Study on Carbon Source Screening Technology for Prevention and Control of Banana Fusarium Wilt

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Corresponding Author: Xiaolin Fan South China Agricultural University, Guangzhou 510642, China Email: xlfan@scau.edu.cn Abstract: With the spread of banana Fusarium wilt and the aggravation of the damage, it has become the biggest threat to the banana industry. Many studies have shown that bio-organic fertilizer plays an important role in the prevention and control of banana wilt, but most studies focused on biocontrol bacterium screening, not screening carbon sources that could prevent banana wilt, which lead to organic fertilizer fertilization without better effect. So that, a pot culture experiment (I) with inoculating Fusarium oxysporum f. sp. Cubense (FOC) on bananas was carried out to screen the carbon sources, and d-malic acid (CS2) to prevent and control banana wilt had been screened out by Biolog technology and Principal Component Analysis of microbial community level physiological profiling. To verify application effects of CS2, another banana pot experiment (II) with FOC inoculation had been done. The results showed that: (1) CS2 can effectively improve the activity of soil microorganisms. (2) CS2 can significantly reduce FOC population in soil, and the FOC population treated with CS2 was respectively lower than that treated with control carbon source (CS1) and without carbon source (CK1) 48% and 59%. (3) The banana wilt infection rate and disease index of CS2 treatment were significantly smaller than those of CS1 and CK1. (4) CS2 can promote the growth and biomass accumulation of bananas in pathogenic soil. In conclusion, the screening technology studied here can be applied to screen out the carbon sources for prevention of banana Fusarium wilt, which has great significance to the improvement of prevention effect of biological organic fertilizers to banana wilt.

Keywords: Carbon Sources, Soil Microbes, FOC, Bananas

Introduction

Banana Fusarium wilt is a fungal disease of banana and a typical soil-borne vascular bundle disease caused by FOC infection (Wu *et al.*, 2013). This kind of pathogen invades the plant mainly from the root or wound of the banana, through the vascular bundle of banana spreads to the upper part of the pseudostem and leaves and causes the xylem tube be blocked, the nutrition cannot be transmitted normally, finally the plant will withers and dies (Huang *et al.*, 2019). Especially the *Fusarium oxysporum* f. sp. *Cubense* species-4 (FOC4), which is the most harmful to bananas, has strong pathogenicity and long survival and latent time, it's difficult to control and can infect almost all kinds of bananas (Li *et al.*, 2015). Therefore, Fusarium wilt caused by FOC4 is currently recognized as a devastating banana disease in the world. It is a worldwide scientific problem that has plagued banana production for many years and also the biggest obstacle to the development of the world banana industry.

At present, the control methods of banana Fusarium wilt mainly include breeding disease-resistant varieties of bananas, killing FOC with chemical agents, and the cultivation measures of preventing and controlling FOC such as rotation, and the biological control methods for inhibiting growth of FOC by exogenous or indigenous antagonistic bacteria. Among them, the biological control represented by biological organic fertilizers with antagonistic bacteria played an important role in inhibiting pathogenic bacteria (Wu *et al.*, 2009), moreover, the carbon sources in organic fertilizers can



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significantly improve activity, functional diversity and community structure of soil microbes, and inhibit FOC through beneficial indigenous microorganisms and prevent soil-borne diseases (Lee, 2010). Therefore, biological organic fertilizer not only is more and more important for prevention and control of banana Fusarium wilt, but also is environmentally friendly and pollutionfree, so many scholars have conducted relevant studies on it (Huang et al., 2017). However, most of these studies focused on screening biocontrol bacteria that could inhibit FOC and did not screen organic carbon sources that could prevent and control banana Fusarium wilt according to the metabolic characteristics of soil microorganisms. Microorganisms in different soil have different preferences for organic carbon sources (Dwivedi et al., 2013, Xue et al., 2015), which leads to great differences in the ecological effects of organic fertilizers on soil microorganisms. In addition, the exogenous antagonistic bacteria could not adapt to the new soil environment and colonize in it when they entered the field with the organic fertilizers, thus they could not survive and reproduce normally or play the role of inhibiting pathogens (Shen et al., 2017), which affected the biocontrol effect of organic fertilizers. Therefore, the results of most studies on biological organic fertilizers are only have good effects in the indoor environment, but there are some problems in the field environment, such as the exogenous antagonistic bacteria cannot colonize in the new environment or the additional carbon sources cannot be utilized efficiently by beneficial indigenous microorganisms, which directly lead to the unsatisfactory effect of biological control in the field conditions.

