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## SINGLE SPORE ISOLATION AND MOLECULAR CHARACTERIZATION OF TRICHODERMA SPECIES FROM AGRICULTURAL FIELDS OF ZHEJIANG PROVINCE, CHINA

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### ABSTRACT

It is well known that *Trichoderma* spp. can be used as a biological control agent against variety of plant pathogens. In this study, *Trichoderma* species were isolated from rhizosphere samples of various sites in Zhejiang province, China. A total of 41 soil samples were processed for the isolation of single cell culture of *Trichoderma* sp. Out of which, 32 species of the genus *Trichoderma*, were isolated and characterized by using single spore isolation method. The isolates were subjected to sequencing, for precise identification up to species level. *Trichoderma harzianum* was isolated from maximum (09) number of soil samples, followed by *T. velutinum*, isolated from 3 soil samples. The other isolated species of the genus *Trichoderma* were; *lixii*, *atroviride*, *koningii*, *longibrachatum*, *rossicum*, *gamsii*, *koningiopsis*, *tomentosum*, *afroharzianum*, *viride*, *citroviride*, *aureoviride*, *simmonsii* and *hamatum*. It was concluded that *Trichoderma* sp. were abundantly found in the agricultural soils of Zhejiang province, China.

**Keywords:** *Trichoderma*, Biocontrol, Fungi, Rhizosphere, single spore culture

### INTRODUCTION

*Trichoderma* is a common filamentous imperfect fungus that belongs to class Deuteromycetes, and family Dematiaceae. It is a common saprophytic fungus in the rhizosphere and found in almost any soil. *Trichoderma* species are cosmopolitan fungi, frequently present in all types of soils (Alexander, 1961). *Trichoderma* spp. are free living fungi that are common in soil and root ecosystems and promote plant growth (Yedidia *et al.*, 2001). The importance of this genus is evident from its biocontrol potential against several soil borne plant pathogens (Papavizas 1985; Elad *et al.* 1993; Elad 2000; Freeman *et al.* 2004; Dubey *et al.* 2007; Windham *et al.* 1989; Sharon *et al.* 2001). The growth of plants with *Trichoderma* spp. alone or in combination with

pathogen was greater than in plants inoculated with pathogen alone (Sreedevi *et al.*, 2011). *Trichoderma* has recently been reported as an effective entomopathogenic fungus against *Bemisia tabaci* (Waheed *et al.*, 2016)

The process of fungal identification does not rely solely on morphological features for identification. The use of fungal cultures, obtained from single spore isolations is fundamental to the identification of many fungi. The importance of careful identification of fungi and obtaining cultures that have been isolated from single spores cannot be over stressed (Smith, 1969). Isolation methods must be simple to perform, even for those who do not have experience in isolation, relatively inexpensive and effective. Simple, inexpensive and effective methods have been described to isolate fungi from single spore cultures (Choi *et al.*, 1999). The phylogenetic species concept, which can include comparison of both morphological and molecular characters, in most cases require single spore cultures (Goh and Hanlin, 1997).

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For the storage of fungal pure culture, it can be maintained on an agar slant at 4°C or stored as a few pieces of agar with mycelia in cryo vials with 10% glycerol in liquid nitrogen. Alternatively, it can be stored as a few pieces of agar with mycelia in small bottle with sterilized paraffin oil or sterilized water at 4°C. There are several other methods to maintain fungal cultures and as reviewed in the "Preservation and Maintenance of Living Fungi" (Smith and Onions, 1994).

The present research was mainly focused on the single spore isolation and molecular characterization of *Trichoderma* sp. from soil of Zhejiang province, China.

#### MATERIALS AND METHODS

**Soil Sampling:** Soil samples were collected from various ecological habitats of cultivated vegetable crops of Zhejiang province, China for the isolation of *Trichoderma* spp. Samples were stored at 4°C until further used.

**Isolation of *Trichoderma* species:** For the isolation of single spore *Trichoderma* strains, a serial dilution technique was followed. Soil suspension was prepared from each sample by adding 1g of soil into 100ml of sterile distilled water, serial dilutions were prepared and 0.5ml from 10<sup>-4</sup> dilution of each soil suspension was pipetted onto water agar (WA) plates. It was gently spread over WA with a glass spreader and incubated at 28°C. The culture plates were examined daily and the individual colony was isolated and sub cultured onto fresh Potato Dextrose Agar (PDA) plates. The pure cultures of *Trichoderma* sp. were stored in 20% glycerol at -80°C for long time storage.

