

## Construction of transgenic *citrus* plants by introduction of fatty acid desaturase genes

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### Abstract

Three kinds of *citrus*, 'Yoshida' navel orange (YN), 'Cara Cara Red' navel orange (CCRN), and kumquat were transformed with fatty acid desaturase genes, *fad3* and *fad7*. To introduce DNA the microprojectile bombardment system was used for the stem microsections of kumquat. The transformants were screened on the selective medium containing kanamycin, and then DNA's of the transformants were analyzed by polymerase chain reaction. Some embryogenic clones were obtained from the transformed callus cells of YN and CCRN. Some planted were regenerated from the kumquat microsections transformed with *fad7*.

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### Introduction

The fatty acid composition of cell membrane is well known to be closely related with the tolerance of plants to low temperature. In general plants with higher content of polyunsaturated fatty acids are more tolerant to cold stress.

The enzymes, fatty acid desaturases, catalyze the process which introduce double bonds into fatty acids. Many reports showed that modifications of the genes encoding these enzymes altered the composition of unsaturated fatty acid in membranes and changed low-temperature resistance of plants (Ishizaki - Nishizawa et al., 1996).

Recently the composition of unsaturated fatty acids in membrane lipids was reported to be important in growth,

respiration and photosynthesis as well as low-temperature resistance of plants (Tasaka et al., 1996). Therefore introduction of desaturase genes might improve fruit qualities via enhancing these physiological activities of plants especially at low temperature.

In this study we tried to transform *citrus* species with fatty acid desaturase genes for improving fruit qualities.

### Materials and methods

The embryogenic callus cells of 'Yoshida' (YN) and 'Cara Cara Red' (CCRN) navel orange (*Citrus sinensis*), and the stem microsections of kumquat (*Fortunella japonica*) were used for transformation.

Two kinds of fatty acid desaturase genes, *fad3* and *fad7*, were used. Each gene was constructed in a plant expression vector, pBI121, harboring nptII.

The PDS-1000 He Biolistic Particle Delivery System (Bio-Rad) was used for introducing DNA into callus cells of YN and CCRN orange. *Agrobacterium* system was used for kumquat stem microsections.

Transformants were screened on a selective MT medium containing 100mg/l of kanamycin. The introduced DNA of each gene in transformants was analyzed by PCR.

### Results and discussion

The embryogenic callus cells of YN and CCRN were transformed by microprojectile bombardment system. For the transformation the DNA of *fad3* or *fad7* reconstructed in pBI121 was coated on M17 tungsten particles. The cells were bombarded with the particles at 1100 psi of helium gas pressure, 1/4" of gap distance and 7.0cm of target distance. The bombarded cells were transferred to the selective medium containing 100mg of kanamycin after two days stabilization. The cells growing continuously on the kanamycin medium were obtained. Fig. 1 showed the selective growth of YN cells transformed with *fad7*. We could obtain some cells of YN and CCRN in the early stage of embryogenesis. Fig. 2 showed the embryogenic

cells of YN transformed with *fad7*.



Fig. 1. Selective growths of YN callus cells on MT medium containing kanamycin after bombardment with pBI121 containing both of *nptII* and *fad7* gene.

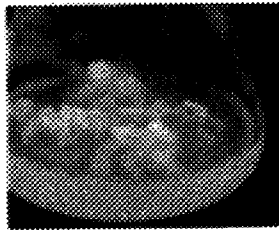


Fig. 2. The transformed YN cells under somatic embryogenesis.

The genomic DNA extracted from the selected callus clones was amplified by PCR with the specific primers for *nptII* or target genes, *fad3* and *fad7*, to confirm that the clones contain the introduced DNA. Fig. 3 showed that the DNA bands of 1.0 kbp for *fad7* were observed in two clones (lane 8 and 9) among five from the YN cells bombarded with pBI121 harboring *nptII* and *fad7*. These two clones also showed the 795 bp DNA bands for *nptII* (lane 13 and 14). Therefore these two clones were appeared to contain both genes of *nptII* and *fad7*. Five clones from the cells bombarded with pBI121 harboring *nptII* and *fad3* were also tested for *fad3* DNA. The two clones showed both DNA of *nptII* and *fad3*.

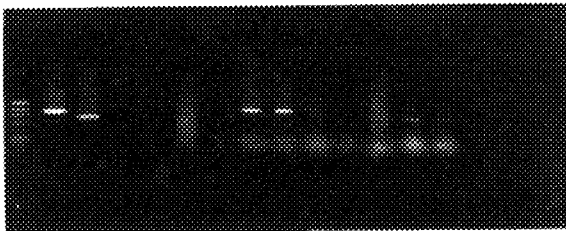


Fig. 3. PCR-amplified DNA's of *nptII* and *fad7* gene from the clones bombarded with pBI121 containing both of *nptII* and *fad7*.

lane 1: marker, 2: positive control (*fad7*), 3: positive control (*nptII*), 4: negative control ( $-fad7$ ), 5: negative control ( $-nptII$ ), 6 and 7: non-bombarded clones, 8-12: *fad7* amplification in bombarded clones, 13-17: *nptII* amplification in bombarded clones.

For the transformation of kumquat, stem microsections were used. The microsections were prepared as 400  $\mu$ m thickness by vibrating microtome. The DNA of pBI121 harboring *nptII* and *fad7* were introduced by *agrobacterium* transformation system. The transformed microsections were cultured on the MT medium containing 2mg/l of BA, 0, 2mg/l of NAA and 50mg/l of kanamycin. We could obtained some plantlets from the kumquat microsections (Fig. 4).

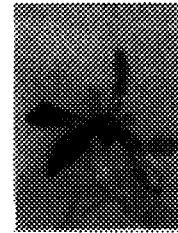


Fig. 4. A plantlet derived from the microsection of kumquat transformed with *fad7*.

## References

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