In view of the above problems, according to the carbon metabolism characteristics of microbes in soils inoculated with FOC4 and without FOC4, the carbon source that could prevent and control banana Fusarium wilt was selected by Biolog technology and R language and studied the effect and microbial mechanism of this carbon source on preventing and controlling banana Fusarium wilt through the banana pot experiment. This study will lay a foundation for the prevention and control of banana Fusarium wilt by screening carbon sources and is expected to improve the prevention and control effect of organic fertilizer on banana Fusarium wilt in the field, which is of great significance to maintain the sustainable productivity of bananas in the affected area.

Materials and Methods

Experimental Materials

The soil used in these experiments was a sandy loam soil from a farm at South China Agricultural University. The pH of the soil was 5.98. The soil contained 11.09g/kg organic matter (Potassium bichromate titrimetric method-outside heating method), 52.59 mg/kg available nitrogen (Alkaline hydrolyzation diffusion method), 56.57 mg/kg available phosphorus (Olsen's method) and 115.40 mg/kg available potassium (NH₄OAc extraction-flame photometry method).

Sterile tissue-cultured banana plants (*Musa* sp., AAA, Giant Cavendish cv. Baxi,) with 6 leaves were used for the experiments.

Spores of *Fusarium oxysporum* f. sp. *cubense* (E. F. Smith) Snyder and Hansen (FOC4) was supplied by the Plant Pathology Lab of the College of Natural Resource and Environment of South China Agriculture University. The concentration of the FOC4 spore suspension was 1×10^7 cfu/ml.

The carbon sources required for experiment II were selected through experiment I, and they were the control carbon source (CS1) and the carbon source for preventing banana wilt (CS2), respectively. Then, we purchased CS1 and CS2 of analytically pure (AR) from Promega Corporation (USA) as test carbon sources.

Study Design

Experiment I

This experiment was a single factor completely randomized design, that is, different FOC concentrations. Thus, there were 3 treatments. Specifically, the tested FOC concentrations were 0, 10^3 and 10^6 cfu per gram of soil, termed C0, C1 and C2 respectively.

The experiment was performed in a greenhouse at South China Agricultural University (latitude $23^{\circ}09'40''$ N, longitude $113^{\circ}21'28''$ E) in November 2014 to January 2015. Each pot was $15 \times 13 \times 12$ cm and contained 500g of the soil with one banana. Each banana had at least 6 leaves at the start of the experiment (12 November 2014). On 22 November 2014, the C0, C1 and C2 FOC spore suspension treatments were added to the 500mg of the soil around each banana pseudostem from a syringe containing 50ml of 0, 1×10^4 , 1×10^7 cfu/ml, respectively.

The 50 ml of 0.6% compound fertilizer (22N:8P:15K) solution were irrigated into each replication every week. One individual plant was considered a single replication in each treatment *and* each treatment was replicated 12 times. Thus, there were 3 treatments and 36 plants in this study.

Experiment II

The experiment was performed in a greenhouse at South China Agricultural University (latitude $23^{\circ}09'40''$ N, longitude $113^{\circ}21'28''$ E) in June to October 2015. Each pot was $16 \times 15 \times 15$ cm and contained 800 g of the soil with one banana. This experiment sets up 4 treatments, CS1 and CS2 were respectively applied in

potted plants with inoculating FOC and marked as treatment CS1 and CS2, respectively. Third treatment was with FOC inoculation and without the carbon source as control treatment 1 (CK1). Fourth treatment was without FOC inoculation and the carbon source as control treatment 2 (CK2). Each treatment was repeated 15 times, and one banana was planted in each pot for one repeat, a total of 60 pots were randomly arranged.

The bananas were transplanted into pots on June 19, 2015, and each banana in pot was irrigated once a week with a concentration of 0.4% NPK compound fertilizer (22N:8P:15K) 100ml. CS1 and CS2 aqueous solutions with concentration of 0.01% were respectively irrigated in pots of CS1 and CS2 treatment every 15~20days and CK1 and CK2 without applying carbon source were instead by clean water. The 100ml/pot carbon source solution was irrigated every time until the end of the experiment, a total of 8 times of solution or water was irrigated.

The FOC spore suspension $(10^{7}$ cfu/ml) cultured in advance was diluted with sterile water on July 4, 2015 until the spore concentration was 5×10^{5} cfu/ml (FOC inoculation solution) and then use a syringe respectively inhaled 100ml FOC inoculation solution and directly injected into the soil around banana pseudostem of treatment needed inoculation. For the treatment without inoculation, inject by 100ml water instead.

