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## Protecting maize from rootworm damage with the combined application of arbuscular mycorrhizal fungi, *Pseudomonas* bacteria and entomopathogenic nematodes

Geoffrey Jaffuel<sup>1</sup>, Nicola Imperiali<sup>2</sup>, Kent Shelby<sup>3</sup>, Raquel Campos-Herrera<sup>1,4</sup>, Ryan Geisert<sup>3</sup>, Monika Maurhofer<sup>5</sup>, Joyce Loper<sup>6,7</sup>, Christoph Keel<sup>1</sup>, Ted C. J. Turlings<sup>8</sup> & Bruce E. Hibbard<sup>8</sup>

*Diabrotica virgifera virgifera* LeConte, the western corn rootworm (WCR), is the most destructive pest of maize in North America, and has recently spread across central Europe. Its subterranean larval stages are hard to reach with pesticides and it has evolved resistance to conventional management practices. The application of beneficial soil organisms is being considered as a sustainable and environmental friendly alternative. In a previous study, the combined application in wheat fields of arbuscular mycorrhizal fungi, entomopathogenic *Pseudomonas* bacteria, and entomopathogenic nematodes was found to promote growth and protection against a natural pest infestation, without negative cross effects. Because of the insect-killing capacity of the bacteria and nematodes, we hypothesized that the application of these organisms would have similar or even greater beneficial effects in WCR-infested maize fields. During three consecutive years (2015–2017), we conducted trials in Missouri (USA) in which we applied the three organisms, alone or in combinations, in plots that were artificially infested with WCR and in non-infested control plots. For two of the three trials, we found that in plots treated with entomopathogenic nematodes and/or entomopathogenic *Pseudomonas* bacteria, roots were less damaged than the roots of plants in control plots. During one year, WCR survival was significantly lower in plots treated with *Pseudomonas* than in control plots, and the surviving larvae that were recovered from these plots were lighter. The bacterial and nematodes treatments also enhanced yield, assessed as total grain weight, in one of the trials. The effects of the treatments varied considerable among the three years, but they were always positive for the plants.

*Diabrotica virgifera virgifera* LeConte, the western corn rootworm (WCR), causes significant damage to maize (*Zea mays* L.) across North America, as well as across Central and Eastern Europe<sup>1,2</sup>. The larval stage is the most damaging, as it feeds on root hairs, cortical tissue, and tunnels inside the roots of maize plants. This can lead to the destruction of roots<sup>3,4</sup>, which hampers the uptake of water and nutrients from the soil<sup>5</sup>, and increases plant's

<sup>1</sup>FARCE Laboratory, Institute of Biology, University of Neuchâtel, Neuchâtel, Switzerland. <sup>2</sup>Department of Fundamental Microbiology, University of Lausanne, Lausanne, Switzerland. <sup>3</sup>Biological Control of Insects Research, US Department of Agriculture, Agricultural Research Service, Columbia, MO, USA. <sup>4</sup>Instituto de Ciencias de la Vid y del Vino, CSIC-Universidad de La Rioja-Gobierno de La Rioja, Logroño, Spain. <sup>5</sup>Institute of Integrative Biology, ETH Zurich, Zurich, Switzerland. <sup>6</sup>Department of Botany and Plant Pathology, Oregon State University, Corvallis, OR, USA. <sup>7</sup>Horticultural Crops Research Laboratory, US Department of Agriculture, Agricultural Research Service, Corvallis, OR, USA. <sup>8</sup>Plant Genetics Research Unit, US Department of Agriculture-ARS, University of Missouri, Columbia, MO, USA. Geoffrey Jaffuel and Nicola Imperiali contributed equally. Correspondence and requests for materials should be addressed to T.C.J.T. (email: [ted.turlings@unine.ch](mailto:ted.turlings@unine.ch)) or B.E.H. (email: [Bruce.Hibbard@ars.usda.gov](mailto:Bruce.Hibbard@ars.usda.gov))

susceptibility to lodging<sup>6</sup>. Often, roots are fully pruned by older larvae that move up to the base of the stalk<sup>7</sup>. In affected areas in the US, WCR larvae can cause tremendous yield losses<sup>1,8–10</sup>.

From the time that it was discovered as a pest<sup>11</sup> until 1946, the only successful management option was crop rotation. Since then, WCR management has also included granular and liquid soil insecticides, and more recently insecticidal seed treatments and transgenic Bt maize<sup>12–14</sup>. Over time, WCR has developed resistance to most insecticides classes<sup>15–17</sup>. Crop rotation is still highly effective against the WCR in most regions, but some populations have apparently lost their ovipositional fidelity to cornfields, and lay eggs in soybean and other crops in addition to maize<sup>18–20</sup>. Beginning in 2003, transgenic maize carrying a gene from the entomopathogenic bacterium *Bacillus thuringiensis* Berliner (Bt) has been effective in controlling the WCR and northern corn rootworms (*D. barberi*). Yet, certain WCR populations have since evolved resistance to some Bt toxins<sup>21</sup>. This ability of WCR to rapidly evolve resistance has significantly reduced the efficacy of these management strategies, at least in certain areas.

Kuhlmann & Van der Burgt<sup>22</sup> recommended biological control as an option for Europe, where genetically modified plants are mostly banned and the use of additional insecticides is not desirable. Classical biological control would involve the importation and the establishment of natural enemies from the WCR area of origin in North America. A more readily available option would be an inundative biological control approach with commercially available native antagonists, such as entomopathogenic nematodes (EPN)<sup>22</sup>.

Soil-dwelling EPN have been successfully used as biological control agents against a range of different insect pests, including WCR<sup>23–25</sup>. EPN are favored because they are harmless to vertebrates, commercially available, and authorized in many countries<sup>26–31</sup>. EPN in the families Steinernematidae and Heterorhabditidae carry mutualistic bacteria of the genera *Xenorhabdus* and *Photorhabdus*, respectively, and together function as obligate parasites of insects<sup>32,33</sup>. The free-living stage of EPN, known as the infective juvenile (IJ), is adapted to persist in the soil where it searches for a suitable insect host<sup>34</sup>. Upon contact with a host, it enters the insect's hemocoel through natural openings and releases their symbiont bacteria. Within 2–3 days, the insect host dies of septicemia caused by the proliferating bacteria. The EPN consume the bacteria and reproduce to form two to three generations, until the resources in the cadaver are depleted. Non-feeding infective stages then emerge and may survive in the surrounding soil for several months in search of a new host<sup>35</sup>.

Various other soil organisms also have the potential to improve plant performance by, for instance, promoting growth, facilitating nutrient acquisition, stimulating defenses, and protecting plants from pathogens and pests<sup>36–38</sup>. Among these are arbuscular mycorrhizal fungi, which colonize roots of many terrestrial plants and can provide these plants with nutrients in exchange for photosynthetic by-products<sup>39,40</sup>. Arbuscular mycorrhizal fungi have also been shown to increase plant tolerance to a variety of stresses, both biotic and abiotic<sup>40</sup>. Some arbuscular mycorrhizal fungi such as *Rhizoglyphus irregularis* are commercialized as inoculates for seedlings or as seed coatings, in order to improve soil fertility and plant performance<sup>41–46</sup>.

Similarly, growth promoting rhizobacteria within the *Pseudomonas fluorescens* group, such as *Pseudomonas protegens* and *Pseudomonas chlororaphis*, have been shown to trigger systemic resistance in colonized plants, and may control soil-borne pathogens with potent antifungal compounds<sup>47–51</sup>. *Pseudomonas protegens* and *Pseudomonas chlororaphis* strains also have insecticidal activity and are particularly effective against Lepidopteran pests<sup>52–55</sup>. Currently there are several products based on plant-beneficial pseudomonads that are commercialized, primarily in the USA<sup>52,56–58</sup>.

A previous study<sup>59</sup> showed that the combined application of the EPN *Heterorhabditis bacteriophora* and the rhizobacteria *Pseudomonas protegens* CHA0 and *Pseudomonas chlororaphis* PCL1391 improved the performance and protection of wheat. This was most evident during a season that the plants were infested by frit fly larvae<sup>59</sup>.

In the current study, we evaluated the singular application of three beneficial soil organisms on maize performance under WCR infestation. Treatments with EPN (*Steinernema feltiae* and *H. bacteriophora*), *Pseudomonas* bacteria, and a commercial formulation of arbuscular mycorrhizal fungi, as well as a treatment with the combination of all three beneficial organisms were applied under realistic field conditions.

## Materials and Methods

**The beneficial soil organisms' origins and formulations.** Strains of *Pseudomonas protegens* Pf-5<sup>60,61</sup> and *Pseudomonas chlororaphis* O6<sup>62</sup> with a spontaneous resistance to the antibiotic rifampicin were used in this study in 2015 (Table 1). In 2016 and 2017 we used two closely related bacterial strains, *Pseudomonas protegens* CHA0<sup>63</sup>, and *Pseudomonas chlororaphis* PCL1391<sup>64</sup> that have been similarly selected for spontaneous resistance to rifampicin following previously described protocols<sup>59,65</sup> (Table 1). To prepare the bacterial inoculum for field application, the strains were grown overnight at 25 °C in LB Broth Miller (Fisher BioReagents) containing 100 µg/ml of rifampicin. Aliquots of 200 µl of each culture were then plated on LB Agar Miller (Fisher BioReagents) without antibiotics. After incubation at 27 °C for 16 h, bacterial cells were harvested and washed in sterile distilled water. The optical density at 600 nm (OD<sub>600</sub>) of the bacterial cell suspensions was adjusted to 0.15 corresponding to a cell density of about 8 × 10<sup>7</sup> CFU/ml. To preserve the bacterial concentrations chosen for application to the field, the bacterial stock suspensions were maintained on ice until final dilution and use.

Entomopathogenic nematodes (EPN) of the species *Steinernema feltiae* and *Heterorhabditis bacteriophora* were provided by the company Koppert Biological Systems (<https://www.koppert.com>, Table 1). EPN were received in vermiculite powders around two weeks before their application to the field. One or two days before field application, IJs concentration was assessed and the powder containing nematodes weighted to reach a concentration of 0.65 Mio of IJs of each species and placed in a 50 ml sterile conical tube (USA Scientific) (Table 1). Tubes containing the IJs were kept at ~5 °C prior to field application.

Beneficial group/species	Strain	Application type	GenBank accession no.	Reference or source
<b>Arbuscular mycorrhizal fungi</b>				
<i>Rhizophagus irregularis</i> <sup>a</sup>		Substrate	n.a. <sup>d</sup>	Mycorrhizal Fungi Products Sarasota, Florida
<i>Funnelformis mosseae</i> <sup>a</sup>		Substrate	n.a.	Mycorrhizal Fungi Products Sarasota, Florida
<i>Septogloium deserticola</i> <sup>a</sup>		Substrate	n.a.	Mycorrhizal Fungi Products Sarasota, Florida
<i>Claroideogloium claroideum</i> <sup>a</sup>		Substrate	n.a.	Mycorrhizal Fungi Products Sarasota, Florida
<i>Claroideogloium etunicatum</i> <sup>a</sup>		Substrate	n.a.	Mycorrhizal Fungi Products Sarasota, Florida
<i>Rhizogloium microaggregatum</i> <sup>a</sup>		Substrate	n.a.	Mycorrhizal Fungi Products Sarasota, Florida
<i>Rhizogloium clarum</i> <sup>a</sup>		Substrate	n.a.	Mycorrhizal Fungi Products Sarasota, Florida
<b>Entomopathogenic nematodes</b>				
<i>Heterorhabditis bacteriophora</i> <sup>b</sup>		Aqueous	KJ938576	Koppert biological systems
<i>Steinernema feltiae</i> <sup>b</sup>		Aqueous	KJ938569	Koppert biological systems
<b><i>Pseudomonas</i> bacteria</b>				
<i>Pseudomonas chlororaphis</i>	PCL1391 <sup>c</sup>	Aqueous	NZ_LFUT01000004	Chin-A-Woeng <i>et al.</i> <sup>64</sup> ; Flury <i>et al.</i> <sup>54</sup>
<i>Pseudomonas protegens</i>	CHA0 <sup>c</sup>	Aqueous	NC_021237	Stutz <i>et al.</i> <sup>63</sup> ; Flury <i>et al.</i> <sup>54</sup>
<i>Pseudomonas chlororaphis</i>	O6 <sup>c</sup>	Aqueous	NZ_CM001490.1	Loper, <i>et al.</i> <sup>93</sup>
<i>Pseudomonas protegens</i>	Pf-5 <sup>c</sup>	Aqueous	NC_004129.6	Loper, <i>et al.</i> <sup>93</sup>

**Table 1.** Beneficial soil organisms applied individually or in combinations in the field experiments. <sup>a</sup>A commercialized treatment (Ecovam™ Vamendo Granular) containing seven species of arbuscular mycorrhiza was used as inoculant in the 2015, 2016 and 2017 field trials. <sup>b</sup>A mixture of the entomopathogenic nematodes *H. bacteriophora* and *S. feltiae* was used in the 2015, 2016 and 2017 field trials. <sup>c</sup>Rifampicin-resistant variants of strains O6 and Pf-5 were used as inoculants in the 2015 field trial, while strains CHA0 and PCL1391 were used as inoculants in the 2016 and 2017 field trials. <sup>d</sup>n.a., not available.

