race (isolates) | Race gene | Resistance combination | | | |
|------------------------|------------|------------------------|-----------|-----------|-----------|
| | | Ve1-/V2- | Ve1+/V2+ | Ve1+/V2- | Ve1-/V2+ |
| Race 1/2 (Le1087, JR2) | Ave1+/Av2+ | Disease | Resistant | Resistant | Resistant |
| Race 2 (To22, Ca36) | Ave1-/Av2+ | Disease | Resistant | Disease | Resistant |
| Race 3 (HoMCF, KJ14a) | Ave1-/Av2- | Disease | Disease | Disease | Disease |
| Race 1/3 (Vdp4) | Ave1+/Av2- | Disease | Resistant | Resistant | Disease |

Table S2. Biocontrol agents (BCAs) and biologicals with biocontrol activity against *Verticillium dahliae*.

| Biocontrol organism | Source | Mechanism of action /efficacy | References |
|--|---------------------------------|--|------------|
| <i>Acinetobacter calcoaceticus</i> FS339 | Root (Tomato) | Reduced mycelial growth, reduced disease index, increased chitinase, siderophores, protease activity | [249] |
| <i>Bacillus axarquiensis</i> TUBP1 protein | Rhizosphere (Cotton) | Anomalies in fungal hyphae and conidia, reduced spore germination, targeted plasma membrane of <i>V. dahliae</i> | [250] |
| <i>Bacillus</i> spp. | Composted ginned cotton residue | Reduce disease severity and wilt symptoms, antibiosis | [161] |
| <i>Bacillus thuringiensis</i> | Rhizosphere (Tomato) | Mycolitic chitinases, polypeptide synthetases, bacillibactin | [251] |
| <i>Bacillus velezensis</i> AL7 | Soil (Cotton) | Synthesize antifungal antibiotics | [153] |

| Biocontrol organism | Source | Mechanism of action /efficacy | References |
|--|---|---|------------|
| <i>Bacillus velezensis</i> C2 | Endophyte (Tomato) | Reduced disease incidence, antifungal activities (synthesis of lipopeptides, presence of volatile metabolites, produce lytic enzymes) | [252] |
| <i>Bacillus velezensis</i> OEE1 | Endophyte (Olive) | Reduced fungal growth (92%), disease severity index, percentage of dead plants, and microsclerotia density | [159] |
| <i>Burkholderia gladioli</i> pv. <i>agaricicola</i> | - | Reduced mycelial growth and disease incidence | [253] |
| Cell-free mix of <i>Sphingobacterium</i> A1 and <i>Bacillus tequilensis</i> C-9 (1:9) | Rhizosphere (A1 from <i>Resina ferulae</i> and C-9 from cotton) | Reduced spore production (97.8%), germination (100%), and virulence protein of <i>V. dahliae</i> | [155] |
| <i>Chaetomium globosum</i> CEF-082 | Endophyte (Cotton) | Biocontrol effect 59%, Regulate multiple metabolic pathways, induce defense responses | [254] |
| <i>Cryptoseria myriophylloides</i> , <i>Laminaria digitata</i> , <i>Fucus spiralis</i> | Algal extracts | Reduced disease severity, Increase defense response in tomato due to polyphenol oxidases, peroxidases | [255] |
| The essential oil from <i>Thymus</i> spp. | Thyme | Inhibition of mycelia and microsclerotia; disease reduction | [157] |