**Molecular Characterization of *Trichoderma* strains:** For the molecular characterization of *Trichoderma* strains, DNA was isolated from the pure culture, amplified through PCR and subjected for sequencing of the amplified product.

**Extraction of fungal DNA:** Isolation of fungal DNA is a multi-step method. It includes growing the fungus in liquid or solid medium, lyophilizing the fungal samples, disrupting cell wall, removing proteins with phenol and chloroform, and precipitating DNA with ethanol or Isopropanol. In this experiment, DNA from fungal mycelia was isolated by adopting a modified technique, as reported by (Saitoh *et al.*, 2006; Tripathy *et al.*, 2017). Steps involved in DNA extraction were as follows:

- Took 700 ul extraction or lyses buffer (Tris-HCl, EDTA, 150 mM NaCl, 1% SDS, pH 8.0) in 1.5 ml Eppendorf tube.

- A small amount from 10-day old fungal culture was added into the buffer, by using a sterile toothpick.
- Quartz sand was added into the culture and the tubes were subjected to tissue lyses at 65,000 Hz for 2 min.
- Centrifuge at 12,000 rpm for 10 min.
- Discarded pellet with sand particles and took 650 ul supernatant into tubes.
- Added 650 ul Chloroform (CIA) to the tubes and shake at 150 rpm for 15 min.
- Centrifuge at 12,000 for 10 min.
- Remove 550 ul of the supernatant into new tubes and added 550 ul of Isopropanol left for 30 min.
- Centrifuge at 12,000 for 10 min.
- Discarded the supernatant and collected the pellet.
- Washed DNA pellet twice with 70% ETOH (500 ul) and air dried.
- Added 50 ul Eluent and left for 10 min before PCR.

**DNA Amplification by PCR:** For DNA amplification, first PCR Mix was prepared, which contained 15ul Taq mix, 1 ul Primer-1, 1ul Primer-2, 1ul fungal DNA, 12 ul ddH<sub>2</sub>O, making it 30ul volume. The PCR protocol, given by (Lee and Taylor 1990) was used for the fungal DNA amplification. The PCR reactions involved 1 cycle at 95°C for 5 min, followed by 34 cycles with a denaturation step at 95°C for 30 sec, an annealing step at 57°C for 30 sec, and an extension step at 72°C for 1 min, Hold: at 4° C.

**Gel Electrophoresis:** After the DNA amplification, the PCR product was subjected for gel electrophoresis. To 25 µl of amplification products obtained after the PCR, 2 µl of loading dye (bromophenol blue) were added and loaded into individual wells of 1.0% Agarose in 1 x TBE buffer. Electrophoresis was carried out at 100 V for 2 h, and thereafter the gel was stained with ethidium bromide (10mg µl<sup>-1</sup>).

Put 1% Agarose gel in a gel tank and Ethidium bromide dye was added into it. A comb was placed gently and allowed the gel to solidify. Removed comb carefully and placed the gel in a gel tank. Loaded 7 ul PCR products into each well and a ladder DNA was loaded to a side. Run the gel at 110A, 400V for 20 min. Detection of DNA was made on a trans-illuminator under UV light. The DNA samples with clear bands were sent to the company for sequencing. After getting the sequencing data, a tree was prepared to check the similarity index among the fungal isolates.

**Storage of Cultures:** The pure culture was maintained on an agar slant at 4°C or stored as a few pieces of agar

with mycelia in cryo vials with 10% glycerol in liquid nitrogen.

## RESULTS AND DISCUSSION

Biological control of plant pathogens by microorganisms has been considered as a natural and environmentally acceptable alternative to the existing chemical treatment methods (Baker and Paulitz, 1996). Intensified use of fungicides resulted in the accumulation of toxic compounds potentially hazardous to humans and environment (Cook and Baker, 1983).