Sample Collection

Soil Sample Collection

Experiment I. On 14 January 2015, this is, 54 days after inoculated FOC, soil samples were collected from each of the 12 replicates in each treatment. For each treatment, 3 sets of soil samples were created by randomly combining 4 replicates (three soil samples). The soil samples were stored at 4° C for conducting microbial Community Level Physiological Profiling (CLPP) by the Biolog EcoPlate method and the principal components analysis of microbial CLPP was conducted by R software for screening carbon sources of prevention and control FOC.

Experiment II. 107 days after inoculated FOC, soil samples were collected from each of the 15 replicates in each treatment. For each treatment, 3 sets of soil samples were created by randomly combining 5 replicates (three soil samples). The soil samples also were stored at 4° C for conducting microbial CLPP and counting the FOC population.

Determination for Plant Growth Indexes and Collection Samples

Observed the plant height and maximum leaf area of bananas for every ten days in experiment II after inoculated FOC. On the 107th day after inoculation FOC, referenced Carlier's *et al.* (2002) to statistic the incidence and disease index of bananas when the incidence of banana Fusarium wilt was obvious, and harvested bananas and measured their plant biomass, including root dry weight (DW), shoot DW and plant DW.

Plate Count Method

Applied the Microbiological Plate Count Method to determine the FOC population in soil and prepared with the selective medium of FOC according to reference literature (Sun *et al.*, 1978).

Biolog Eco Plate Method

The Biolog technique can be used to investigate the metabolic functional diversity of soil microbes reflected by the utilization ratio of carbon sources (Preston-Mafham *et al.*, 2002). The protocols for creating serially diluted spore suspensions of the soil samples and Biolog EcoPlate method were performed in accordance with Classen's methods (Classen *et al.*, 2010).

Each Biolog EcoPlate consisted of 3 identical sections (3 replicates), each section containing 31 micro wells with distinct dissolved organic carbon substrates and one well with no substrate to serve as negative control (blank; Table 1). For analysis, each plate was treated as 1 individual soil samples for the same treatment was used to fill all wells of the plate. Three replicates in one EcoPlate for each soil samples were detected. The optical densities and Average Well Color Development (AWCD) in the inoculated Biolog EcoPlate was measured at culture times 3, 24, 36, 48, 72, 96, 120, 144 and 168h and using a ELx808-Biolog absorbance reader (BioTek Instruments, USA) under a light wavelength of 590 nm.

Statistical Analyses

Readings from individual plate were inputted into Microsoft Excel for compilation and manipulation. Data from each plate were parsed into 3 data lines for the 3 replicates. For each plate, the Optical Density (OD) of the blank well was subtracted from the OD of each of the 31 wells of its replicate to calculate a net substrate (normalized) OD value (Christian and Lind, 2007). When blanking resulted in negative values, these values were set to zeros for data analysis (Kurten and Barkoh, 2016).

The OD and AWCD of the wells in the Biolog EcoPlate are considered positively associated with the utilization of carbon sources, and reflect the activity of soil microbes. The AWCD conforms to a time-dependent asymptotic sigmoidal curve (Qin *et al.*, 2017). For each replicate reading, the AWCD was calculated as the sum of all normalized OD values divided by 31 (Garland, 1997), as follows:

$$AWCD = \sum (ODi - R)/n,$$

where, ODi represents the OD value of the substrate in each well, R represents the OD of the control (blank), ODi-R is the corrected absorbance value (normalized) and n is the 31 types of carbon substrates (Kurten and Barkoh, 2016).

The Principal Components Analysis (PCA) of the OD values at 72 h of cultivation time was performed with ADE-4 software (R software package) to study the difference of substrate utilization profiles between the inoculation and non-inoculation treatments. A permutation test was used to test the significance of the groupings suggested by the PCA. Values were considered as different when the probability to reject the nil hypothesis was lower than 0.05 (Dai *et al.*, 2004). Duncan's multiple range tests were performed using SAS statistical software.

Results and Discussions

FOC Prevention and Control with Carbon Sources Screening

PCA of soil microbial CLPP was performed to screen the carbon sources for prevention and control FOC. PCA was conducted to understand better the microbial CLPP in the soil microenvironment in response to the different treatments (Hackett and Griffiths, 1997), and microbial utilizing profiles of 31 carbon substrates from different inoculation treatments to analyze associations among these carbon sources and screen the carbon sources (Fig. 1 and Table 1).