Arbuscular mycorrhizal fungi (AMF) were provided by Evocam™ (<https://horticulturalalliance.com/product/ecovam-vam-endo-granular/>) that contains seven species of arbuscular mycorrhizal fungi, belonging to the genera *Rhizophagus*, *Funnelformis*, *Septogloium*, *Claroideogloium* and *Rhizogloium* (Table 1). The product richness was estimated to 150 spores per gram of substrate. Moreover, a “mock” inoculum, which consisted of the substrate without arbuscular mycorrhizal fungi spores was prepared by autoclaving the original arbuscular mycorrhizal fungi inoculum for 2 h at 110–120 °C, two weeks before field application. Bags containing the inoculum and the “mock” inoculum were stored at room temperature prior to field application.

**Field experiments.** Field experiments were conducted during three consecutive springs, in 2015, 2016 and 2017, at the Bradford Research and Extension Centre (38.8929376 N, –92.2009539 W, Columbia, MO, USA). The soil type at this location is a Mexico silt loam made up of 12.5% sand, 65% silt, and 22.5% clay as determined by the University of Missouri Soil Testing Facility, Columbia, MO.

In plots of 1.5 m we planted a row with 8 seeds of the maize cultivar Pioneer 33T55. Each of these experimental rows was separated with a buffer row of the same size planted with the same maize cultivar. Row spacing was 0.76 m, hence, rows were separated from each other by 0.76 m. Experimental plots were hand planted in May of each year. The treatments applied to the field were: (1) EPN suspension, (2) plant-growth promoting rhizobacteria (PGPR) suspension, (3) AMF inoculum, (4) a combination of the EPN, AMF and PGPR, (5) AMF “mock” inoculum, and (6) control (no application). Each year, the experiment was conducted in different fields on the same experimental farm.

Bacterial cell suspensions were applied directly on the maize seeds after they were placed in the furrows using treatment-specific watering cans. Concentrated bacterial stock suspensions (OD<sub>600</sub> 0.15; corresponding to ~8 × 10<sup>7</sup> CFU/ml) were diluted in ca. 5 L of water for each plot directly at the field site before soil inoculation. In the field trial performed in 2015, the bacterial inoculum was a mixture of *P. protegens* Pf-5 and *P. chlororaphis* O6, while in 2016 and 2017 the chosen strains were *P. protegens* CHA0 and *P. chlororaphis* PCL1391.

For EPN application (*S. feltiae* and *H. bacteriophora*) the nematodes that had been stored in 50 ml sterile tubes were mixed in treatment-specific watering cans in which water was added to a final volume of ca. 5 L per plot and applied in the furrows at a final concentration of 1.3 × 10<sup>6</sup> IJs/m<sup>2</sup>.

Finally, 400 ml of substrate per plot, containing approximately 4.8 × 10<sup>7</sup> AMF spores were evenly applied on the seeds using a 500 ml glass beaker. AMF-control plots were inoculated with the same amount of substrate without AMF propagules. Control plots were treated with the same volume of water without the beneficial organisms. After treatments, the seeds were immediately covered with soil by closing the seed furrows. All material which entered into contact with the different inoculants was cleaned and disinfested with 70% ethanol.

When plants were at the two-leaf stage, half of experimental plots were artificially infested with WCR eggs as previously described in El Khishen *et al.*<sup>66</sup>. The WCR eggs were obtained from the USDA-ARS facility in Brookings (SD, USA). We used their primary diapausing strain, which was maintained at ~8 °C until application. The eggs were applied when the plants reached the V2 stage as described above. Eggs were mixed into a solution of water containing agar at the final concentration of 0.15%, and each plant was exposed to ~800 viable eggs delivered evenly down both sides of the row with a tractor-mounted system. The number of replicates for

each treatment in the experiment carried out in 2015 was 8, for a total of 96 experimental plots, while in field experiments performed in 2016 and 2017 the experiment was doubled to facilitate data collection, for a total of 192 experimental plots (Supplementary Material 1). All replicates were arranged in a randomized complete block design with 8 blocks (each containing 12 treatments in a split-plot design WCR vs no WCR) for the field trial 2015, and with 16 blocks for the experiments carried out the following two years. The 16 blocks in 2016 and 2017 still resulted in 8 replications because half were used for damage plus larval recovery and half for yield.

**Evaluation of WCR damage severity and maize yield.** About six weeks after the WCR eggs were applied, approximately 500 degree-days post infestation as calculated with the techniques of Hibbard *et al.*<sup>67</sup>, root damage was evaluated on three plants per plot. Maize plants were dug out from the soil, their roots were washed, and damage caused by WCR larval feeding was rated using the node injury score<sup>5</sup>.

We also evaluated the presence of WCR larvae on the roots. For this, two additional plants were removed from each plot at approximately 410 degree days<sup>67</sup> post infestation, when most larvae should be at the early third instar. Following Hibbard *et al.*<sup>68</sup>, the entire root system of each collected plant was placed into onion bags and the bags were suspended in a greenhouse (38–50 °C). A water pan was positioned under each bag to collect all larvae that fell down. Larvae were collected and counted until no additional larvae were recovered for three consecutive days. To estimate the impact of the different treatments on the WCR fitness, collected larvae were counted and weighed.

At the end of the season, maize cobs from the three remaining plants per plot (2015) or from the yield portion of the study were harvested and grain yield was determined and expressed in total grain weight.

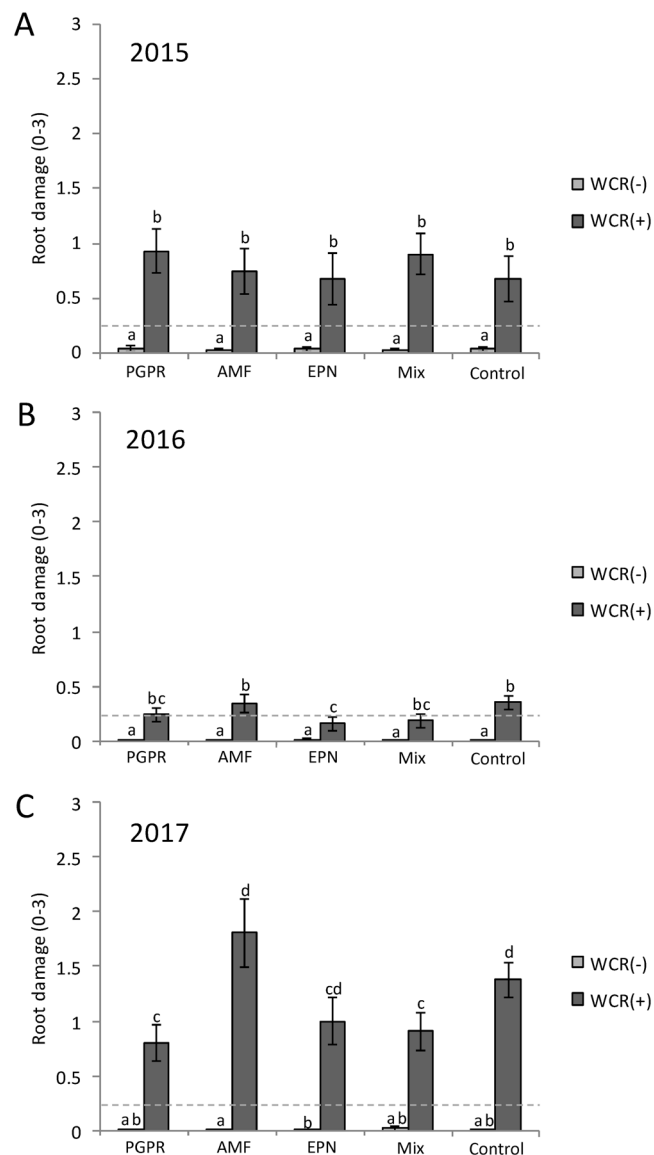
**Monitoring of beneficial organisms.** *Pseudomonas bacteria.* In 2017, we monitored the presence of the *Pseudomonas* strains in the different plots. Maize roots were sampled about 5 weeks after the application of WCR eggs. For this, the root systems from six maize plants (i.e. two plants taken from three extra-plots specifically planted to assess *Pseudomonas* survival during the field experiment) were dug up, pooled, washed and gently dried using paper towels. To avoid cross-contamination between samples, all material used for the sampling at the field site was cleaned with 70% ethanol. Roots were placed in 15 ml sterile conical screw cap centrifuge tubes (Basix) containing 40 ml of sterile water and vigorously agitated on a rotary shaker at 180 rpm for 15–20 min. Subsequently roots were removed from the tubes, dried at 80 °C for three days and weighed. The remaining suspensions were transferred to fresh sterile tubes on ice and centrifuged at 8500 rpm (9300 g) at 4 °C. The obtained pellet was re-suspended in 1 ml of sterile water. Each sample was then serially diluted and dilutions plated on LB Agar Miller containing 100 µg/ml of cycloheximide (Sigma-Aldrich) and 100 µg/ml of rifampicin<sup>69</sup>. The colonies were counted and the results were expressed as colony forming units (CFU) per gram of dry root weight.

*Entomopathogenic nematodes.* In 2017, soil samples were taken from each of the plots inoculated with the EPN mix. Approximately 2,000 cm<sup>3</sup> of soil was sampled from the plots near the plants by taking multiple scoops approximately 12 cm deep into the soil. Individual plot samples were mixed and two subsamples of approximately 120 ml were placed into 236 ml plastic containers (Solo Cup Company, Lake Forest, IL, USA) and baited with two last-instar *Galleria mellonella* L. (Lepidoptera: Pyralidae) larvae each. Samples were maintained in the dark at 20 °C and checked daily for *G. mellonella* mortality. If cadavers were found with nematodes present as typical EPN symptom<sup>70</sup>, the plots tested were recorded as having an active nematode population.

**Statistical analysis.** All statistical analyses were performed using the software package R<sup>71</sup>, version 3.2.3. Data were checked for normal distribution with the Shapiro-Wilk test and by plotting QQ-Plots. Equality of variance was verified performing Bartlett's test. Most of the data failed the normality and equality of variance assumptions, therefore non-parametric Kruskal-Wallis analysis of variance on ranks (H-tests) were carried out. Post-hoc test analyses were conducted using Fisher's least significant difference with a Benjamini-Hochberg correction of *P*-values (package "agricolae")<sup>72</sup>. Results obtained in the control experiment in which the carrier substrate for the AMF was tested alone (AMF "mock" inoculum) were not significantly different from those obtained in the untreated control. Therefore, the control and AMF-control were pooled to facilitate the interpretation of the results. Moreover, the effect of the WCR infestation (infested *versus* non-infested) was so high compared to the effect of the beneficial organisms treatments, that, in order to detect differences among application treatments, the effect of the WCR infestation was assessed separately.

## Results

**Impact of beneficial soil organisms' application on maize root damage.** In 2015, all plots that were artificially infested with WCR showed significantly more damage than non-infested plots, revealing the efficiency of the infestation (Chisq = 53.65, *P* < 0.001). Average damage was 0.78 on the Oleson scale for infested plots (Fig. 1A). The application of the beneficial soil organisms did not reduce the damage caused by WCR larvae to maize roots (Fig. 1A). For the 2016 field trial, average root damage in the infested plots was 0.25 on the Oleson scale, which was again significantly greater than in non-infested plots (Chisq = 63.8, *P* < 0.001) (Fig. 1B). Maize roots from plots treated with PGPR, EPN and the combination of PGPR, EPN and AMF (Mix), were slightly less damaged compared to untreated plants, although the observed difference was statistically significant only for the EPN treatment, which showed significantly less root damage than the control plots (*P* = 0.03). Maize roots were most damaged in plots treated with AMF, and data were not significantly different compared to the control plots (*P* = 0.88) (Fig. 1B). In 2017, the average value of root damage was 1.18, the highest observed over the three consecutive years. Once again, all plots infested with WCR showed significantly more damage than non-infested plots (Chisq = 73.1, *P* < 0.001). Maize roots from plots treated with PGPR, EPN and the Mix were slightly less damaged as compared to untreated plants. The observed differences were statistically significant for the PGPR and the Mix treatment as compared to the AMF and the control plots (PGPR: AMF, *P* = 0.01; PGPR: control, *P* = 0.02; Mix: AMF, *P* = 0.02; Mix: control, *P* = 0.05) (Fig. 1C).

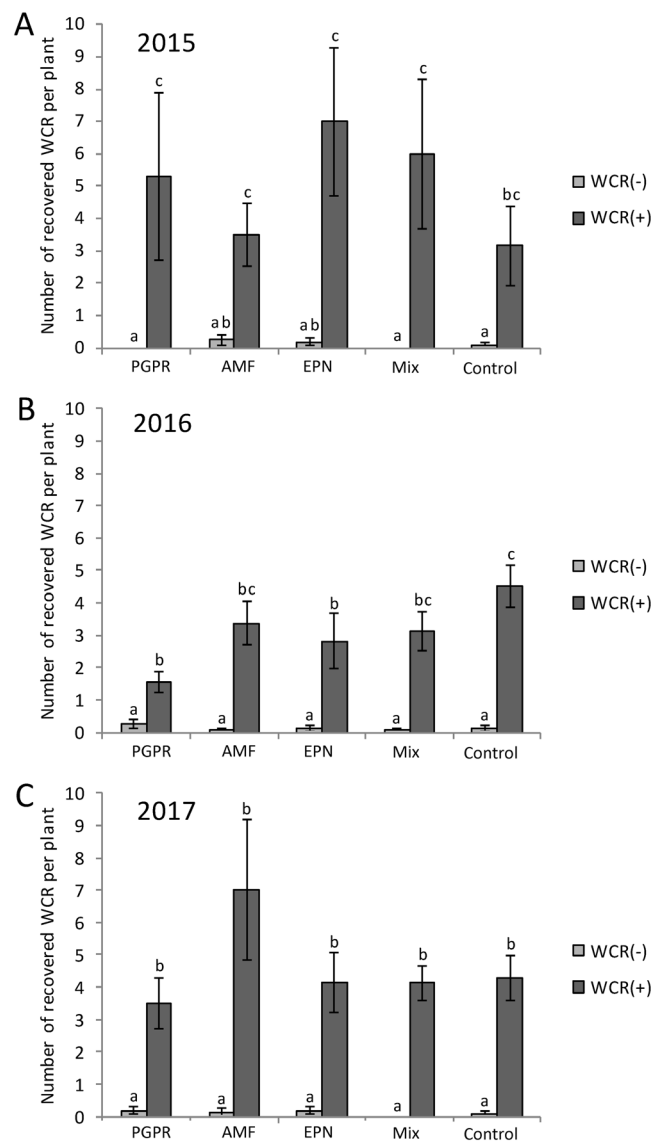


**Figure 1.** Root damage measured on the node-injury scale (Oleson *et al.*<sup>5</sup>) depending on the beneficial organisms applied and the western corn root worm (WCR) infestation status. (A) in 2015, (B) in 2016 and (C) in 2017. The dash line represents the economical threshold of root damage. PGPR: plant-growth promoting rhizobacteria, EPN: entomopathogenic nematodes, AMF: arbuscular mycorrhizial fungi, Mix: PGPR + EPN + AMF. Bars represent mean percentage  $\pm$  SE. Means denoted by different letters are significantly different ( $P < 0.05$ , Fisher's least significant difference test).

