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## Phylogenetic analysis by RAPD analysis of Isolates of *Fusarium oxysporum* formae speciales in Korea

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The goal of this study was to determine that the phylogenetic analysis of *Fusarium oxysporum* and formae speciales. Identification of *Fusarium oxysporum* and formae speciales has been difficult due to confusing phenotypic classification systems. The genetic relationship of forty-one formae speciales of *Fusarium oxysporum* causing plant disease was determined by Random amplified polymorphic DNA (RAPD) markers. A high level of genetic variation was found in *F. oxysporum* and formae speciales. Clustal G analysis of RAPD did not show any specific classification to explain phylogenetic relation of *F. oxysporum* and formae speciales. In phylogenetic analysis, *F. oxysporum* and formae speciales formed a short cluster, whereas *F. oxysporum* f. sp. *radicus-lycopersici* 2 strains, *F. oxysporum* f. sp. *niveum* occupied different positions in the trees.