To study the effect of C0, C1 and C2 treatments on soil microbial utilizing profiles to the substrates, a PCA

was performed (Fig. 1a and 1b). The results showed that the first principal component (Fig. 1, axis X[PC1]) was mainly and positively determined by the utilization rates of D-Malic acid (30), β -Methyl-D-glucoside (1) and D-Cellobiose (24) etc and mainly negatively determined by the utilization rates of 4-Hydroxy benzoic acid (14), its relative inertia was 59.20%. PC2's relative inertia was 10.85% (Fig. 1, axis Y[PC2]) and the total relative inertia of PC1 and PC2 was 70.05%.

The permutation test of the PCA showed that the differences in the substrate utilization profiles among C0, C1 and C2 were significant (P < 0.0001). The C0 treatment was clearly distinguished from C1 and C2 by their respective substrate utilization profiles, with different utilization rates (Fig. 1b): C0 had higher utilization rates for most carbon sources than did C1 and C2, especially for D-Malic acid, β -Methyl-D-glucoside and D-Cellobiose and their distribution of contribution scores on PC1 were 9274, 9237 and 8907, respectively (Table 1). By contrast, utilization rate for 4-Hydroxy benzoic acid of C0 treatment was considerably lower than those of C2 and C3, and distribution of contribution scores from 4-Hydroxy benzoic acid on PC1 was negative 31 (-31). Therefore, C0 in the positive half of AxisX could be significantly differentiated from C1 and C2 both in the negative half of AxisX, and D-Malic acid (the highest score, 9274) was selected as the carbon source of FOC prevention and control (CS2), 4-Hydroxy benzoic acid (the lowest score, -31) was selected as the carbon source of control (CS1).



AxisX(PC1):59.20%

Fig. 1: PCA of soil microbial CLPP in C0, C1 and C2 treatments. (a) Loading plot of 31 substrates by PCA. (b) PCA scores plot of microbial substrate utilization in different treatments. Note: The key for the identification numbers of the 31 substrates are listed in Table 1. The encircled carbon substrate in red circle was selected as the tested carbon source (CS2)

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Table	 Relative contr 	ibutions of the car	bon substrates on PC1		
96 Eco	Plate cells		Substrate	Substrate ID	Contribution score
A1	A5	A9	None (water blank)	None	None
A2	A6	A10	β-Methyl-D-glucoside	1	9237
A3	A7	A11	D-Galactonic acid γ-Lactone	2	8566
A4	A8	A12	L-Arginine	3	7365
B1	B5	B9	Pyruvic acid methyl ester	4	6270
B2	B6	B10	D-Xylose	5	5145
B3	B7	B11	D-Galacturonic acid	6	6365
B4	B8	B12	L-Asparagine	7	7421
C1	C5	C9	Tween 40	8	649
C2	C6	C10	I-Erythritol	9	2981
C3	C7	C11	2-Hydroxy benzoic acid	10	1278
C4	C8	C12	L-Phenylalanine	11	6908
D1	D5	D9	Tween 80	12	3688
D2	D6	D10	D-Mannitol	13	6143
D3	D7	D11	4-Hydroxy benzoic acid	14	-31 ^b
D4	D8	D12	L-Serine	15	7149
E1	E5	E9	α-Cyclodextrin	16	7053
E2	E6	E10	N-Acetyl-D-glucosamine	17	8602
E3	E7	E11	γ-Hydroxybutyric acid	18	4495
E4	E8	E12	L-Threonine	19	8787
F1	F5	F9	Glycogen	20	8368
F2	F6	F10	D-Glucosaminic acid	21	3553
F3	F7	F11	Itaconic acid	22	5171
F4	F8	F12	Glycyl-L-glutamic acid	23	6481
G1	G5	G9	D-Cellobiose	24	8907
G2	G6	G10	Glucose-1-phosphate	25	8543
G3	G7	G11	α-Ketobutyric acid	26	3902
G4	G8	G12	Phenylethylamine	27	2777
H1	Н5	H9	α-D-Lactose	28	7205
H2	H6	H10	D, L-α-Glycerol phosphate	29	4080
H3	H7	H11	D-Malic acid	30	9274 ^a
H4	H8	H12	Putrescine	31	7098

Note: Each Biolog EcoPlate contains 96 wells consisting of three replicates (first 3 columns) of 31 substrates and 1 water blank. ^a is the carbon substrate that contributed the most on PC1, ^b is the carbon substrate that contributed the least on PC1.