**Impact of treatments on WCR survival and weight.** Almost no WCR larvae were recovered from non-infested plots as compared to infested plots (2015:  $\text{Chisq} = 34.3$ ,  $P < 0.001$ ; 2016:  $\text{Chisq} = 68.4$ ,  $P < 0.001$ ; 2017:  $\text{Chisq} = 76.1$ ,  $P < 0.001$ ) (Fig. 2). In 2015, the number of recovered WCR larvae was not affected by the various soil applications, but tended to be slightly higher compared to the control plots (Fig. 2). In 2016, however, the number of recovered WCR larvae was significantly lower in plots with PGPR application and in plots with EPN application, but not in plots with the mixture containing the two plus AMF (PGPR: control,  $P = 0.003$ ; EPN: control,  $P = 0.03$ ) (Fig. 2B). In 2017, the beneficial soil organisms did not negatively affect the number of recovered WCR larvae (Fig. 2C).

In 2015, WCR larvae weight was not affected by the applications (Fig. 3A). In 2016, WCR larvae in PGPR plots weight significantly less compared to those recovered from control plots (PGPR: control,  $P = 0.02$ ), while no significant differences were observed for the other treatments (Fig. 3B). In 2017, no differences in larval weight were observed among the different treatment plots (Fig. 3C).

**Impact of treatments on yield.** The infestation with WCR did not have any impact on yield (expressed as gram of seed per plot) in any of the field trials (2015:  $\text{Chisq} = 0.01$ ,  $P = 0.9$ ; 2016:  $\text{Chisq} = 0.1$ ,  $P = 0.7$ ; 2017:  $\text{Chisq} = 0.1$ ,  $P = 0.7$ ) (Fig. 4). However, in 2015, yield was positively impacted by PGPR and EPN applications



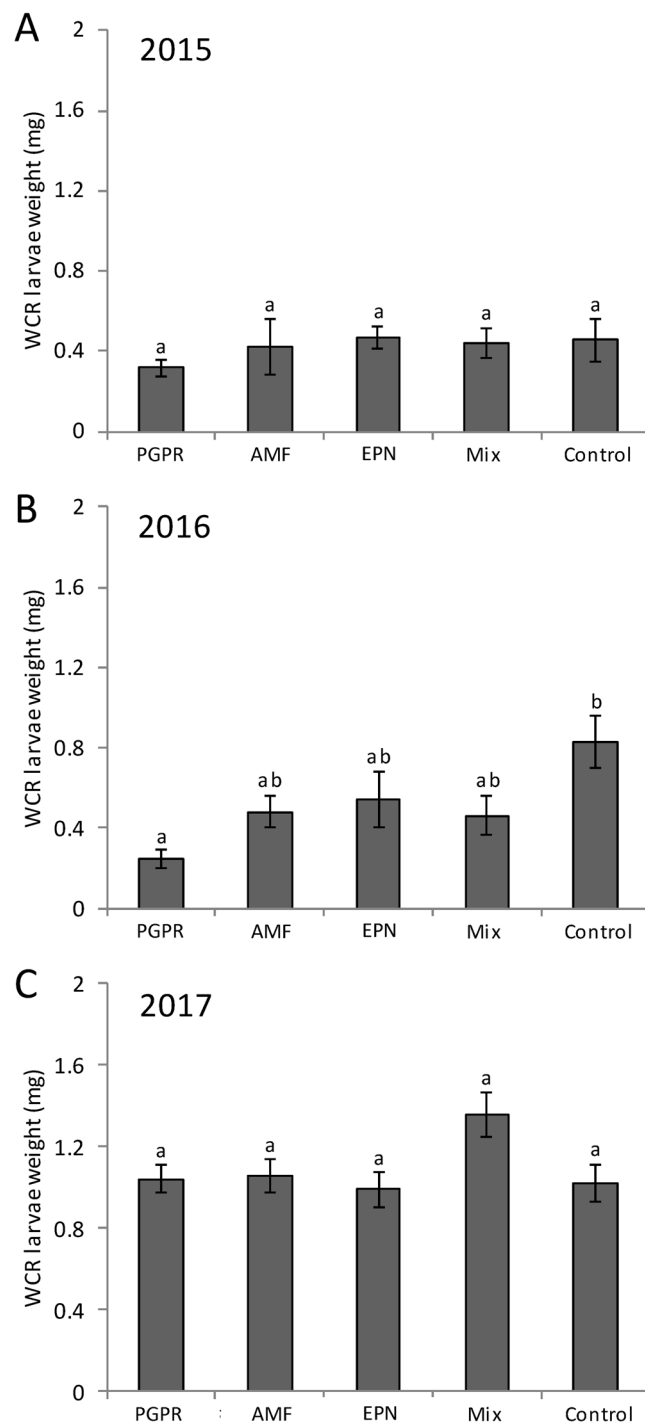
**Figure 2.** Number of western corn root worm larvae recovered from root system depending on the beneficial organisms applied and the western corn root worm (WCR) infestation status. (A) In 2015, (B) in 2016 and (C) in 2017. PGPR: plant-growth promoting rhizobacteria, EPN: entomopathogenic nematodes, AMF: arbuscular mycorrhizal fungi, Mix: PGPR + EPN + AMF. Bars represent mean percentage  $\pm$  SE. Means denoted by different letters are significantly different ( $P < 0.05$ , Fisher's least significant difference test).

(independently of the WCR infestation) (PGPR: control,  $P = 0.01$ , EPN: control,  $P = 0.01$ ). In 2016 and 2017, the beneficial soil organisms had no impact on yield (Fig. 4B,C).

**Persistence of the applied rhizobacteria and nematodes.** In 2017, nematodes were found in all of the corresponding augmented plots. In plots that had been treated with rhizobacteria, the numbers of rifampicin-resistant *Pseudomonas* varied between  $1.05 \times 10^3$  and  $3.81 \times 10^5$  CFU.g<sup>-1</sup> of dry root weight. No rifampicin-resistant bacteria were found in control plots.

## Discussion

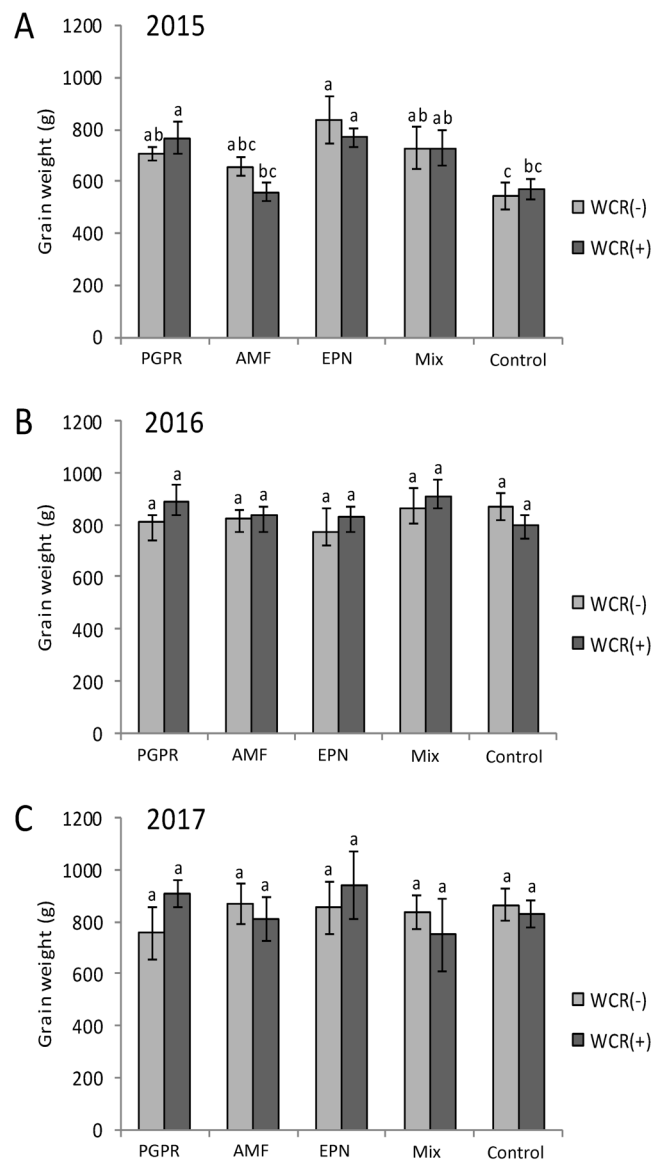
Overall, our results confirm that PGPR and EPN can protect maize roots from WCR, as observed through a reduction in root damage in plots where they were applied separately or in combination with AMF. In 2016, in plots treated with EPNs, root damage was reduced below the economic threshold. This was expected because both EPN species used in this study (*S. feltiae* and *H. bacteriophora*) are known to readily kill WCR<sup>23–26,73–75</sup>, and in a previous study on the same experimental farm, root damage by WCR was reduced by the application of a slightly lower dose of *H. bacteriophora* (50 IJs/cm<sup>2</sup>)<sup>76</sup>. From our 2017 trial, we can conclude that PGPR application can also significantly reduce WCR-inflicted root damage. For the PGPR, the observed reduction of root feeding may be explained by induced systemic resistance<sup>77,78</sup>, as well as by direct insecticidal effects. Ours is the first field trial



**Figure 3.** Western corn rootworm weight in response to the application of beneficial organisms. (A) In 2015, (B) in 2016 and (C) in 2017. PGPR: plant-growth promoting rhizobacteria, EPN: entomopathogenic nematodes, AMF: arbuscular mycorrhizial fungi, Mix: PGPR + EPN + AMF. Bars represent mean percentage  $\pm$  SE. Means denoted by different letters are significantly different ( $P < 0.05$ , Fisher's least significant difference test).

to test if *Pseudomonas* strains application can reduce WCR pressures. Yet, from a study using transgenic maize plants expressing an insecticidal protein that is naturally produced by a *P. chlororaphis* isolate, it is known that it strongly affects WCR feeding and survival<sup>79</sup>.

PGPR application in the 2016 trial was the only treatment that reduced WCR weight, possibly explained by enhanced plant defense or increased infection of the insects. In 2016 and 2017, the yield was not affected by WCR infestation, nor by any of the applications of soil organisms. Apparently, despite significant damage to the root system, the plants were able to somehow compensate and still be fully productive<sup>80</sup>. Although PGPR and EPN



**Figure 4.** Yield, expressed as maize grain weight, in response to the application of beneficial organisms and the western corn root worm (WCR) infestation status. (A) in 2015, (B) in 2016 and (C) in 2017. PGPR: plant-growth promoting rhizobacteria, EPN: entomopathogenic nematodes, AMF: arbuscular mycorrhizal fungi, Mix: PGPR + EPN + AMF. Bars represent mean percentage  $\pm$  SE. Means denoted by different letters are significantly different ( $P < 0.05$ , Fisher's least significant difference test).

did not have any detectable impact on WCR in 2015, this was the only year where yield was increased following their application. Impact of WCR was minimal during 2015, probably due to waterlogging of the plots. We speculate that the positive impact of EPN and PGPR application may have been the result of induced resistance against pathogens and growth promotion, which are known properties of these organisms<sup>77,78,81,82</sup>. For instance, *Pseudomonas* spp. produce antimicrobial compounds that can stimulate systemic resistance in plants<sup>82</sup>, or act as growth promoters or inhibitors and increase stress tolerance<sup>83</sup>. EPN have also been shown to induce such resistance in plants, but the mechanisms that are involved remain to be elucidated<sup>81</sup>. Also, phytohormones like auxin, cytokinin, gibberellin or ethylene of microbial and fungal origin can affect growth, root development, immune response and hormonal pathways in plants<sup>38,51</sup>. PGPR are also involved in the solubilization of mineral phosphates and other nutrients that can facilitate their access by the plant<sup>84,85</sup>. We should stress that the bacterial strains used in 2015 (i.e., *P. protegens* Pf-5 and *P. chlororaphis* O6) were not the same as those used in 2016 and 2017 (i.e., *P. protegens* CHA0 and *P. chlororaphis* PCL1391) and, although they were very similar, possible differences between strains may have had an impact on their effectiveness. All four strains possess the cluster *fit* that directs the synthesis of FitD, the insecticidal protein that enables these bacteria to kill different insect pests<sup>53,54</sup>.

