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# Production of intergeneric somatic hybrids between round kumquat (Fortunella japonica Swingle) and 'Morita navel' orange (Citrus sinensis Osbeck)

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Abstract Intergeneric somatic hybrids between embryogenic callus-derived protoplasts of round kumquat (Fortunella japonica Swingle) and 'Morita navel' orange (Citrus sinensis Osbeck) were produced by electrofusion. Among the eight different fusion strains obtained, six showed normal morphology, whereas the remaining two showed malformation. All the regenerated plants were intermediate in leaf morphology and had thick and round leaves, which are typical characteristics of polyploids. Ploidy analyses by flow cytometry and chromosome counting in root-tip cells revealed that these plants are amphidiploid  $(2n=4\times1=36)$ . Hybridity of the fusion products was confirmed by random amplified polymorphic DNA and cleaved amplified polymorphic sequence (CAPS) analyses. Furthermore, analyses of chloroplast (cp) and mitochondrial (mt) DNA by CAPS showed that these somatic hybrids contained cp- and mt-DNA of round kumquat without recombination in the regions analyzed.

Keywords Citrus · Fortunella · Protoplast fusion · Universal primers · Cytoplasmic DNA

Abbreviations BA: 6-Benzylaminopurine · CAPS: Cleaved amplified polymorphic sequence · RAPD: Random amplified polymorphic DNA

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# Introduction

The kumquat (Fortunella), a close relative of Citrus, consists of a few species and is primarily grown in China and Japan. As well as being an ornamental plant, kumquat fruit is used for preserve, candy, syrup and fresh consumption because of its special flavor and thick edible rind (Hodgson 1967). In the southern part of Japan, kumquat is one of the most important fruit trees and is a regional specialty product crop. However, breeding of kumquat has lagged far behind that of Citrus species, and only one cultivar—Meiwa kumquat (Fortunella crassifolia Swingle)—is under cultivation. Production of new cultivars with unique characteristics such as seedlessness and attractive flavor is desired.

Somatic hybridization can overcome barriers of sexual incompatibility to allow production of hybrids between distantly related species, offering the chance to generate novel cultivars possessing traits from both parents. In Citrus and related species, somatic hybridization techniques are well developed and have been employed to facilitate improvement of Citrus scion and root stock (Grosser and Chandler 2000; Grosser et al. 2000). To date, more than 150 citrus somatic hybrids have been produced throughout the world, including intergeneric combinations (Moreira et al. 2000; Guo et al. 2002; Liu et al. 2002). However, only a few somatic hybrids have been produced between Fortunella and Citrus species, i.e., Meiwa kumquat with sweet orange (Deng et al. 1992; Shi et al. 1998; Cheng et al. 2003), Meiwa kumquat with 'Succari' sweet orange (Grosser et al. 1996), round kumquat with Mediterranean mandarin (Ollitrault et al. 1996), and Kinzu kumquat with trifoliate orange (Miranda et al. 1997). In these examples of intergeneric somatic hybrids between Fortunella and Citrus species, abnormal growth was reported, the reasons for which are still unclear due to the limited number of examples. Cheng et al. (2003) reported that nuclear-cytoplasmic incompatibility might be the cause of such abnormal growth.

In the present study, we describe regeneration of intergeneric somatic hybrids produced by electrofusion between embryogenic callus-derived protoplasts of round kumquat (Fortunella japonica Swingle) and 'Morita navel' orange (Citrus sinensis Osbeck) with male and female sterility, and discuss possible reasons for abnormal growth of these intergeneric somatic hybrids.

