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## Isolation and characterization of virulence-deficient mutants Xanthomonas oryzae pv. oryzae by transposon mutagenesis

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## **Objectives**

This study was to construct a mutant pool of *Xanthomonas oryzae* pv. *oryzae* (KACC10331) by transposon mutagenesis. Virulence-deficient mutantsof Xoo were to isolate and characterize

## Materials and Methods

- 1. Materials: Xanthomonas oryzae pv. oryzae strain- KACC10331 (K-1 race), Transposon mutagenesis: Electroporation using Transposome
- 2. Methods: Tn5 mutants of Xoo were inoculated in susceptiblerice variety, Milyang 23.

## Results and Discussion

Xanthomonas oryzae pv. oryzae(KACC10331) is causal agent of bacterial leaf blight on rice. Recently, whole genome sequence of the Xoo strain was determined and the functional studies of Xoo have been done on basis of the sequencing information. In this study, transposon was randomly introduced into the genome to identify virulence genes in X. oryzae pv. oryzae genome (KACC10331). Transposon mutant clones of Xoo were selected from media containing Kanamycin and inoculated on rice variety, Milyang 23 which is susceptibleto Xoo for investigating their pathogenicity. Of primary screened 24,540 Xoo-mutant strains, 542 mutant strains showed intermediate or non-pathogenic feature. Furthermore, ideal non-pathogenic mutant strains were finally selected by second and third pathogenic screening. Insertion copy numbers of transposon in the mutant genomes were analyzed by Southern blot analysis. The transposon insertion sequences in the mutant genomes were determined by direct sequencing based on Tailing-PCR technique.

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