The application of AMF, which were included to confirm their compatibility with the nematodes and bacteria, did not significantly affect WCR survival and performance, nor did it affect plant performance. AMF were applied before the development of the first roots because the spores can persist for a long time in the soil until



they form hyphae to colonize roots<sup>86</sup>. Unfortunately, for practical reasons, we could not assess the establishment success of the AMF, but the same commercial inoculum is commonly used to enhance crop performance and has been shown to successfully colonize rice roots<sup>87</sup>. It is, however, possible that our inoculant was not effective and failed to persist in the soil and to colonize roots, as was reported for another trial with similar AMF species for arable maize<sup>88</sup>. From a Swiss study in wheat plots with a different AMF inoculum we know that after application at the seedling stage, it has the potential to persist and can successfully colonize the roots<sup>59</sup>. Although we can safely conclude that the applied AMF had no effect on the efficacy of the PGPR and EPN, for future studies, it would be desirable to gather information on the inoculant fate over the field seasons to confirm persistence and colonization status.

We observed no synergistic or additive effect of the soil organisms as their combined application did not result in a higher efficacy to protect maize roots from damage or increase yield. Therefore, it is not excluded that interactions between the applied soil organisms in some ways limit their full potential. The weather, and therefore the field conditions, were very different from one year to another (Supplementary Material 2), in particular in terms of precipitation, which surely affected the results. To compare the weather over the entire period of each trial, we took into account several parameters from 1<sup>st</sup> May to 1<sup>st</sup> November, for each year. Rainfall, with 625 mm, was intermediate for 2015, whereas 2016 was wetter (700 mm) and 2017 much drier (506 mm). Temperature was quite stable with an average of 20.5 °C, 21.1 °C, and 19.9 °C for 2015, 2016 and 2017, respectively. Based on these values we can assume that soil moisture was quite different among years and this must have had a significant impact on the organisms that we applied to the field. For EPNs, soil moisture determines the thickness of the water film that the nematodes need to move and survive, and it also affects the surface tension and the amount of oxygen present in the soil. These parameters influence the efficacy and survival of EPN<sup>89</sup> and could explain why EPN application was most effective in 2016, the year with the highest precipitation rate. Soil moisture is also one of the best predictors of soil microbial biomass: wet soils normally contain a greater bacterial biomass than dry soils<sup>90</sup>. Yet, Burr *et al.*<sup>91</sup> found that specific strains of *P. fluorescens* and *P. putida* were able to persist under field conditions for many weeks when the soil was “relatively dry”. Soil moisture was also a key factor in a field study on stress tolerance of *P. protegens* Pf-5<sup>59</sup>. Interestingly, the level of irrigation can have contrasting effects on the abundance of different *Pseudomonas* strains. For instance, strains that produce the antimicrobial compounds 2,4-diacetylphloroglucinol are more present in irrigated soils, whereas phenazines producers are more abundant in drier terrains<sup>92</sup>. Soil moisture effects on the persistence and performance of the strains used in our study have not yet been specifically tested.

In this study we chose to apply the soil organisms at seeding as a strategy to reduce the field work-load, using a single event for seeding and the application of the biocontrol agents. The infestation with WCR eggs occurred about two weeks after the application of the soil organisms and the WCR larvae started feeding on the roots about three weeks after application. Therefore, the soil organisms must persist in the soil for this period of time to have an effect. Applying the soil organisms after WCR infestation can be expected to be more effective in controlling the pest, but would be much more labor intensive.

## Conclusion

As is often the case with field studies, the results were quite different for the different years. Yet, each year at least one of the treatments was significantly better compared to the control (Figs 1–3). Depending on the year, the treatments had a direct impact on corn plant performance, but also impacted the survival and performance of the WCR larvae. The results obtained in 2016 were particularly encouraging, although the effect on WCR was not as evident as the two other two years. The fact that the yield was not been significantly compromised by the artificial WCR infestation explains at least partially why we found no significant impact on plant productivity, except for 2015. During years 2016 and 2017, the level of root damage proved to be a suitable parameter to measure treatment effects. We think that studies such as this one can be the basis for the development of effective soil treatments that can replace the use of pesticides, and provide a more sustainable control of WCR and other soil pests.

## Data Availability

The datasets generated during and/or analyzed during the current study are provided in the supplementary materials.

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## Author Contributions

Geoffrey Jaffuel and Nicola Imperiali performed the field experiment and wrote the manuscript. Kent Shelby helps to prepare the field experiment and performed the bacterial isolation. Raquel Campos-Herrera, Christoph Keel, Monika Maurhofer, Ted. C.J. Turlings and Bruce E. Hibbard designed the experimental set up. Christoph Keel, Ted. C.J. Turlings and Bruce E. Hibbard provided funds and materials necessary to perform the field experiment. Ryan Geisert performed the nematode isolation and help performing the field experiment. Joyce Loper provided the bacterial strain in 2015. All authors contributed to writing the manuscript.

## Additional Information

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# Protecting maize from rootworm damage with the combined application of arbuscular mycorrhizal fungi, *Pseudomonas* bacteria and entomopathogenic nematodes

Nicola Imperiali<sup>+1</sup>, Geoffrey Jaffuel<sup>+2</sup>, Kent Shelby<sup>3</sup>, Raquel Campos-Herrera<sup>2,4</sup>, Ryan Geisert<sup>3</sup>, Monika Maurhofer<sup>5</sup>, Joyce Loper<sup>6,7</sup>, Christoph Keel<sup>1</sup>, Ted. C.J Turlings<sup>2</sup>, Bruce E. Hibbard<sup>8</sup>

<sup>1</sup>Department of Fundamental Microbiology, University of Lausanne, Lausanne, Switzerland

<sup>2</sup>FARCE Laboratory, Institute of Biology, University of Neuchâtel, Neuchâtel, Switzerland

<sup>3</sup>Biological Control of Insects Research, US Department of Agriculture, Agricultural Research Service, Columbia, MO, USA

<sup>4</sup>Instituto de Ciencias de la Vid y del Vino, CSIC-Universidad de La Rioja-Gobierno de La Rioja, Logroño, Spain

<sup>5</sup>Institute of Integrative Biology, ETH Zurich, Zurich, Switzerland

<sup>6</sup>Department of Botany and Plant Pathology, Oregon State University, Corvallis, OR, USA

<sup>7</sup>Horticultural Crops Research Laboratory, US Department of Agriculture, Agricultural Research Service, Corvallis, OR, USA

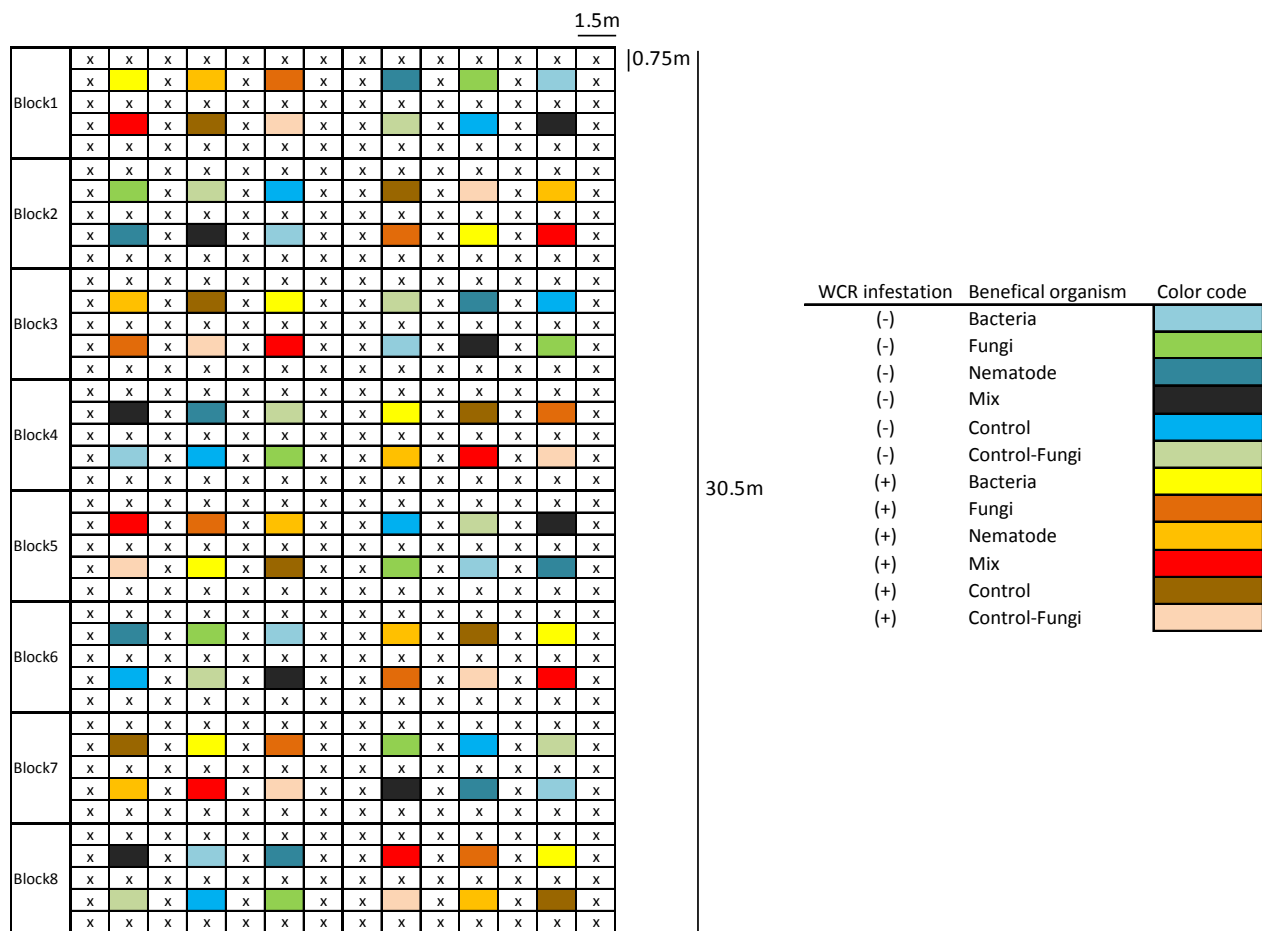
<sup>8</sup>Plant Genetics Research Unit, US Department of Agriculture-ARS, University of Missouri, Columbia, MO, USA

<sup>+</sup>These authors contributed equally to the work

Correspondence: T. Turlings & B. Hibbard; email: [ted.turlings@unine.ch](mailto:ted.turlings@unine.ch) & [Bruce.Hibbard@ars.usda.gov](mailto:Bruce.Hibbard@ars.usda.gov)

# Supplementary material 1:

## Field plot design



Experimental design for the field trial conducted in 2015, resulting in 8 blocks of 12 plots each, for a total of 96 plots. In 2016 and 2017, the size of the experiments was doubled, with 16 blocks of 12 plots each, for a total of 192 plots.

## Supplementary Material 2

Field trials (year)	2015	2016	2017
<b>Parameters recorded from 1<sup>st</sup> May to 1<sup>st</sup> November</b>			
Total precipitation (mm)	625.11	700.24	506.22
Average maximum air temperature (°C)	26.2	27.0	26.1
Average minimum air temperature (°C)	15.3	15.7	14.0
Average air temperature (°C)	20.5	21.1	19.9
Total solar radiation (MJ / m <sup>2</sup> )	17.41	17.97	18.81
Vapor pressure (kPa)	1.884	1.958	1.749
Average maximum wind speed (m/s)	8.6	8.2	8.8

**Main physical parameters recorded between the months of May and November in 2015, 2016 and 2017, at the Bradford Research and Extension Centre, Columbia, Missouri.** Data were taken from the "Missouri Historical Agricultural Weather Database" platform. (<http://agebb.missouri.edu/weather/history>).