| Biocontrol organism | Source | Mechanism of action /efficacy | References |
|---|-------------------------|--|------------|
| <i>Fusarium solani</i> CEF559 | Endophyte (cotton) | Reduced colony growth (75%) and sporulation (80), PR genes and genes in lignin metabolism pathway upregulated, greenhouse control efficacy (60%) and field efficacy (30-56%) | [160] |
| <i>Metarhizium brunneum</i> | - | Reduce germination of microsclerotia, inhibited the hyphal formation | [256] |
| <i>Paenibacillus polymyxa</i> ShX301 | Rhizosphere (Cotton) | Reduced disease incidence and disease severity | [257] |
| <i>Pseudomonas fluorescences</i> FS167 | Root endophyte (Tomato) | Reduced mycelial growth, reduced disease index, chitinase, siderophores, protease | [249] |
| <i>Pseudomonas mosselli</i> FS67 | Root endophyte (Tomato) | Reduced mycelial growth, reduced disease index, siderophores | [249] |
| <i>Pseudomonas</i> spp. | Rhizosphere (Olive) | Reduced disease onset and development, disease incidence, plant mortality, phytase, and catalase activities | [258] |
| <i>Purpureocillium lilacinum</i> QLP12 | Soil (Eggplant) | Reduce disease index in greenhouse and field | [259] |
| <i>Stenotrophomonas maltophilia</i> FS300 | Root endophyte (Tomato) | Reduced mycelial growth, reduced disease index, chitinase, siderophores, protease | [249] |
| <i>Trichoderma atroviride</i> | | Antifungal activity | [154] |

| Biocontrol organism | Source | Mechanism of action /efficacy | References |
|---|----------------------------|---|------------|
| <i>Trichoderma harzianum</i> | Rhizosphere (Pistachio) | Reduced mycelial growth and disease severity, mycoparasitism | [260] |
| Triterpene derivatives (at 10 µg/ml) | Latex (Euphorbia) | Reduced disease severity, stunting, and vascular discoloration; enhanced peroxidase and polyphenol oxidase activities in tomato | [158] |

Table S3. Some examples of organic amendments (OAs) with suppressiveness against *Verticillium dahliae*.

| Organic amendments | Test crop/source | Mechanism of action | References |
|-------------------------------|------------------------|--|------------|
| Solid olive oil waste compost | Olive | Reduce mycelial growth; and microsclerotia viability (52%-76%) | [261] |
| Olive mill compost | Cotton | Low β-glucosidase activity; high oligotrophic actinomycete populations, Mycelial growth inhibition, Reduced disease severity, lower microsclerotial concentrations | [262] |
| Broccoli residue | Rotation with eggplant | Reduced disease incidence by 53%, reduced <i>V. dahliae</i> DNA in soil | [263] |
| Compost tea | Corn straw | Reduced mycelial growth and conidial germination by 91% and 78% respectively in strawberry; control efficacy of 42% in greenhouse | [264] |

| Organic amendments | Test crop/source | Mechanism of action | References |
|---|------------------|--|------------|
| Grape marc compost | Olive | Reduced microsclerotia density; disease incidence; | [156,265] |
| Plant compost (grape pomace compost; olive pomace/dairy manure compost; mixed crop residue compost) | Bell pepper | reduced pathogen population in fields (29%-42%) up to 14 weeks post-application; Compost extracts reduced <i>V. dahliae</i> growth by 25-50% in Petri dish assay | [163] |
| Fresh manure (4 kg m ⁻² sheep manure + 1 kg m ⁻² composted poultry litter) covered with transparent plastic | Bell pepper | Reduced disease incidence to less than 1% | [167] |
| 4% Dried spearmint or oregano | Tomato | Reduced disease to visually no symptoms | [266] |
| Composts with various compositions | Eggplant | Reduced disease indicated by disease suppression index | [267] |
| Tomato waste compost | Eggplant | Reduced disease severity and fungal colonization | [268] |
| Turkey litter compost | Eggplant | Reduced disease severity | [268] |
| Various compositions of steam-explosion liquid waste, Agro-industrial residues/waste, and plant green waste composts | Eggplant | Reduced disease indicated by disease suppression index (43%-65%), showed richness in potential biocontrol agents | [166] |

| Organic amendments | Test crop/source | Mechanism of action | References |
|---|------------------|--|------------|
| Mustard | Tomato | Suppressed verticillium wilt with “biofumigant” properties | [164] |
| Sudangrass | Tomato | Suppressed verticillium wilt with “biofumigant” properties | [164] |
| Plant-based compost | Tomato | Reduced disease intensity, showed richness in culturable biocontrol agents | [162] |
| Compost of winery residues including grape stalks and grape pomace | Eggplant | Reduced disease incidence, severity and plant mortality, high total phenol content | [269] |
| Compost of tomato pulp with sawdust and chipping wood as bulking agents with a ratio of 2:1:1 v/v/v | Eggplant | Reduced disease incidence, severity and plant mortality, high total phenol content | [269] |
| Compost of organic fraction of municipal solid waste | Eggplant | Reduced disease incidence, severity and plant mortality, high total phenol content | [269] |
| Compost consisted of 73% olive mill extracted press cake, 24% wastewater, and 3% olive leaves | Eggplant | Reduced disease incidence, severity and plant mortality, high total phenol content | [269] |