## Materials and methods

## Plant materials

Round kumquat (F. japonica Swingle) and 'Morita navel' orange (C. sinensis Osbeck) were used in this study. To obtain embryogenic calli from these species, mature fruits were surface-sterilized and then cut open aseptically. Unfertilized ovules were carefully dissected and placed on MT (Murashige and Tucker 1969) medium containing 5 mg l<sup>-1</sup> adenine, 500 mg l<sup>-1</sup> malt extract, 30 g l<sup>-1</sup> sucrose and 2 g  $I^{-1}$  gellan gum (Kunitake et al. 1991). White, friable nucellar calli, with high embryogenic ability, were induced after approximately 3 months of culture. Embryogenic calli of both species were maintained for 3 years by subculturing on MT medium containing 10 mg  $1^{-1}$  6-benzylaminopurine (BA), 30 g  $1^{-1}$ sucrose and 2  $g$  l<sup>-1</sup> gellan gum at 25 $^{\circ}$ C under continuous illumination (38  $\mu$ mol m<sup>-2</sup> s<sup>-1</sup>).

#### Protoplast isolation

Protoplasts were isolated from embryogenic calli of 'Morita navel' and round kumquat according to the method of Kunitake et al. (2002). Prior to protoplast isolation, embryogenic calli of both species were pretreated for 10 days by transferring to MT medium containing  $50 \text{ g } l^{-1}$  lactose and  $2 \text{ g } l^{-1}$  gellan gum to prevent bursting of protoplasts. The pretreated calli were gently squashed and incubated with an enzyme solution containing 0.3% (w/v) Cellulase Onozuka R-10 (Yakult, Tokyo, Japan), 0.3% (w/v) Macerozyme R-10 (Yakult), 0.1% Driselase (Kyowa, Tokyo, Japan), one-half strength MT macro elements and 0.7 M sorbitol, on a rotary shaker (70 rpm min<sup>-1</sup>) for 16 h at 25°C to liberate protoplasts. Protoplasts were purified by filtration through Miracloth (Calbiochem, San Diego, Calif.) and washed twice with 0.6 M mannitol solution after centrifugation (800 rpm, 3 min).

## Electrofusion and protoplast culture

Isolated protoplasts were fused by electrofusion according to Kunitake et al. (2002). After fusion, protoplasts were cultured in MT medium containing 0.6 *M* sucrose and 2 g  $1^{-1}$  gellan gum, according to Ohgawara et al. (1985). After 5 months of culture, several green somatic embryos were transferred onto MT medium containing 500 mg  $l^{-1}$  malt extract, 30 g  $l^{-1}$  sucrose and 2 g  $l^{-1}$ gellan gum. After a further 1 month, mature somatic embryos were transferred onto one-half strength MT medium containing  $1 \text{ mg } 1^{-1}$ gibberellic acid (GA<sub>3</sub>), 10 g l<sup>-1</sup> sucrose and 3 g l<sup>-1</sup> gellan gum for germination.

## Flow cytometry

Young leaves of approximately 1  $\text{cm}^2$  were collected from each of the parents as well as the regenerated plants, and chopped with a razor blade. The leaf samples were treated for 5 min in 1 ml buffer solution containing  $1.0\%$  (v/v) Triton X-100, 140 mM mercaptoethanol, 50 mM Na<sub>2</sub>SO<sub>3</sub> and 50 mM Tris-HCl at pH 7.5, according to the method of Harusaki et al. (2000). Crude samples were filtered through Miracloth and stained with  $25 \mu g l^{-1}$  propidium iodide. The relative fluorescence of total DNA was measured for each nucleus with an EPICS XL flow cytometry system (Beckman-Coulter, Munich, Germany) equipped with an argon laser (488 nm, 15 mW).

Chromosome number

Radicular tips of about 1.5–2.0 cm long were excised from the putative somatic hybrid plants, immersed in 2 mM 8-hydroxyquinoline for 15 h at  $\hat{4}^{\circ}$ C and fixed in a mixed solution of ethanol: acetic acid  $(3:1)$  for 24 h at 4 $\rm{°C}$ .