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<sup>3</sup>Biological Control of Insects Research, US Department of Agriculture, Agricultural Research Service, Columbia, MO, USA

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<sup>6</sup>Department of Botany and Plant Pathology, Oregon State University, Corvallis, OR, USA

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Field design 2015

Block	Treat-Number	WCR	Beneficial	Missouri Plot Number
1	7	YES	Bacteria	1
1	9	YES	Nematode	2
1	8	YES	Fungi	3
1	3	NO	Nematode	4
1	2	NO	Fungi	5
1	1	NO	Bacteria	6
1	10	YES	Mix	7
1	11	YES	Control	8
1	12	YES	Control-Fungi	9
1	6	NO	Control-Fungi	10
1	5	NO	Control	11
1	4	NO	Mix	12
<hr/>				
2	2	NO	Fungi	13
2	6	NO	Control-Fungi	14
2	5	NO	Control	15
2	11	YES	Control	16
2	12	YES	Control-Fungi	17
2	9	YES	Nematode	18
2	3	NO	Nematode	19
2	4	NO	Mix	20
2	1	NO	Bacteria	21
2	8	YES	Fungi	22
2	7	YES	Bacteria	23
2	10	YES	Mix	24
<hr/>				
3	9	YES	Nematode	25
3	11	YES	Control	26
3	7	YES	Bacteria	27
3	6	NO	Control-Fungi	28
3	3	NO	Nematode	29
3	5	NO	Control	30
3	8	YES	Fungi	31
3	12	YES	Control-Fungi	32
3	10	YES	Mix	33
3	1	NO	Bacteria	34
3	4	NO	Mix	35
3	2	NO	Fungi	36
<hr/>				
4	4	NO	Mix	37
4	3	NO	Nematode	38
4	6	NO	Control-Fungi	39
4	7	YES	Bacteria	40
4	11	YES	Control	41
4	8	YES	Fungi	42
4	1	NO	Bacteria	43
4	5	NO	Control	44
4	2	NO	Fungi	45
4	9	YES	Nematode	46
4	10	YES	Mix	47

4	12	YES	Control-Fungi	48
5	10	YES	Mix	49
5	8	YES	Fungi	50
5	9	YES	Nematode	51
5	5	NO	Control	52
5	6	NO	Control-Fungi	53
5	4	NO	Mix	54
5	12	YES	Control-Fungi	55
5	7	YES	Bacteria	56
5	11	YES	Control	57
5	2	NO	Fungi	58
5	1	NO	Bacteria	59
5	3	NO	Nematode	60
6	3	NO	Nematode	61
6	2	NO	Fungi	62
6	1	NO	Bacteria	63
6	9	YES	Nematode	64
6	11	YES	Control	65
6	7	YES	Bacteria	66
6	5	NO	Control	67
6	6	NO	Control-Fungi	68
6	4	NO	Mix	69
6	8	YES	Fungi	70
6	12	YES	Control-Fungi	71
6	10	YES	Mix	72
7	11	YES	Control	73
7	7	YES	Bacteria	74
7	8	YES	Fungi	75
7	2	NO	Fungi	76
7	5	NO	Control	77
7	6	NO	Control-Fungi	78
7	9	YES	Nematode	79
7	10	YES	Mix	80
7	12	YES	Control-Fungi	81
7	4	NO	Mix	82
7	3	NO	Nematode	83
7	1	NO	Bacteria	84
8	4	NO	Mix	85
8	1	NO	Bacteria	86
8	3	NO	Nematode	87
8	10	YES	Mix	88
8	8	YES	Fungi	89
8	7	YES	Bacteria	90
8	6	NO	Control-Fungi	91
8	5	NO	Control	92
8	2	NO	Fungi	93
8	12	YES	Control-Fungi	94
8	9	YES	Nematode	95
8	11	YES	Control	96

Root damage 2015

Plot	Beneficial	WCR	Damage (1-3)
2006	Bacteria	NO	0.005
2021	Bacteria	NO	0.025
2034	Bacteria	NO	0.0125
2043	Bacteria	NO	0.0625
2059	Bacteria	NO	0
2063	Bacteria	NO	0
2084	Bacteria	NO	0
2086	Bacteria	NO	0.1875
2005	Fungi	NO	0
2013	Fungi	NO	0.025
2036	Fungi	NO	0.0625
2045	Fungi	NO	0
2058	Fungi	NO	0
2062	Fungi	NO	0
2076	Fungi	NO	0.0675
2093	Fungi	NO	0.005
2004	Nematode	NO	0
2019	Nematode	NO	0
2029	Nematode	NO	0
2038	Nematode	NO	0.0675
2060	Nematode	NO	0
2061	Nematode	NO	0.125
2083	Nematode	NO	0.025
2087	Nematode	NO	0.0375
2012	Mix	NO	0
2020	Mix	NO	0.025
2035	Mix	NO	0
2037	Mix	NO	0.025
2054	Mix	NO	0.0625
2069	Mix	NO	0
2082	Mix	NO	0.0375
2085	Mix	NO	0.05
2011	Control	NO	0.125
2077	Control	NO	0
2015	Control	NO	0
2030	Control	NO	0
2052	Control	NO	0.005
2092	Control	NO	0.066666667
2044	Control	NO	0
2067	Control	NO	0
2010	Control-Fungi	NO	0.275
2014	Control-Fungi	NO	0
2028	Control-Fungi	NO	0.033333333
2039	Control-Fungi	NO	0.006666667
2053	Control-Fungi	NO	0
2068	Control-Fungi	NO	0.005
2078	Control-Fungi	NO	0

2091 Control-Fungi	NO	0.0025
2001 Bacteria	YES	0.6875
2023 Bacteria	YES	0.0025
2027 Bacteria	YES	0.5625
2040 Bacteria	YES	0.5
2056 Bacteria	YES	1.3125
2066 Bacteria	YES	1.5625
2074 Bacteria	YES	1.1875
2090 Bacteria	YES	1.625
2003 Fungi	YES	0.8333333333
2022 Fungi	YES	0
2031 Fungi	YES	1.6875
2042 Fungi	YES	1.125
2050 Fungi	YES	1.125
2070 Fungi	YES	0.1375
2075 Fungi	YES	0.1925
2089 Fungi	YES	0.8125
2002 Nematode	YES	0.025
2018 Nematode	YES	1.5
2025 Nematode	YES	1.416666667
2046 Nematode	YES	0.04
2051 Nematode	YES	0.075
2064 Nematode	YES	0.0875
2079 Nematode	YES	1.25
2095 Nematode	YES	1
2007 Mix	YES	0.9375
2024 Mix	YES	1.625
2033 Mix	YES	0.275
2047 Mix	YES	0.0033333333
2049 Mix	YES	0.875
2072 Mix	YES	1.125
2080 Mix	YES	0.9375
2088 Mix	YES	1.4375
2057 Control	YES	0.1375
2008 Control	YES	0.175
2065 Control	YES	1.3333333333
2041 Control	YES	0.0775
2026 Control	YES	0.255
2016 Control	YES	0.09
2073 Control	YES	0.625
2096 Control	YES	1.125
2009 Control-Fungi	YES	0.5875
2017 Control-Fungi	YES	0.0125
2032 Control-Fungi	YES	1.3125
2048 Control-Fungi	YES	1.5625
2055 Control-Fungi	YES	1.75
2071 Control-Fungi	YES	0.275
2081 Control-Fungi	YES	0.325
2094 Control-Fungi	YES	1.166666667

Larval recovery 2015

Plot	Beneficial	WCR	larval recovery (mean nb)
2001	Bacteria	YES	0.5
2023	Bacteria	YES	0
2027	Bacteria	YES	0
2040	Bacteria	YES	0
2056	Bacteria	YES	11.5
2066	Bacteria	YES	5.5
2074	Bacteria	YES	4.5
2090	Bacteria	YES	20.5
2006	Bacteria	NO	0
2021	Bacteria	NO	0
2034	Bacteria	NO	0
2043	Bacteria	NO	0
2059	Bacteria	NO	0
2063	Bacteria	NO	0
2084	Bacteria	NO	0
2086	Bacteria	NO	0
2008	Control	YES	0
2016	Control	YES	2
2026	Control	YES	8
2041	Control	YES	0
2057	Control	YES	0
2065	Control	YES	0
2073	Control	YES	4
2096	Control	YES	10.5
2011	Control	NO	0
2015	Control	NO	0
2030	Control	NO	0
2044	Control	NO	0
2052	Control	NO	1
2067	Control	NO	0.5
2077	Control	NO	0
2092	Control	NO	0
2009	Control-Fungi	YES	7
2017	Control-Fungi	YES	0
2032	Control-Fungi	YES	1
2048	Control-Fungi	YES	1.5
2055	Control-Fungi	YES	16.5
2071	Control-Fungi	YES	0
2081	Control-Fungi	YES	0
2094	Control-Fungi	YES	0
2010	Control-Fungi	NO	0
2014	Control-Fungi	NO	0
2028	Control-Fungi	NO	0
2039	Control-Fungi	NO	0
2053	Control-Fungi	NO	0
2068	Control-Fungi	NO	0
2078	Control-Fungi	NO	0

2091 Control-Fungi	NO	0
2003 Fungi	YES	1.5
2022 Fungi	YES	0
2031 Fungi	YES	5.5
2042 Fungi	YES	4
2050 Fungi	YES	7.5
2070 Fungi	YES	4
2075 Fungi	YES	0
2089 Fungi	YES	5.5
2005 Fungi	NO	0
2013 Fungi	NO	1
2036 Fungi	NO	0
2045 Fungi	NO	0
2058 Fungi	NO	0
2062 Fungi	NO	1
2076 Fungi	NO	0
2093 Fungi	NO	0
2007 Mix	YES	4.5
2024 Mix	YES	2
2033 Mix	YES	1
2047 Mix	YES	0
2049 Mix	YES	8
2072 Mix	YES	18
2080 Mix	YES	13
2088 Mix	YES	1.5
2012 Mix	NO	0
2020 Mix	NO	0
2035 Mix	NO	0
2037 Mix	NO	0
2054 Mix	NO	0
2069 Mix	NO	0
2082 Mix	NO	0
2085 Mix	NO	0
2002 Nematode	YES	0
2018 Nematode	YES	18
2025 Nematode	YES	11
2046 Nematode	YES	0
2051 Nematode	YES	8.5
2064 Nematode	YES	0.5
2079 Nematode	YES	11
2095 Nematode	YES	7
2004 Nematode	NO	0.5
2019 Nematode	NO	0
2029 Nematode	NO	0
2038 Nematode	NO	0
2060 Nematode	NO	0
2061 Nematode	NO	1
2083 Nematode	NO	0
2087 Nematode	NO	0

Larval weight 2015

Plot	Beneficial	WCR	larvae weight (mg)
2001	Bacteria	YES	1
2023	Bacteria	YES	0
2027	Bacteria	YES	0
2040	Bacteria	YES	0
2056	Bacteria	YES	11.5
2066	Bacteria	YES	5.5
2074	Bacteria	YES	4.5
2090	Bacteria	YES	20.5
2008	Control	YES	0
2016	Control	YES	4
2026	Control	YES	8
2041	Control	YES	0
2057	Control	YES	0
2065	Control	YES	0
2073	Control	YES	4
2096	Control	YES	10.5
2009	Control-Fungi	YES	14
2017	Control-Fungi	YES	0
2032	Control-Fungi	YES	1
2048	Control-Fungi	YES	1.5
2055	Control-Fungi	YES	16.5
2071	Control-Fungi	YES	0
2081	Control-Fungi	YES	0
2094	Control-Fungi	YES	0
2003	Fungi	YES	1.5
2022	Fungi	YES	0
2031	Fungi	YES	5.5
2042	Fungi	YES	4
2050	Fungi	YES	7.5
2070	Fungi	YES	4
2075	Fungi	YES	0
2089	Fungi	YES	5.5
2007	Mix	YES	4.5
2024	Mix	YES	2
2033	Mix	YES	1
2047	Mix	YES	0
2049	Mix	YES	8
2072	Mix	YES	18
2080	Mix	YES	26
2088	Mix	YES	1.5
2002	Nematode	YES	0
2018	Nematode	YES	18
2025	Nematode	YES	11
2046	Nematode	YES	0
2051	Nematode	YES	17
2064	Nematode	YES	1
2079	Nematode	YES	11

2095	Nematode	YES	7
2006	Bacteria	NO	0
2021	Bacteria	NO	0
2034	Bacteria	NO	0
2043	Bacteria	NO	0
2059	Bacteria	NO	0
2063	Bacteria	NO	0
2084	Bacteria	NO	0
2086	Bacteria	NO	0
2011	Control	NO	0
2015	Control	NO	0
2030	Control	NO	0
2044	Control	NO	0
2052	Control	NO	2
2067	Control	NO	0.5
2077	Control	NO	0
2092	Control	NO	0
2010	Control-Fungi	NO	0
2014	Control-Fungi	NO	0
2028	Control-Fungi	NO	0
2039	Control-Fungi	NO	0
2053	Control-Fungi	NO	0
2068	Control-Fungi	NO	0
2078	Control-Fungi	NO	0
2091	Control-Fungi	NO	0
2005	Fungi	NO	0
2013	Fungi	NO	2
2036	Fungi	NO	0
2045	Fungi	NO	0
2058	Fungi	NO	0
2062	Fungi	NO	2
2076	Fungi	NO	0
2093	Fungi	NO	0
2012	Mix	NO	0
2020	Mix	NO	0
2035	Mix	NO	0
2037	Mix	NO	0
2054	Mix	NO	0
2069	Mix	NO	0
2082	Mix	NO	0
2085	Mix	NO	0
2004	Nematode	NO	1
2019	Nematode	NO	0
2029	Nematode	NO	0
2038	Nematode	NO	0
2060	Nematode	NO	0
2061	Nematode	NO	2
2083	Nematode	NO	0
2087	Nematode	NO	0



Yield 2015

Plot	Beneficial	WCR	Weight (g)
2182	Bacteria	NO	600.7263288
2159	Bacteria	NO	640.1321616
2102	Bacteria	NO	664.2292536
2117	Bacteria	NO	676.136052
2139	Bacteria	NO	717.5263512
2130	Bacteria	NO	754.097232
2155	Bacteria	NO	761.4681072
2180	Bacteria	NO	849.068124
2140	Control	NO	319.4990904
2107	Control	NO	346.1476392
2111	Control	NO	346.9981248
2126	Control	NO	448.4894064
2173	Control	NO	517.6622352
2163	Control	NO	679.8214896
2188	Control	NO	686.6253744
2148	Control	NO	841.6972488
2135	Control-Fungi	NO	392.640852
2124	Control-Fungi	NO	403.6971648
2187	Control-Fungi	NO	415.6039632
2106	Control-Fungi	NO	438.000084
2164	Control-Fungi	NO	555.650592
2110	Control-Fungi	NO	572.3768088
2149	Control-Fungi	NO	701.9341152
2174	Control-Fungi	NO	1051.767192
2109	Fungi	NO	536.9399088
2172	Fungi	NO	564.4389432
2158	Fungi	NO	592.2214728
2189	Fungi	NO	601.5768144
2154	Fungi	NO	676.136052
2141	Fungi	NO	707.320524
2132	Fungi	NO	779.3283048
2101	Fungi	NO	806.2603488
2165	Mix	NO	362.5903608
2133	Mix	NO	552.5321448
2116	Mix	NO	622.271964
2178	Mix	NO	709.0214952
2131	Mix	NO	740.4894624
2181	Mix	NO	797.4719976
2108	Mix	NO	925.0448376
2150	Mix	NO	1113.28565
2183	Nematode	NO	551.1146688
2115	Nematode	NO	612.0661368
2134	Nematode	NO	679.8214896
2157	Nematode	NO	747.009852
2179	Nematode	NO	845.0991912
2125	Nematode	NO	898.1127936
2100	Nematode	NO	1047.798259