Table S4. Examples of sources of carbon from recent studies for anaerobic soil disinfestation of soil-borne pathogens including *Verticillium dahliae*.

| The carbon source for ASD | Host plant | Targeted pathogen | References |
|---------------------------|-------------------------------------|--|-----------------------|
| Alfalfa | Cucumber | <i>Rhizoctonia solani</i> | [174,270] |
| Bean dregs | - | <i>Fusarium oxysporum</i> | [137] |
| Broccoli + Rice bran | Strawberry | <i>Fusarium oxysporum</i> f. sp. <i>fragariae</i> ; <i>Verticillium dahliae</i> | [271] |
| Corn straw | - | - | [174] |
| Ethanol | Tomato; Cucumber; Strawberry | <i>Fusarium oxysporum</i> f. sp. <i>lycopersici</i> ; <i>Fusarium oxysporum</i> ; <i>Verticillium dahliae</i> ; <i>Pyrenochaeta lycopersici</i> ; <i>Colletotrichum coccodes</i> ; <i>Rhizoctonia solani</i> <i>Meloidogyne</i> spp. | [137,140,270,272,273] |
| FL-104 rye + Rice bran | Strawberry | <i>Fusarium oxysporum</i> f. sp. <i>fragariae</i> | [271] |
| Grape pomace | Strawberry | <i>Verticillium dahliae</i> | [140,175,274] |
| Italian rye + Rice bran | Strawberry | <i>Fusarium oxysporum</i> f. sp. <i>fragariae</i> | [271] |
| Molasses | Tomato Lettuce; Mustard green | <i>Plasmodiophora brassicae</i> ; <i>Meloidogyne</i> spp. | [169,273,275] |

| The carbon source for ASD | Host plant | Targeted pathogen | References |
|---|------------|---|---------------|
| | | <i>Verticillium dahliae</i> ; <i>Pyrenochaeta lycopersici</i> ; <i>Colletotrichum coccodes</i> | |
| Molasses + composted poultry litter (CPL) | Tomato | <i>Meloidogyne</i> spp. | [276] |
| Mustard + Rice bran | Strawberry | <i>Fusarium oxysporum</i> f. sp. <i>fragariae</i> | [271] |
| Mustard seed meal | Strawberry | <i>Verticillium dahliae</i> | [140,169] |
| Onion waste | Strawberry | <i>Verticillium dahliae</i> | [140] |
| Rice bran | Strawberry | <i>Verticillium dahliae</i> ; <i>Fusarium oxysporum</i> f. sp. <i>fragariae</i> | [140,169,271] |
| Sudan grass + Rice bran | Strawberry | <i>Fusarium oxysporum</i> f. sp. <i>fragariae</i> | [271] |
| Sugarcane bagasse | - | <i>Fusarium oxysporum</i> | [137] |

| The carbon source for ASD | Host plant | Targeted pathogen | References |
|---------------------------|---------------------------|--|---------------|
| Tomato pomace | - | - | [169,175,274] |
| Triticale + rice bran | Strawberry | <i>Fusarium oxysporum</i> f. sp. <i>fragariae</i> | [271] |
| Wheat bran | Tomato; Strawberry | <i>Plasmodiophora brassicae</i> ; <i>Meloidogyne</i> spp. <i>Verticillium dahliae</i> ; <i>Pyrenochaeta lycopersici</i> ; <i>Colletotrichum coccodes</i> ; <i>Meloidogyne</i> spp. | [140,273,275] |
| Wheat bran + Molasses | Lettuce; Mustard green | <i>Plasmodiophora brassicae</i> ; <i>Meloidogyne</i> spp. | [275] |