

The Construction of Infectious Tomato Yellow Leaf Curl Virus(TYLCV) and the Analysis of Resistance on Transgenic ...

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ABSTRACT

Tomato yellow leaf curl virus(TYLCV) is a member of the Begomovirus in the Geminiviridae family. TYLCV contains a monopartite genome and is transmitted by whitefly *Bemisia argentifoli* and results in severe losses in tomato productions. In this study, naturally infected tomato samples were collected from eight tomato fields in Changhua County and polymerase chain reaction (PCR) were used for the detection of TYLCV using primers pair that is specific to TYLCV genome. Five in the forty samples from Sihu and one in eight samples collected from Yongjin, Dacun, and Yuanlin, respectively, showed positive in PCR analyses. The average infection rate is 12.5%. The PCR products were cloned and sequenced. Sequences alignment with the C1 gene and the full-length genome revealed that TYLCV Sihu isolate is most closely related to a TYLCV Thailand strain, with the nucleotide sequence identity of 98%. By the concept of parasite-derived resistance (PDR), a part of TYLCV replicase gene (C1) was used to engineer into three different transgene constructs for plant transformation, which include the TYLCV C1 gene arranged in a sense, antisense and double antisenses manners. These three constructs were transformed into tobacco (*Nicotiana benthamiana*) by *Agrobacterium tumefaciens*. The putative transgenic tobacco plants were selected in Kanamycin medium and confirmed by PCR with the C1 and NPTII specific primer pairs. Ten to fifteen putative transgenic lines were obtained from the transformed of each sense, antisense, and double antisenses constructs. Since it was not possible to analyses the resistant ability of the TYLCV transgenic tobacco by mechanical inoculation of natural TYLCV. An infectious TYLCV clones was constructed. Three overlapping DNA fragment that covered the full-length of TYLCV genome of Sihu isolate were obtained from PCR amplification. The TYLCV clone contains approximately 3.5 Kb that is about 1.27 times longer than virus genome unit size were inoculated into non-transgenic tobacco by argoinoculation. The mild mosaic and less leaf curve symptom were observed 14 days post inoculation. The ability of the infectivity in the inoculated plants were further confirmed by PCR amplification and sequencing. The TYLCV infectious clone was then used as a virus source for challenge into transgenic tobacco and the resistant ability of the transgenic is under investigation. The study of the resistant ability of TYLCV transgenic tobacco can provide the information for the engineering of TYLCV transgenic tomato in the future.

Keywords : Tomato yellow leaf curl virus(TYLCV) ; transformation ; *Agrobacterium tumefaciens* ; polymerase chain reaction

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