2156	Nematode	NO	1330.442974
2123	Bacteria	YES	372.2291976
2170	Bacteria	YES	735.1030536
2136	Bacteria	YES	744.1749
2152	Bacteria	YES	770.5399536
2119	Bacteria	YES	833.475888
2186	Bacteria	YES	868.3457976
2097	Bacteria	YES	908.3186208
2162	Bacteria	YES	912.854544
2104	Control	YES	422.9748384
2137	Control	YES	458.9787288
2122	Control	YES	511.708836
2169	Control	YES	599.8758432
2153	Control	YES	607.5302136
2112	Control	YES	633.611772
2161	Control	YES	657.9923592
2192	Control	YES	680.1049848
2128	Control-Fungi	YES	333.9573456
2190	Control-Fungi	YES	393.4913376
2177	Control-Fungi	YES	453.0253296
2167	Control-Fungi	YES	485.3437824
2105	Control-Fungi	YES	516.5282544
2151	Control-Fungi	YES	629.9263344
2113	Control-Fungi	YES	803.1419016
2144	Control-Fungi	YES	922.776876
2185	Fungi	YES	385.553472
2138	Fungi	YES	499.5185424
2118	Fungi	YES	531.5535
2146	Fungi	YES	553.6661256
2099	Fungi	YES	555.9340872
2127	Fungi	YES	579.1806936
2171	Fungi	YES	667.9146912
2166	Fungi	YES	699.6661536
2145	Mix	YES	477.4059168
2176	Mix	YES	555.650592
2129	Mix	YES	573.5107896
2103	Mix	YES	691.4447928
2184	Mix	YES	705.903048
2143	Mix	YES	843.39822
2120	Mix	YES	983.1613536
2168	Mix	YES	995.9186376
2121	Nematode	YES	625.3904112
2147	Nematode	YES	697.9651824
2098	Nematode	YES	700.233144
2160	Nematode	YES	738.504996
2191	Nematode	YES	769.9729632
2142	Nematode	YES	815.6156904
2175	Nematode	YES	851.903076
2114	Nematode	YES	961.3322232

Field design 2016

Plant damage/larval recovery plots

Block	Plot	Treatment	WCR		Beneficial	Code
1	3001	5	NO	1	Control	C
1	3002	4	NO	2	Mix	MIX
1	3003	3	NO	3	Nematode	N
1	3004	2	NO	4	Fungi	F
1	3005	1	NO	5	Bacteria	B
1	3006	6	NO	6	Control-Fungi	C-F
1	3007	9	YES	1	Nematode	N*
1	3008	10	YES	2	Mix	MIX*
1	3009	12	YES	3	Control-Fungi	C-F*
1	3010	11	YES	4	Control	C*
1	3011	7	YES	5	Bacteria	B*
1	3012	8	YES	6	Fungi	F*
2	3013	9	YES	1	Nematode	N*
2	3014	11	YES	2	Control	C*
2	3015	8	YES	3	Fungi	F*
2	3016	10	YES	4	Mix	MIX*
2	3017	12	YES	5	Control-Fungi	C-F*
2	3018	7	YES	6	Bacteria	B*
2	3019	2	NO	1	Fungi	F
2	3020	5	NO	2	Control	C
2	3021	6	NO	3	Control-Fungi	C-F
2	3022	1	NO	4	Bacteria	B
2	3023	3	NO	5	Nematode	N
2	3024	4	NO	6	Mix	MIX
3	3025	5	NO	1	Control	C
3	3026	6	NO	2	Control-Fungi	C-F
3	3027	3	NO	3	Nematode	N
3	3028	1	NO	4	Bacteria	B
3	3029	2	NO	5	Fungi	F
3	3030	4	NO	6	Mix	MIX
3	3031	9	YES	1	Nematode	N*
3	3032	12	YES	2	Control-Fungi	C-F*
3	3033	10	YES	3	Mix	MIX*
3	3034	11	YES	4	Control	C*
3	3035	7	YES	5	Bacteria	B*
3	3036	8	YES	6	Fungi	F*
4	3037	12	YES	1	Control-Fungi	C-F*
4	3038	11	YES	2	Control	C*
4	3039	9	YES	3	Nematode	N*
4	3040	8	YES	4	Fungi	F*
4	3041	10	YES	5	Mix	MIX*
4	3042	7	YES	6	Bacteria	B*
4	3043	6	NO	1	Control-Fungi	C-F
4	3044	5	NO	2	Control	C
4	3045	2	NO	3	Fungi	F
4	3046	4	NO	4	Mix	MIX

4	3047	1	NO	5	Bacteria	B
4	3048	3	NO	6	Nematode	N
5	3049	11	YES	1	Control	C*
5	3050	7	YES	2	Bacteria	B*
5	3051	9	YES	3	Nematode	N*
5	3052	8	YES	4	Fungi	F*
5	3053	10	YES	5	Mix	MIX*
5	3054	12	YES	6	Control-Fungi	C-F*
5	3055	4	NO	1	Mix	MIX
5	3056	6	NO	2	Control-Fungi	C-F
5	3057	3	NO	3	Nematode	N
5	3058	5	NO	4	Control	C
5	3059	1	NO	5	Bacteria	B
5	3060	2	NO	6	Fungi	F
6	3061	4	NO	1	Mix	MIX
6	3062	3	NO	2	Nematode	N
6	3063	1	NO	3	Bacteria	B
6	3064	5	NO	4	Control	C
6	3065	2	NO	5	Fungi	F
6	3066	6	NO	6	Control-Fungi	C-F
6	3067	11	YES	1	Control	C*
6	3068	9	YES	2	Nematode	N*
6	3069	12	YES	3	Control-Fungi	C-F*
6	3070	10	YES	4	Mix	MIX*
6	3071	8	YES	5	Fungi	F*
6	3072	7	YES	6	Bacteria	B*
7	3073	5	NO	1	Control	C
7	3074	4	NO	2	Mix	MIX
7	3075	2	NO	3	Fungi	F
7	3076	1	NO	4	Bacteria	B
7	3077	6	NO	5	Control-Fungi	C-F
7	3078	3	NO	6	Nematode	N
7	3079	10	YES	1	Mix	MIX*
7	3080	11	YES	2	Control	C*
7	3081	7	YES	3	Bacteria	B*
7	3082	9	YES	4	Nematode	N*
7	3083	12	YES	5	Control-Fungi	C-F*
7	3084	8	YES	6	Fungi	F*
8	3085	11	YES	1	Control	C*
8	3086	7	YES	2	Bacteria	B*
8	3087	8	YES	3	Fungi	F*
8	3088	9	YES	4	Nematode	N*
8	3089	10	YES	5	Mix	MIX*
8	3090	12	YES	6	Control-Fungi	C-F*
8	3091	2	NO	1	Fungi	F
8	3092	1	NO	2	Bacteria	B
8	3093	6	NO	3	Control-Fungi	C-F
8	3094	5	NO	4	Control	C
8	3095	4	NO	5	Mix	MIX
8	3096	3	NO	6	Nematode	N

Root damage 2016

Plot	Beneficial	WCR	Damage (1-3)
3005	Bacteria	NO	0.00666667
3022	Bacteria	NO	0.04
3028	Bacteria	NO	0
3047	Bacteria	NO	0
3059	Bacteria	NO	0
3063	Bacteria	NO	0.03333333
3076	Bacteria	NO	0
3092	Bacteria	NO	0
3004	Fungi	NO	0
3019	Fungi	NO	0
3029	Fungi	NO	0
3045	Fungi	NO	0
3060	Fungi	NO	0.04
3065	Fungi	NO	0
3075	Fungi	NO	0
3091	Fungi	NO	0.01666667
3003	Nematode	NO	0
3023	Nematode	NO	0
3027	Nematode	NO	0
3048	Nematode	NO	0.09
3057	Nematode	NO	0
3062	Nematode	NO	0
3078	Nematode	NO	0
3096	Nematode	NO	0
3002	Mix	NO	0
3024	Mix	NO	0
3030	Mix	NO	0
3046	Mix	NO	0
3055	Mix	NO	0
3061	Mix	NO	0
3074	Mix	NO	0.03333333
3095	Mix	NO	0
3001	Control	NO	0
3020	Control	NO	0
3025	Control	NO	0.00666667
3044	Control	NO	0
3058	Control	NO	0
3064	Control	NO	0.08333333
3073	Control	NO	0
3094	Control	NO	0.00666667
3006	Control-Fungi	NO	0.03333333
3021	Control-Fungi	NO	0
3026	Control-Fungi	NO	0
3043	Control-Fungi	NO	0
3056	Control-Fungi	NO	0
3066	Control-Fungi	NO	0.00666667
3077	Control-Fungi	NO	0

3093	Control-Fungi	NO	0
3011	Bacteria	NO	0.28333333
3018	Bacteria	YES	0.58333333
3035	Bacteria	YES	0.35
3042	Bacteria	YES	0.2
3050	Bacteria	YES	0.05666667
3072	Bacteria	YES	0.13333333
3081	Bacteria	YES	0.21666667
3086	Bacteria	YES	0.1
3012	Fungi	YES	0.19
3015	Fungi	YES	0.00666667
3036	Fungi	YES	0.2
3040	Fungi	YES	0.66666667
3052	Fungi	YES	0.66666667
3071	Fungi	YES	0.25
3084	Fungi	YES	0.42333333
3087	Fungi	YES	0.28333333
3007	Nematode	YES	0.1
3013	Nematode	YES	0
3031	Nematode	YES	0.15
3039	Nematode	YES	0.00666667
3051	Nematode	YES	0.18333333
3068	Nematode	YES	0.25666667
3082	Nematode	YES	0.03333333
3088	Nematode	YES	0.51666667
3008	Mix	YES	0.5
3016	Mix	YES	0.09
3033	Mix	YES	0.09
3041	Mix	YES	0.03333333
3053	Mix	YES	0.11666667
3070	Mix	YES	0.33333333
3079	Mix	YES	0.05
3089	Mix	YES	0.25
3010	Control	YES	0.91666667
3014	Control	YES	0
3034	Control	YES	0.7
3038	Control	YES	0.09
3049	Control	YES	0.33333333
3067	Control	YES	0.45
3080	Control	YES	0.20666667
3085	Control	YES	0.36666667
3009	Control-Fungi	YES	0.16666667
3017	Control-Fungi	YES	0.5
3032	Control-Fungi	YES	0.25
3037	Control-Fungi	YES	0.17333333
3054	Control-Fungi	YES	0.58333333
3069	Control-Fungi	YES	0.5
3083	Control-Fungi	YES	0.16666667
3090	Control-Fungi	YES	0.18333333

Larval recovery 2016

Plot	Beneficial	WCR	Larval recovery (mean nb)
3005	Bacteria	NO	0
3022	Bacteria	NO	0
3028	Bacteria	NO	0
3047	Bacteria	NO	1
3059	Bacteria	NO	0.5
3063	Bacteria	NO	0
3076	Bacteria	NO	0
3092	Bacteria	NO	0.5
3004	Fungi	NO	0
3019	Fungi	NO	0
3029	Fungi	NO	0
3045	Fungi	NO	0
3060	Fungi	NO	0.5
3065	Fungi	NO	0
3075	Fungi	NO	0
3091	Fungi	NO	0
3003	Nematode	NO	0
3023	Nematode	NO	0.5
3027	Nematode	NO	0
3048	Nematode	NO	0.5
3057	Nematode	NO	0
3062	Nematode	NO	0
3078	Nematode	NO	0
3096	Nematode	NO	0
3002	Mix	NO	0
3024	Mix	NO	0
3030	Mix	NO	0
3046	Mix	NO	0
3055	Mix	NO	0.5
3061	Mix	NO	0
3074	Mix	NO	0
3095	Mix	NO	0
3001	Control	NO	0.5
3020	Control	NO	0
3025	Control	NO	0
3044	Control	NO	0
3058	Control	NO	1.5
3064	Control	NO	0
3073	Control	NO	0
3094	Control	NO	0
3006	Control-Fungi	NO	0
3021	Control-Fungi	NO	0
3026	Control-Fungi	NO	0
3043	Control-Fungi	NO	0
3056	Control-Fungi	NO	0
3066	Control-Fungi	NO	0
3077	Control-Fungi	NO	0

3093	Control-Fungi	NO	0
3011	Bacteria	YES	2.5
3018	Bacteria	YES	0.5
3035	Bacteria	YES	1.5
3042	Bacteria	YES	0.5
3050	Bacteria	YES	3
3072	Bacteria	YES	2
3081	Bacteria	YES	1
3086	Bacteria	YES	1.5
3012	Fungi	YES	6.5
3015	Fungi	YES	4.5
3036	Fungi	YES	0.5
3040	Fungi	YES	2.5
3052	Fungi	YES	3
3071	Fungi	YES	2
3084	Fungi	YES	3
3087	Fungi	YES	5
3007	Nematode	YES	4
3013	Nematode	YES	0.5
3031	Nematode	YES	4
3039	Nematode	YES	2
3051	Nematode	YES	5
3068	Nematode	YES	0
3082	Nematode	YES	0.5
3088	Nematode	YES	6.5
3008	Mix	YES	6
3016	Mix	YES	2
3033	Mix	YES	3.5
3041	Mix	YES	1.5
3053	Mix	YES	2
3070	Mix	YES	4.5
3079	Mix	YES	1.5
3089	Mix	YES	4
3010	Control	YES	7
3014	Control	YES	1.5
3034	Control	YES	4.5
3038	Control	YES	1.5
3049	Control	YES	3.5
3067	Control	YES	5
3080	Control	YES	6
3085	Control	YES	5.5
3009	Control-Fungi	YES	2.5
3017	Control-Fungi	YES	6.5
3032	Control-Fungi	YES	9
3037	Control-Fungi	YES	1.5
3054	Control-Fungi	YES	8.5
3069	Control-Fungi	YES	3.5
3083	Control-Fungi	YES	0.5
3090	Control-Fungi	YES	5.5