Enzymatic maceration and air-drying were performed according to Fukui (1996) with some modifications. The root tips were washed in distilled water to remove the fixative and macerated in an enzyme mixture containing 2% (w/v) Cellulase Onozuka RS (Yakult), 1%(w/v) Macerozyme R-200 (Yakult), 0.3% Pectolyase (w/v) (Kikkoman, Tokyo, Japan), and 200 mM EDTA at  $37^{\circ}$ C for 10 min.

Chromosomes were stained with 2% Giemsa solution (Merck, Darmstadt, Germany) in 1/30 phosphate buffer (pH 6.8) for 30 min, rinsed with distilled water, air-dried and observed under an optical microscope.

#### Extraction of total DNA

Total DNA was extracted from plants and cotyledonary tissues that failed to germinate normally, according to the method of Doyle and Doyle (1987). The total DNA was used for analyses of nuclear and cytoplasmic DNA by random amplified polymorphic DNA (RAPD) and cleaved amplified polymorphic sequence (CAPS) analyses.

#### RAPD analysis of nuclear DNA

RAPD analysis of nuclear DNA was performed by a method modified from that of Williams et al. (1990). The reaction mixtures (25  $\mu$ l) contained 10 mM Tris-HCl pH 8.9, 80 mM KCl, 1.5 mM MgCl<sub>2</sub>, 100  $\mu$ M dNTPs, 0.3  $\mu$ M primer, 2.5 U Tth Taq DNA polymerase and 10 ng genomic DNA. Reactions were carried out by repeating 45 cycles of the following thermal treatments: 94°C for 30 s,  $37^{\circ}$ C for 2 min and  $72^{\circ}$ C for 3 min, in an ASTEC Program Control System PC-700. Random 10-mer primers OPA-6, OPA-9 and OPA-10 (Operon Technology, Alameda, Calif.) were used for PCR. Reaction products were resolved by electrophoresis on 1.3% agarose gels containing  $0.5 \mu g$  ml<sup>-1</sup> ethidium bromide and subsequently photographed under UV light (360 nm). For each combination of samples and primers, PCR was carried out twice, and only stable polymorphism was taken into account.

## CAPS analysis of nuclear DNA

The internal transcribed spacer (ITS) region in nuclear ribosomal RNA (rRNA) was used for nuclear DNA analysis using primers ITS1 (5'-TCCGTAGGTGAACCTGCGG-3') and ITS4 (5'-TCCTC-CGCTTATTGATATGC-3') (Yasui and Ohnishi 1998). The reaction mixture (50  $\mu$ l) contained 10 mM KCl, 2.0 mM MgCl<sub>2</sub>, 0.2  $\mu$ M dNTPs, 0.2  $\mu$ M of each primer, 1 U Ex Taq DNA polymerase and 10 ng genomic DNA. Amplification was conducted by the following thermal treatments: pre-denature at 95°C for 5 min and 30 cycles of 1 min at 95°C, 1 min annealing at 60°C, 2 min at 72°C and further extension at 72°C for 10 min in a TaKaRa PCR thermal cycler MP TP3000. PCR products were digested with ten restriction endonucleases (EcoRI, XbaI, MspI, HaeIII, HinfI, MboI, XhoI, PvuI, PvuII and KpnI) for 15 h at  $37^{\circ}$ C. The digested DNA samples were resolved by electrophoresis on a 3.0% agarose gel containing 0.1  $\mu$ g ml<sup>-1</sup> ethidium bromide and visualized under UV light.