*Trichoderma* is an important filamentous fungus, which has recently gained immense importance due to its biological control ability against several plant pathogens, in decomposition of organic wastes, production of enzymes like xylanases, cellulases. In the present study, single spore cultures of *Trichoderma* isolates were subjected for the molecular identification of different *Trichoderma* species isolates present in different fields of Zhejiang province, China. Ranga *et al.*, (2017) studied the genetic diversity of *Trichoderma* sp. from Rhizosphere regions of different cropping systems using RAPD markers. Results show that 32 species of *Trichoderma* have been isolated from agriculture soil of various vegetable fields in Zhejiang province, China (Table 1). *Trichoderma harzianum* was isolated from maximum (09) number of soil samples, followed by *T. velutinum*, isolated from 3 soil samples. *Trichoderma harzianum* has been reported as the most among common, among the biological control agents of the genus *Trichoderma* (Hermosa *et al.*, 2000). Five *Trichoderma* species i.e. *lixii*, *atroviride*, *koningii*, *longibrachiatum* and *rossicum* were isolated from the samples 607, 47, 252, 13 and 561 respectively. The other *Trichoderma* sp. like *gamsii*, *koningiopsis*, *tomentosum*, *afroharzianum*, *viride*, *citroviride*, *aureoviride*, *simmonsii* and *hamatum* were isolated from the soil samples, designated as 498, 505, 584, 10, 52, 542, 51, 531 and 494, respectively as shown

in Table 1. The Agarose gel electrophoresis of PCR products of ITS region is shown in Figure 1. A Dendrogram, depicting variation among isolates of *Trichoderma* spp., is given in Figure 2.

Table 1. List of species of *Trichoderma* sp., collected from various agricultural fields in Zhejiang province, China

S.No.	Sample	<i>Trichoderma</i> Species
1	498	<i>Trichoderma gamsii</i>
2	505	<i>Trichoderma koningiopsis</i>
3	584	<i>Trichoderma tomentosum</i>
4	406	<i>Trichoderma harzianum</i>
5	33-2	<i>Trichoderma velutinum</i>
6	607	<i>Trichoderma lixii</i>
7	47	<i>Trichoderma atroviride</i>
8	540	<i>Trichoderma harzianum</i>
9	10	<i>Trichoderma afroharzianum</i>
10	52	<i>Trichoderma viride</i>
11	252	<i>Trichoderma koningii</i>
12	33-1	<i>Trichoderma velutinum</i>
13	53	<i>Trichoderma koningii</i>
14	13	<i>Trichoderma longibrachiatum</i>
15	542	<i>Trichoderma citrinoviride</i>
16	561	<i>Trichoderma rossicum</i>
17	51	<i>Trichoderma aureoviride</i>
18	531	<i>Trichoderma simmonsii</i>
19	495	<i>Trichoderma lixii</i>
20	17	<i>Trichoderma harzianum</i>
21	33	<i>Trichoderma harzianum</i>
22	37	<i>Trichoderma harzianum</i>
23	637	<i>Trichoderma harzianum</i>
24	33-1	<i>Trichoderma velutinum</i>
25	492	<i>Trichoderma harzianum</i>
26	494	<i>Trichoderma hamatum</i>
27	36	<i>Trichoderma longibrachiatum</i>
28	35	<i>Trichoderma harzianum</i>
29	6	<i>Trichoderma atroviride</i>
30	353	<i>Trichoderma harzianum</i>
31	26	<i>Trichoderma rossicum</i>
32	541	<i>Trichoderma stromaticum</i>