Effects of the Carbon Sources on the Plant Height and Maximum Leaf Area

Plant height and leaf area of banana were important indicators of banana growth. The trend of plant height and maximum leaf area of banana with the extension of inoculation days was shown in Fig. 2a and 2b. The plant height of each treatment increased with the growth time and the plant height of CS2 treatment was significantly higher than that of CS1, CK2 and CK1 treatments during the whole culture period (Fig. 2a). The bananas treated by CS2 were respectively higher than CS1, CK2 and CK1 3%, 5% and 11% at the end of the experiment (107th day after inoculation).

The maximum leaf area of each treatment increased rapidly with the growth time for 57days after inoculation (Fig. 2b). However, the maximum leaf area of the inoculated banana (CS1, CS2, CK1) grew slowly or stopped after 57days of inoculation, but the leaf area of the non-inoculated banana (CK2) still increased rapidly. Because the inoculated bananas did not get sick or the disease was relatively mild in first 57days and the growth of banana leaves was not seriously affected. However, the banana began to get sick and the disease seriously affected the growth of banana leaves after 57days. From 57th day to the end of the experiment (107th day), the area of banana leaves without inoculated (CK2) was significantly larger than that inoculated, while that inoculated with CS2 was significantly larger than that inoculated with CS1 and CK1. In conclusion, the growth of bananas treated with carbon source CS2 and those not inoculated (CK2) were better than those treated with other treatments, indicating that the treatment with carbon source CS2 was beneficial to the growth of bananas in the diseased soil and the effect of promoting the growth of bananas was better than that with carbon source CS1.

Effects of the Carbon Sources on Banana Biomass

As shown in Table 2, there was no significant difference in banana root DW among treatments, however, shoot DW and plant DW of bananas of CS2 and CK2 were significantly larger than those of CS1 and CK1, the plant DW of CS2 was respectively larger than

CS1 and CK1 16.82% and 15.90%. These results indicated that carbon source CS2 could promote banana growth and improve biomass in the disease soil, but carbon source CS1 could not. This may be because the application of carbon source CS2 can effectively reduce the incidence and harm of bananas wilt and contribute to the healthy growth of bananas.

Effects of the Carbon Sources on Wilt Infection and Disease Index of Fusarium wilt in Banana Plants

Wilt Infection and disease index of bananas can directly reflect the disease incidence of bananas (Carlier *et al.*, 2002). As shown in Fig. 3, the bananas of non-inoculated treatment (CK2) did not infect disease at the end of the pot experiment, therefore wilt infection and disease index was significantly smaller than other inoculation

treatments. Wilt infection of bananas treated with CS2 was significantly lower than those of CS1 and CK1 during inoculation treatment, which respectively lowers than CS1 and CK1 40 and 46 percentage points. The disease index of bananas treated with CS2 was also significantly lower than that of CK1, lower than that of CK1 70%, and the difference between CS1 and CK1 was not significant. As a result, the bananas treated with CS2 were less infectious than CS1 and CK1, this indicates that the application of CS2 can reduce the occurrence of banana Fusarium wilt, which may be because the CS2 improved the microbial structure by regulating the metabolic activity of soil contributed microorganisms and to beneficial microorganisms to inhibit the growth of FOC, thus decreased FOC population and reducing the incidence of Fusarium wilt (Wu bananas et al., 2009).



Fig. 2: (a) Effects of carbon sources on the plant height; (b) Effects of carbon sources on the maximum leaf area



Fig. 3: Effects of carbon sources on wilt infection and disease index of Fusarium wilt

Table 2: Effects of the carbon sources on banana dry weight (g)

Treatment	Shoot DW	Root DW	Plant DW
CK1	18.65±1.16b	1.68±0.10a	20.32±1.21b
CS1	18.75±0.54b	1.41±0.11a	20.16±0.61b
CS2	21.94±0.46a	1.61±0.11a	23.55±0.48a
CK2	21.24±0.52a	1.55±0.09a	22.80±0.52a

Note: Number in the table is mean \pm standard deviation. Different letters in each column meant significant differences among treatments (P<0.05).