Table 1 Pairs of universal primers used for amplification of specific chloroplast (cp) and mitochondrial (mt) DNA regions



Table 2 Germination rate and ploidy level of somatic embryo strains obtained from electrofusion treatment



<sup>a</sup> Data were obtained after 3 months of culture

<sup>b</sup> Ploidy level of embryoids were evaluated using flow cytometry and chromo-some counting of root tip cells

<sup>c</sup> Number of germinated plants/number of somatic embryos cultured

CAPS analysis of cytoplasmic DNA

Amplification of chloroplast (cp) and mitochondrial (mt) DNA using cp and mt universal primer pairs (Table 1) was performed in a TaKaRa PCR thermal cycler MP TP3000. For analysis of cp-DNA, three primer pairs (RbcL-PSAI, TrnD-TrnT and trnK3914FtrnK2R) were used for amplification according to the methods of Cheng et al. (2002) and Ureshino and Mayajima (2002). For analysis of mt-DNA, three primer pairs of 18SrRNA-5SrRNA, nad5/1-nad5/2r and nad7/1-nad7/2r were used for amplification according to the method of Cheng et al. (2002) and Dumolin-Lapegue et al. (1997). PCR products were digested with ten restriction endonucleases, as described above, resolved by electrophoresis on a 3.0% agarose gel, and visualized under UV light as described above.

# Results and discussion

The first cell division occurred 1 week after electrofusion, and visible colonies were observed after 3 months of culture. Some of these colonies formed between one and

five green somatic embryos per plastic dish after a further 2 months. These embryos, approximately 2 mm in diameter, were transferred individually to a hormone-free medium as independent strains. One month after transfer, most of these embryos formed several secondary somatic embryos. The 13 independent strains thus obtained (R1– R13) were transferred to one-half strength MT medium containing 1 mg  $l^{-1}$  GA<sub>3</sub>, 10 g  $l^{-1}$  sucrose and 3 g  $l^{-1}$ gellan gum to promote normal germination. The germination rate of these somatic embryos varied from 0 to 60% (Table 2). Finally, 32 regenerated plants were obtained from eight strains 2 months after transfer (Fig. 1A). In other strains, somatic embryos showed only enlargement of cotyledonary tissue, and normal germination was not observed (Fig. 1B). The regenerated plantlets were transferred to pots and grown under greenhouse conditions. Six of the eight strains (R1, R3–R7) had normal morphological characteristics, but the remaining two (R2 and R8) showed malformation (Fig. 2). Normal-



Fig. 1 Normal (A) and abnormal (B) germination from somatic embryos



Fig. 2 Morphological characteristics and flow cytometry analysis of putative somatic hybrids. Strains R3 and R6 showed normal morphology, while R8 was malformed

ly, regenerated plants exhibited leaf size intermediate between that of both parents. They also had thick and round leaves, which are typical characters of somatic hybrids and polyploids.

Flow cytometry analysis of all regenerated plants showed that their fluorescence intensity coincided with that of the  $4 \times$  control (Fig. 2), while fluorescent intensity of cotyledonary tissues of five strains that did not

germinate also corresponded to that of the 4x control. Chromosome counts of root tip cells revealed that the chromosome number of R1–R8 plants was 36 (Fig. 3), which is the sum of chromosome numbers of round kumquat (2n=18) and 'Morita navel' orange (2n=18).

To confirm the hybridity of regenerated plants, we employed RAPD analysis in R1, R3, R5–R7 and their parents. As shown in Fig. 4, these strains yielded bands



Fig. 3 Chromosomes of putative somatic hybrid R3 in a root tip cell  $(2n=4\times 36)$ 

specific to both parents. Hybridity of these strains was further confirmed using CAPS in the regenerated plants (R5, R6 and R7) and their parents. Amplification of the ITS region of nuclear DNA resulted in a fragment of the same size from somatic hybrids and both parents. After digestion of the fragment with MspI, the regenerated plants had specific bands derived from both parents (Fig. 5). Furthermore, hybridity of R10 and R12 somatic embryos, which failed to germinate normally, was confirmed using RAPD analysis. Primers OPA-6, OPA-9 and OPA-10 amplified specific bands from both parents (data not shown). These results indicate that seven strains, including R10 and R12, were somatic hybrids.

Cp- and mt-DNA amplifications were performed on three somatic hybrids (R5, R6 and R7) and both parents using three cp- and mt-DNA universal primer pairs. While every primer pair amplified bands satisfactorily, they did not reveal any polymorphism on agarose gels.

