Soil Biology Report Performed By:

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Date Observed: 11-16-2023

Sample Name: Vermicompost Tea Sample Type: Compost Tea

Plants Present/Desired: Brassicas

Plant Succession: Early Successional Brassica

## **Beneficial Microorganisms**

		mended inge	Sample Results	
Fungi (ug/g)	32	90	0	None Detected: Please contact your Soil Biology Consultant.
Standard Deviation			0	Distribution of the target organisms in the sample was uniform; variation was small.
Bacteria (ug/g)	135	270	9	Low: The bacterial biomass is below the recommended minimum level for your plant's stage in succession. Please contact your Soil Biology Consultant.
Standard Deviation			1	Distribution of the target organisms in the sample was uniform; variation was small.
Actinobacteria (ug/g)	10	16	0.14	Low: The actinobacterial biomass is below the recommended minimum level for brassicas. Please contact your Soil Biology Consultant.
Standard Deviation			0.08	Target organisms were present in the sample, but extremely patchy in distribution.  Precision is poor.
F:B Ratio	0.2:1	0.4:1	0.0	The F:B ratio is low. Increase fungal biomass or reduce bacterial biomass, and check predators to assess balance. Please contact your Soil Biology Consultant.

## Minimum Value

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Protozoa (Total)	> 10,000	9,918	Low: The number of beneficial protozoa is below the minimum requirement. Please contact your Soil Biology Consultant.
Standard Deviation		9,054	Few target organism were present and variability was very high. Precision is very low.
Flagellate (#/g)	(See Total)	3,306	
Standard Deviation		7,392	
Amoebae (#/g)	(See Total)	6,612	
Standard Deviation		9,054	

#### Nematodes

Bacterial-feeding (#/g)	100	0	None detected: Bacterial-feeding nematodes help keep bacterial populations in balance and enhance nutrient cycling.
Fungal-feeding (#/g)	0	0	None detected: Fungal-feeding nematodes help to release nutrients from fungal hyphae to the plants.
Predatory (#/g)	0	0	None detected: Predatory nematodes help reduce root-feeding nematode numbers.

# **Detrimental Microorganisms**

Disease-Causing Fungi	Maximum Value	Sample Results	
Oomycetes (ug/g)	0	0	None detected: No disease-causing fungi were observed in the sample. Great!
Standard Deviation		0	Distribution of the target organisms in the sample was uniform; variation was small.
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Anaerobic Protozoa			
Ciliate (#/g)	0	0	None detected: No ciliates were observed in the sample. Aerobic conditions prevail.  Great!
Standard Deviation		0	Distribution of the target organisms in the sample was uniform; variation was small.
Nematode			
Root-feeding (#/g)	0	0	None detected: No root-feeding nematodes were observed. Great!

### **Additional Comments:**