Larval weight 2016

Plot	Beneficial	WCR	larval weight (mg)
3005	Bacteria	NO	.
3022	Bacteria	NO	.
3028	Bacteria	NO	.
3047	Bacteria	NO	0.234
3059	Bacteria	NO	.
3063	Bacteria	NO	.
3076	Bacteria	NO	.
3092	Bacteria	NO	0.247
3004	Fungi	NO	.
3019	Fungi	NO	.
3029	Fungi	NO	.
3045	Fungi	NO	.
3060	Fungi	NO	0.342
3065	Fungi	NO	.
3075	Fungi	NO	.
3091	Fungi	NO	.
3003	Nematode	NO	.
3023	Nematode	NO	0.106
3027	Nematode	NO	.
3048	Nematode	NO	0.217
3057	Nematode	NO	.
3062	Nematode	NO	.
3078	Nematode	NO	.
3096	Nematode	NO	.
3002	Mix	NO	.
3024	Mix	NO	.
3030	Mix	NO	.
3046	Mix	NO	.
3055	Mix	NO	0.092
3061	Mix	NO	.
3074	Mix	NO	.
3095	Mix	NO	.
3001	Control	NO	0.241
3020	Control	NO	.
3025	Control	NO	.
3044	Control	NO	.
3058	Control	NO	0.552
3064	Control	NO	.
3073	Control	NO	.
3094	Control	NO	.
3006	Control-Fungi	NO	.
3021	Control-Fungi	NO	.
3026	Control-Fungi	NO	.
3043	Control-Fungi	NO	.
3056	Control-Fungi	NO	.
3066	Control-Fungi	NO	.
3077	Control-Fungi	NO	.

3093 Control-Fungi	NO	
3011 Bacteria	YES	0.482
3018 Bacteria	YES	0.231
3035 Bacteria	YES	0.2275
3042 Bacteria	YES	0.055
3050 Bacteria	YES	0.09
3072 Bacteria	YES	0.314
3081 Bacteria	YES	0.268
3086 Bacteria	YES	0.276
3012 Fungi	YES	0.769
3015 Fungi	YES	0.572
3036 Fungi	YES	0.18
3040 Fungi	YES	0.3465
3052 Fungi	YES	0.759
3071 Fungi	YES	0.266
3084 Fungi	YES	0.307
3087 Fungi	YES	0.6565
3007 Nematode	YES	0.6145
3013 Nematode	YES	0.123
3031 Nematode	YES	0.376
3039 Nematode	YES	0.925
3051 Nematode	YES	0.4655
3068 Nematode	YES	
3082 Nematode	YES	0.187
3088 Nematode	YES	1.1145
3008 Mix	YES	0.757
3016 Mix	YES	0.2375
3033 Mix	YES	0.3925
3041 Mix	YES	0.201
3053 Mix	YES	0.2405
3070 Mix	YES	0.8075
3079 Mix	YES	0.2625
3089 Mix	YES	0.8045
3010 Control	YES	1.985
3014 Control	YES	0.1705
3034 Control	YES	0.665
3038 Control	YES	0.2185
3049 Control	YES	0.447
3067 Control	YES	0.6805
3080 Control	YES	1.696
3085 Control	YES	0.984
3009 Control-Fungi	YES	0.989
3017 Control-Fungi	YES	0.915
3032 Control-Fungi	YES	1.1335
3037 Control-Fungi	YES	0.345
3054 Control-Fungi	YES	1.0185
3069 Control-Fungi	YES	0.689
3083 Control-Fungi	YES	0.196
3090 Control-Fungi	YES	1.1115

Yield 2016

Plot	Beneficial	WCR	Weight (g)
3099	Bacteria	NO	1167.4
3117	Bacteria	NO	812.1
3127	Bacteria	NO	904.3
3134	Bacteria	NO	717.3
3150	Bacteria	NO	537.4
3157	Bacteria	NO	607.3
3180	Bacteria	NO	832.1
3184	Bacteria	NO	752.3
3100	Fungi	NO	1054.9
3119	Fungi	NO	886.8
3129	Fungi	NO	822.7
3137	Fungi	NO	915.4
3146	Fungi	NO	663.6
3158	Fungi	NO	745
3175	Fungi	NO	741.7
3185	Fungi	NO	604.9
3097	Nematode	NO	1013
3120	Nematode	NO	577.3
3131	Nematode	NO	855.6
3136	Nematode	NO	831.1
3145	Nematode	NO	700.3
3162	Nematode	NO	642
3179	Nematode	NO	868.7
3183	Nematode	NO	601.5
3101	Mix	NO	1120.5
3116	Mix	NO	850.1
3130	Mix	NO	727.6
3135	Mix	NO	856.8
3149	Mix	NO	602.5
3159	Mix	NO	700.3
3177	Mix	NO	1024.4
3186	Mix	NO	870.6
3098	Control	NO	1031.8
3115	Control	NO	1194
3128	Control	NO	627.6
3133	Control	NO	853.9
3147	Control	NO	605
3161	Control	NO	646.3
3178	Control	NO	817
3181	Control	NO	1156.5
3102	Control-Fungi	NO	1164.8
3118	Control-Fungi	NO	909
3132	Control-Fungi	NO	713.1
3138	Control-Fungi	NO	664.9
3148	Control-Fungi	NO	896.7
3160	Control-Fungi	NO	505.9
3176	Control-Fungi	NO	833.3

3182	Control-Fungi	NO	1053.5
3103	Bacteria	YES	1131
3114	Bacteria	YES	882.4
3123	Bacteria	YES	672.9
3143	Bacteria	YES	955.5
3151	Bacteria	YES	758.5
3166	Bacteria	YES	670.2
3170	Bacteria	YES	982
3187	Bacteria	YES	941.6
3105	Fungi	YES	1044.6
3109	Fungi	YES	1004.2
3122	Fungi	YES	748.5
3144	Fungi	YES	791.4
3153	Fungi	YES	764.7
3163	Fungi	YES	638.6
3174	Fungi	YES	992
3190	Fungi	YES	567
3107	Nematode	YES	899.2
3112	Nematode	YES	983.6
3124	Nematode	YES	742.9
3139	Nematode	YES	1035.3
3155	Nematode	YES	764.3
3167	Nematode	YES	503.7
3173	Nematode	YES	896.7
3189	Nematode	YES	690.1
3104	Mix	YES	1062.2
3111	Mix	YES	804
3121	Mix	YES	945.7
3141	Mix	YES	807
3154	Mix	YES	761.6
3164	Mix	YES	750.8
3169	Mix	YES	954.4
3188	Mix	YES	1004.1
3106	Control	YES	613.1
3113	Control	YES	820.9
3125	Control	YES	641.2
3142	Control	YES	874.6
3152	Control	YES	726.6
3165	Control	YES	776.6
3171	Control	YES	1000.9
3192	Control	YES	676.5
3108	Control-Fungi	YES	928.1
3110	Control-Fungi	YES	1003.8
3126	Control-Fungi	YES	600.8
3140	Control-Fungi	YES	1072.8
3156	Control-Fungi	YES	615.4
3168	Control-Fungi	YES	575.9
3172	Control-Fungi	YES	1016.6
3191	Control-Fungi	YES	505.2

Field design 2017

Plant damage/larval recovery plots

Block	Plot	Treatment	WCR		Beneficial	Code
1	1001	12	add	1	Control-Fungi	C-F*
1	1002	8	add	2	Fungi	F*
1	1003	11	add	3	Control	C*
1	1004	9	add	4	Nematode	N*
1	1005	7	add	5	Bacteria	B*
1	1006	10	add	6	Mix	MIX*
1	1007	4	neg	1	Mix	MIX
1	1008	6	neg	2	Control-Fungi	C-F
1	1009	2	neg	3	Fungi	F
1	1010	1	neg	4	Bacteria	B
1	1011	3	neg	5	Nematode	N
1	1012	5	neg	6	Control	C
2	1013	9	add	1	Nematode	N*
2	1014	8	add	2	Fungi	F*
2	1015	12	add	3	Control-Fungi	C-F*
2	1016	11	add	4	Control	C*
2	1017	10	add	5	Mix	MIX*
2	1018	7	add	6	Bacteria	B*
2	1019	5	neg	1	Control	C
2	1020	3	neg	2	Nematode	N
2	1021	1	neg	3	Bacteria	B
2	1022	4	neg	4	Mix	MIX
2	1023	2	neg	5	Fungi	F
2	1024	6	neg	6	Control-Fungi	C-F
3	1025	4	neg	1	Mix	MIX
3	1026	1	neg	2	Bacteria	B
3	1027	5	neg	3	Control	C
3	1028	3	neg	4	Nematode	N
3	1029	2	neg	5	Fungi	F
3	1030	6	neg	6	Control-Fungi	C-F
3	1031	8	add	1	Fungi	F*
3	1032	9	add	2	Nematode	N*
3	1033	7	add	3	Bacteria	B*
3	1034	11	add	4	Control	C*
3	1035	10	add	5	Mix	MIX*
3	1036	12	add	6	Control-Fungi	C-F*
4	1037	8	add	1	Fungi	F*
4	1038	9	add	2	Nematode	N*
4	1039	12	add	3	Control-Fungi	C-F*
4	1040	11	add	4	Control	C*
4	1041	10	add	5	Mix	MIX*
4	1042	7	add	6	Bacteria	B*
4	1043	5	neg	1	Control	C
4	1044	2	neg	2	Fungi	F
4	1045	1	neg	3	Bacteria	B
4	1046	6	neg	4	Control-Fungi	C-F

4	1047	4	neg	5	Mix	MIX
4	1048	3	neg	6	Nematode	N
5	1049	12	add	1	Control-Fungi	C-F*
5	1050	10	add	2	Mix	MIX*
5	1051	8	add	3	Fungi	F*
5	1052	11	add	4	Control	C*
5	1053	9	add	5	Nematode	N*
5	1054	7	add	6	Bacteria	B*
5	1055	1	neg	1	Bacteria	B
5	1056	2	neg	2	Fungi	F
5	1057	4	neg	3	Mix	MIX
5	1058	6	neg	4	Control-Fungi	C-F
5	1059	3	neg	5	Nematode	N
5	1060	5	neg	6	Control	C
6	1061	9	add	1	Nematode	N*
6	1062	11	add	2	Control	C*
6	1063	8	add	3	Fungi	F*
6	1064	10	add	4	Mix	MIX*
6	1065	7	add	5	Bacteria	B*
6	1066	12	add	6	Control-Fungi	C-F*
6	1067	6	neg	1	Control-Fungi	C-F
6	1068	5	neg	2	Control	C
6	1069	1	neg	3	Bacteria	B
6	1070	3	neg	4	Nematode	N
6	1071	4	neg	5	Mix	MIX
6	1072	2	neg	6	Fungi	F
7	1073	2	neg	1	Fungi	F
7	1074	4	neg	2	Mix	MIX
7	1075	6	neg	3	Control-Fungi	C-F
7	1076	3	neg	4	Nematode	N
7	1077	1	neg	5	Bacteria	B
7	1078	5	neg	6	Control	C
7	1079	11	add	1	Control	C*
7	1080	7	add	2	Bacteria	B*
7	1081	12	add	3	Control-Fungi	C-F*
7	1082	10	add	4	Mix	MIX*
7	1083	8	add	5	Fungi	F*
7	1084	9	add	6	Nematode	N*
8	1085	2	neg	1	Fungi	F
8	1086	5	neg	2	Control	C
8	1087	6	neg	3	Control-Fungi	C-F
8	1088	4	neg	4	Mix	MIX
8	1089	3	neg	5	Nematode	N
8	1090	1	neg	6	Bacteria	B
8	1091	7	add	1	Bacteria	B*
8	1092	8	add	2	Fungi	F*
8	1093	9	add	3	Nematode	N*
8	1094	10	add	4	Mix	MIX*
8	1095	11	add	5	Control	C*
8	1096	12	add	6	Control-Fungi	C-F*