Effects of the Carbon Sources on the FOC Population in Soil

The FOC population is directly related to the incidence of banana Fusarium wilt (Sun et al., 1978). In general, the more FOC population in soil, the more serious the incidence of banana disease, therefore, it is of great significance to analyze the FOC population in different treatments soil for the study of carbon sources prevention and control of banana Fusarium wilt. As shown in Fig. 4, FOC population in soil without inoculated (CK2) was significantly smaller than that in other inoculated. In the inoculation treatment, the FOC population in the soil treated with CS2 was significantly lower than that treated without carbon source (CK1), and respectively less than that treated with CK1 and CS1 59% and 48%, while the difference between CS1 and CK1 was not significant. This result verified that carbon source CS2 could prevent and control the occurrence of banana Fusarium wilt by reducing the FOC population in soil, because CS2 could provide preferred carbon source nutrients for soil microorganisms, improved their overall activity and diversity, and thus inhibited the growth of FOC (Lee, 2010).



Fig. 4: Effects of carbon sources on the FOC population



Fig. 5: AWCD variation of total carbon substrates with cultivation time prolonged

Effects of the Carbon Sources on AWCD Variation of Total Carbon Substrates

The AWCD can reflect the carbon utilization profile and activity of soil microbes (Qin *et al.*, 2017). In general, the AWCD in this study increased in "S" curve during cultivation (Fig. 5). The AWCD curves of 4 treatments showed that carbon utilization increased slowly in the first 24h, rapidly from 24 to 72h *and* was fairly stable from 72 to 168h. An analysis according to the different treatments showed that carbon utilization in soil treated with CS2 was significantly higher than that of CS1, CK1 and CK2 in interval of 48h to 168h, and the carbon utilization in CK2 soil was significantly higher than that of CS1 and CK1 soils. This result means that the activity of soil microbes is lowered by the presence of FOC (CK1), but is improved with the use of carbon source CS2. This result may be due to the FOC is the main dominant population under the inoculation condition and it will inevitably compete with other microorganisms for the habitats such as carbon sources, thus resulting in insufficient nutrition and space for the growth of other microorganisms. It is also possible that the large amount of FOC produces a substance that inhibits the growth of other microorganisms (Huang *et al.*, 2015). And applying CS2 could provide carbon nutrient preferred by most microorganisms in the soil for the growth of microorganisms (Fig. 1), thus improved the overall activity of soil microbes (Lee, 2010).

Conclusion

In this paper, studied carbon source screening technology for the prevention and control of banana Fusarium wilt, and screened carbon source CS2 to control of banana wilt according to this technology. This carbon source could effectively improve the overall activity of microorganisms in pathogenic soil, and the AWCD treated with CS2 was significantly higher than that treated with other treatments. Meanwhile, CS2 could significantly reduce the FOC population in soil, the FOC population treated with CS2 was respectively lower than that treated with control carbon source (CS1) and without carbon source (CK1) 48% and 59%. Therefore, the banana wilt infection of CS2 was significantly smaller than that of CS1 and CK1, which respectively lower than CS1 and CK1 40 and 46 percentage points. The disease index of bananas wilt treated by CS2 was also significantly lower than that of CK1, which lower than that of CK1 70%, and the difference between CS1 and CK1 was not significant. Why could CS2 reduce the FOC population, wilt infection rate and disease index of bananas? Those details were not covered in this paper and will be covered in a future study.

Due to apply CS2 could effectively prevent and control the occurrence of banana Fusarium wilt and reduce the damage of Fusarium wilt to bananas, the CS2 could promote the growth and biomass accumulation of bananas in the pathogenic soil. The results showed that the growth vigor of bananas treated with CS2 was significantly better than that of CS1 and CK1 and the shoot DW and plant DW of bananas were significantly larger than those of CS1 and CK1. The plant DW of CS2 was respectively larger than that of CS1 and CK1 16.82% and 15.90%. In conclusion, this paper studied the carbon source screening technology according to microbial CLPP in different soil, and this screening technology can be used to screen out high-quality carbon sources for the prevention and control of banana wilt, which has great significance to the production or selection of organic fertilizers containing high-quality carbon sources and the improvement of prevention effect of biological organic fertilizers to banana wilt.

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Author's Contributions

Jin Li and Tingting Duan: Designed and performed the experiments, analyzed the data and prepared the paper. Both authors contributed equally to this work.

Zhong Lin and Chao Zheng: Participated to collect the materials related to the experiment.

Xiaolin Fan: Designed the experiments and revised the manuscript.

Ethics

The authors declare their responsibility for any ethical issues that may arise after the publication of this manuscript.

Conflict of Interest

The authors declare that they have no competing interests. The corresponding author affirms that all of the authors have read and approved the manuscript.

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