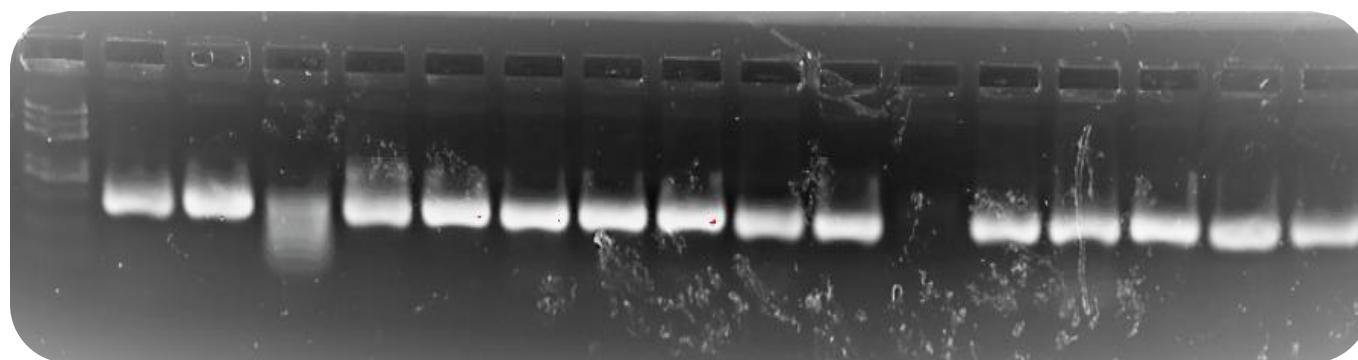


Figure 1. Agarose gel electrophoresis of PCR products of ITS region.

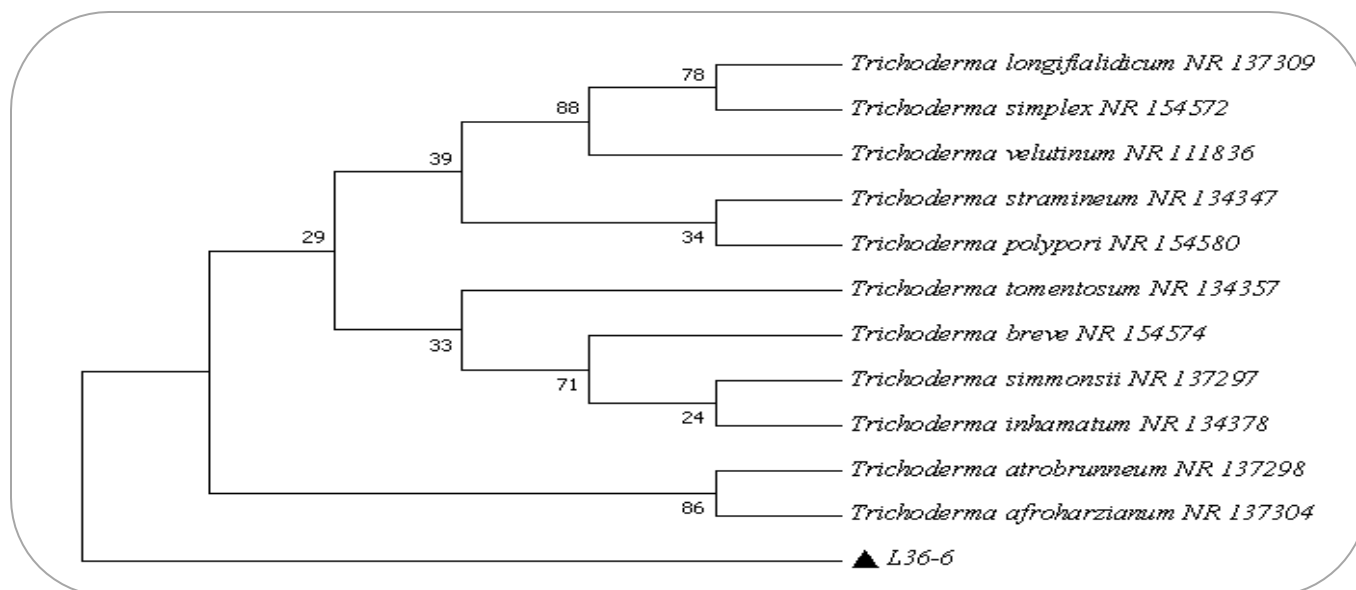


Figure 2. Dendrogram, depicting variation among isolates of *Trichoderma* spp.

#### CONCLUSION AND RECOMMENDATIONS

Single spore cultures of *Trichoderma* isolates were subjected for the molecular characterization of *Trichoderma* sp., collected from the agriculture fields of Zhejiang province, China. Out of the 32 isolated species of *Trichoderma*, 9 were reported as *T. harzianum*, 3 were *T. velutinum*. The biocontrol potential of these isolates of *Trichoderma* against various pathogens needs to be investigated.

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