Root damage 2017

Plot	WCR	Beneficial	Damage (1-3)
1005	add	Bacteria	0.766666667
1018	add	Bacteria	0.7
1033	add	Bacteria	1.166666667
1042	add	Bacteria	0.166666667
1054	add	Bacteria	1
1065	add	Bacteria	1.5
1080	add	Bacteria	0.116666667
1091	add	Bacteria	0.933333333
1010	neg	Bacteria	0
1021	neg	Bacteria	0
1026	neg	Bacteria	0.003333333
1045	neg	Bacteria	0.005
1055	neg	Bacteria	0
1069	neg	Bacteria	0.003333333
1077	neg	Bacteria	0
1090	neg	Bacteria	0.006666667
1003	add	Control	1.35
1016	add	Control	0.916666667
1034	add	Control	1.333333333
1040	add	Control	0.216666667
1052	add	Control	2.083333333
1062	add	Control	1.083333333
1079	add	Control	0.75
1095	add	Control	1.666666667
1012	neg	Control	0
1019	neg	Control	0
1027	neg	Control	0.003333333
1043	neg	Control	0
1060	neg	Control	0
1068	neg	Control	0
1078	neg	Control	0
1086	neg	Control	0
1001	add	Control-Fungi	0.253333333
1015	add	Control-Fungi	2.333333333
1036	add	Control-Fungi	1.166666667
1039	add	Control-Fungi	2.083333333
1049	add	Control-Fungi	2
1066	add	Control-Fungi	1.25
1081	add	Control-Fungi	1.666666667
1096	add	Control-Fungi	1.916666667
1008	neg	Control-Fungi	0
1024	neg	Control-Fungi	0.003333333
1030	neg	Control-Fungi	0
1046	neg	Control-Fungi	0.003333333
1058	neg	Control-Fungi	0.016666667
1067	neg	Control-Fungi	0.016666667
1075	neg	Control-Fungi	0

1087 neg	Control-Fungi	0.016666667
1002 add	Fungi	1.083333333
1014 add	Fungi	3
1031 add	Fungi	1.75
1037 add	Fungi	2.416666667
1051 add	Fungi	0.683333333
1063 add	Fungi	2.75
1083 add	Fungi	0.866666667
1092 add	Fungi	1.916666667
1009 neg	Fungi	0.006666667
1023 neg	Fungi	0
1029 neg	Fungi	0.006666667
1044 neg	Fungi	0.003333333
1056 neg	Fungi	0.03
1072 neg	Fungi	0.003333333
1073 neg	Fungi	0
1085 neg	Fungi	0.033333333
1006 add	Mix	0.25
1017 add	Mix	0.95
1035 add	Mix	2
1041 add	Mix	0.7
1050 add	Mix	0.683333333
1064 add	Mix	1
1082 add	Mix	0.833333333
1094 add	Mix	0.833333333
1007 neg	Mix	0.003333333
1022 neg	Mix	0
1025 neg	Mix	0.173333333
1047 neg	Mix	0
1057 neg	Mix	0
1071 neg	Mix	0
1074 neg	Mix	0
1088 neg	Mix	0
1004 add	Nematode	1.1
1013 add	Nematode	0.283333333
1032 add	Nematode	1.5
1038 add	Nematode	1.083333333
1053 add	Nematode	1.833333333
1061 add	Nematode	1.5
1084 add	Nematode	0.333333333
1093 add	Nematode	0.333333333
1011 neg	Nematode	0
1020 neg	Nematode	0
1028 neg	Nematode	0
1048 neg	Nematode	0.003333333
1059 neg	Nematode	0
1070 neg	Nematode	0
1076 neg	Nematode	0
1089 neg	Nematode	0



Larval recovery 2017

Plot	WCR	Beneficial	Larval recovery (mean nb)	
	1010	NO	Bacteria	0
	1021	NO	Bacteria	0
	1026	NO	Bacteria	0
	1045	NO	Bacteria	0
	1055	NO	Bacteria	0
	1069	NO	Bacteria	0.5
	1077	NO	Bacteria	0
	1090	NO	Bacteria	1
	1009	NO	Fungi	0
	1023	NO	Fungi	0
	1029	NO	Fungi	1
	1044	NO	Fungi	0
	1056	NO	Fungi	0
	1072	NO	Fungi	0
	1073	NO	Fungi	0
	1085	NO	Fungi	0
	1011	NO	Nematode	0
	1020	NO	Nematode	0
	1028	NO	Nematode	0
	1048	NO	Nematode	0
	1059	NO	Nematode	0
	1070	NO	Nematode	1
	1076	NO	Nematode	0.5
	1089	NO	Nematode	0
	1007	NO	Mix	0
	1022	NO	Mix	0
	1025	NO	Mix	0
	1047	NO	Mix	0
	1057	NO	Mix	0
	1071	NO	Mix	0
	1074	NO	Mix	0
	1088	NO	Mix	0
	1012	NO	Control	0
	1019	NO	Control	0
	1027	NO	Control	0
	1043	NO	Control	1
	1060	NO	Control	0.5
	1068	NO	Control	0
	1078	NO	Control	0
	1086	NO	Control	0
	1008	NO	Control-Fungi	0
	1024	NO	Control-Fungi	0
	1030	NO	Control-Fungi	0
	1046	NO	Control-Fungi	0
	1058	NO	Control-Fungi	0
	1067	NO	Control-Fungi	0
	1075	NO	Control-Fungi	0

1087 NO	Control-Fungi	0
1005 YES	Bacteria	4
1018 YES	Bacteria	4
1033 YES	Bacteria	1.5
1042 YES	Bacteria	2
1054 YES	Bacteria	3
1065 YES	Bacteria	5.5
1080 YES	Bacteria	7.5
1091 YES	Bacteria	0.5
1002 YES	Fungi	1.5
1014 YES	Fungi	6.5
1031 YES	Fungi	7.5
1037 YES	Fungi	21.5
1051 YES	Fungi	3
1063 YES	Fungi	4.5
1083 YES	Fungi	6
1092 YES	Fungi	5.5
1004 YES	Nematode	6.5
1013 YES	Nematode	3.5
1032 YES	Nematode	3.5
1038 YES	Nematode	5
1053 YES	Nematode	2
1061 YES	Nematode	2.5
1084 YES	Nematode	1
1093 YES	Nematode	9
1006 YES	Mix	1.5
1017 YES	Mix	2
1035 YES	Mix	4.5
1041 YES	Mix	5
1050 YES	Mix	4.5
1064 YES	Mix	4.5
1082 YES	Mix	6
1094 YES	Mix	5
1003 YES	Control	4
1016 YES	Control	1.5
1034 YES	Control	3
1040 YES	Control	1
1052 YES	Control	1.5
1062 YES	Control	4
1079 YES	Control	4
1095 YES	Control	6
1001 YES	Control-Fungi	5
1015 YES	Control-Fungi	11.5
1036 YES	Control-Fungi	3
1039 YES	Control-Fungi	4.5
1049 YES	Control-Fungi	2.5
1066 YES	Control-Fungi	3.5
1081 YES	Control-Fungi	3.5
1096 YES	Control-Fungi	10

Larval weight 2017

Plot	WCR	Beneficial	larval weight (mg)
1005	YES	Bacteria	1.2575
1018	YES	Bacteria	1.02
1033	YES	Bacteria	0.78333333
1042	YES	Bacteria	1.095
1054	YES	Bacteria	0.85666667
1065	YES	Bacteria	1.26454545
1080	YES	Bacteria	1.02466667
1091	YES	Bacteria	.
1002	YES	Fungi	0.84
1014	YES	Fungi	0.95307692
1031	YES	Fungi	0.98733333
1037	YES	Fungi	0.93348837
1051	YES	Fungi	0.81666667
1063	YES	Fungi	1.20333333
1083	YES	Fungi	1.38916667
1092	YES	Fungi	1.34909091
1004	YES	Nematode	1.04307692
1013	YES	Nematode	1.34285714
1032	YES	Nematode	0.80428571
1038	YES	Nematode	1.01
1053	YES	Nematode	0.935
1061	YES	Nematode	1.228
1084	YES	Nematode	1.005
1093	YES	Nematode	0.56388889
1006	YES	Mix	1.26666667
1017	YES	Mix	2.0025
1035	YES	Mix	0.94777778
1041	YES	Mix	1.39
1050	YES	Mix	1.29444444
1064	YES	Mix	1.08222222
1082	YES	Mix	1.47166667
1094	YES	Mix	1.414
1003	YES	Control	1.1825
1016	YES	Control	1.63333333
1034	YES	Control	0.75333333
1040	YES	Control	0.64
1052	YES	Control	1.08666667
1062	YES	Control	1.3825
1079	YES	Control	0.5075
1095	YES	Control	1.09
1001	YES	Control-Fungi	1.554
1015	YES	Control-Fungi	1.13565217
1036	YES	Control-Fungi	0.49333333
1039	YES	Control-Fungi	0.88222222
1049	YES	Control-Fungi	1.12
1066	YES	Control-Fungi	1.40857143
1081	YES	Control-Fungi	0.64428571

1096 YES	Control-Fungi	0.8685
1010 NO	Bacteria .	
1021 NO	Bacteria .	
1026 NO	Bacteria .	
1045 NO	Bacteria .	
1055 NO	Bacteria .	
1069 NO	Bacteria .	0.06
1077 NO	Bacteria .	
1090 NO	Bacteria .	1.215
1009 NO	Fungi .	
1023 NO	Fungi .	
1029 NO	Fungi .	0.47
1044 NO	Fungi .	
1056 NO	Fungi .	
1072 NO	Fungi .	
1073 NO	Fungi .	
1085 NO	Fungi .	
1011 NO	Nematode .	
1020 NO	Nematode .	
1028 NO	Nematode .	
1048 NO	Nematode .	
1059 NO	Nematode .	
1070 NO	Nematode .	0.065
1076 NO	Nematode .	1.6
1089 NO	Nematode .	
1007 NO	Mix .	
1022 NO	Mix .	
1025 NO	Mix .	
1047 NO	Mix .	
1057 NO	Mix .	
1071 NO	Mix .	
1074 NO	Mix .	
1088 NO	Mix .	
1012 NO	Control .	
1019 NO	Control .	
1027 NO	Control .	
1043 NO	Control .	2.17
1060 NO	Control .	1.78
1068 NO	Control .	
1078 NO	Control .	
1086 NO	Control .	
1008 NO	Control-Fungi .	
1024 NO	Control-Fungi .	
1030 NO	Control-Fungi .	
1046 NO	Control-Fungi .	
1058 NO	Control-Fungi .	
1067 NO	Control-Fungi .	
1075 NO	Control-Fungi .	
1087 NO	Control-Fungi .	

Yield 2017

Plot	WCR	Beneficial	Weight (g)
1102	NO	Bacteria	443.783688
1110	NO	Bacteria	1227.53008
1128	NO	Bacteria	755.98227
1138	NO	Bacteria	740.594697
1148	NO	Bacteria	816.292
1164	NO	Bacteria	936.902344
1180	NO	Bacteria	305.454887
1186	NO	Bacteria	841.72695
1101	NO	Fungi	725.445255
1109	NO	Fungi	835.875
1130	NO	Fungi	1190.05556
1133	NO	Fungi	726.046763
1146	NO	Fungi	587.652174
1167	NO	Fungi	1056.24809
1177	NO	Fungi	1112.24609
1185	NO	Fungi	712.369919
1098	NO	Nematode	1008.368
1112	NO	Nematode	568.036885
1131	NO	Nematode	544.665441
1135	NO	Nematode	740.582677
1147	NO	Nematode	572.934307
1168	NO	Nematode	1147.24409
1176	NO	Nematode	976.150376
1181	NO	Nematode	1282.45652
1100	NO	Mix	824.272358
1111	NO	Mix	897.25188
1132	NO	Mix	429.951493
1134	NO	Mix	829.46831
1150	NO	Mix	907.556
1163	NO	Mix	991.03125
1179	NO	Mix	794.267361
1183	NO	Mix	1017.172
1097	NO	Control	1290.43852
1114	NO	Control	771.776
1127	NO	Control	838.357664
1137	NO	Control	888.666667
1149	NO	Control	587.304688
1165	NO	Control	505.92
1178	NO	Control	1196.38372
1184	NO	Control	677.651079
1099	NO	Control-Fungi	664.30315
1113	NO	Control-Fungi	682.834615
1129	NO	Control-Fungi	970.164234
1136	NO	Control-Fungi	626.488372
1145	NO	Control-Fungi	1197.23358
1166	NO	Control-Fungi	1101.78276
1175	NO	Control-Fungi	727.190141

1182 NO	Control-Fungi	1117.14815
1108 YES	Bacteria	999.566929
1118 YES	Bacteria	924.639098
1125 YES	Bacteria	1053.44643
1139 YES	Bacteria	948.023256
1155 YES	Bacteria	684.165441
1161 YES	Bacteria	655.822222
1171 YES	Bacteria	1007.748
1190 YES	Bacteria	988.620301
1107 YES	Fungi	842.698529
1115 YES	Fungi	688.774074
1124 YES	Fungi	665.503571
1144 YES	Fungi	639.79927
1152 YES	Fungi	658.467153
1159 YES	Fungi	666.172535
1173 YES	Fungi	1009.66783
1192 YES	Fungi	1322.956
1105 YES	Nematode	1179.48162
1119 YES	Nematode	990.277778
1122 YES	Nematode	924.985294
1142 YES	Nematode	1209.58271
1153 YES	Nematode	496.120155
1157 YES	Nematode	290.914179
1169 YES	Nematode	1316.91729
1189 YES	Nematode	1117.612
1106 YES	Mix	1190.32677
1117 YES	Mix	1183.32813
1123 YES	Mix	928.189781
1143 YES	Mix	313.640152
1151 YES	Mix	493.58156
1160 YES	Mix	907.433824
1174 YES	Mix	147.306569
1191 YES	Mix	837.446043
1103 YES	Control	1072.82963
1116 YES	Control	551.406716
1126 YES	Control	874.507634
1140 YES	Control	797.865248
1156 YES	Control	889.043796
1162 YES	Control	1055.59559
1172 YES	Control	913.995935
1188 YES	Control	1158.53759
1104 YES	Control-Fungi	494.518382
1120 YES	Control-Fungi	1026.19466
1121 YES	Control-Fungi	610.240741
1141 YES	Control-Fungi	900.419847
1154 YES	Control-Fungi	905.753571
1158 YES	Control-Fungi	479.869919
1170 YES	Control-Fungi	770.434109