 $MspI$ 



Fig. 5 Restriction pattern of the MspI-digested ribosomal RNA (rRNA) internal transcribed spacer (ITS) of nuclear genomes. Lanes: *M* Size marker; *N* 'Morita navel' orange;  $5-7$  putative somatic hybrids R5, R6, R7; K round kumquat. Arrowheads Bands specific to each parent

When the PCR products were digested with ten restriction endonucleases, cp-DNA polymorphism was observed in nine primer/enzyme combinations, RbcL-PSAI/MspI, HinfI (Fig. 6A), HaeIII and MboI, TrnD-TrnT/MspI, HinfI, HaeIII and MboI(Fig. 6B), and trnK3914F-trnK2R/ HinfI(Fig. 6C). All of the three somatic hybrids had uniform bands identical to those of round kumquat. On the other hand, mt-DNA polymorphism was seen in only one combination of 18SrRNA-5SrRNA/MspI (Fig. 7), and the remaining combinations showed no polymorphism. With this combination, these somatic hybrids showed the same band patterns as round kumquat. These results

Fig. 4 Random amplified polymorphic DNA (RAPD) analysis of putative somatic hybrids and their parents. Lanes: M Size marker; N 'Morita navel' orange; 1, 3, 5–7 somatic hybrids R1, R3, R5, R6, R7; K round kumquat. Arrowheads Bands specific to each parent. OPA-6, OPA-9 Random 10-mer primers used for PCR

 $OPA-6$ 

 $OPA-9$ 



Fig. 6 Restriction patterns of the HinfI-, MboI, and HinfIdigested RbcL-PSAI (A), TrnD- $TrnT$  (B), and  $TrnK3914F-$ TrnK2R (C) regions of chloroplast genomes. Lanes: N 'Morita navel' orange; 5–7 putative somatic hybrids R5, R6, R7;  $K$  round kumquat;  $M$  size marker. Arrowheads Bands specific to round kumquat





Fig. 7 Restriction pattern of the MspI-digested 18SrRNA-5SrRNA regions of mitochondrial genomes. Lanes: N 'Morita navel' orange; 5–7 putative somatic hybrids R5, R6, R7; K round kumquat; M size marker. Arrowheads Bands specific to round kumquat

showed that three somatic hybrids had cp- and mt-DNA of round kumquat.

The cytoplasmic composition of somatic hybrids has been investigated mostly through Southern analysis of cytoplasmic or total DNA with cp- and mt-DNA probes, or through cp- and mt-DNA restriction-profile analysis (Earle 1995; Cardi et al. 1999; Bastia et al. 2000). These methods are usually expensive and/or time-consuming. Moreover, a considerable amount of DNA is required. Recently, new types of PCR-based molecular markers for cp- and mt-DNA—CAPS markers—have been developed. CAPS is simple, more rapid and less expensive when compared to the direct analysis of restricted cp- and mt-DNA. In citrus, CAPS markers were employed to characterize the cp- and mt-DNA of somatic hybrids or cybrids between Microcitrus papuana and Citrus jambhiri (Liu et al. 2002), Citrus reticulata and Poncirus trifoliata (Guo et al. 2002), and C. sinensis and F. crassifolia (Cheng et al. 2003). In the present study, cp- and mt-DNA analyses using CAPS markers revealed that cp- and mt-DNA of the somatic hybrids were derived from round kumquat without recombination in the regions analyzed.

Deng et al. (1992) produced intergeneric somatic hybrids between 'Valencia' sweet orange and Meiwa kumquat. These somatic hybrids grew less vigorously and showed dieback of their shoots annually (Shi et al. 1998). Cheng et al. (2003) described that both the cp- and mt-DNA in these somatic hybrids were derived from 'Valencia' sweet orange. Furthermore, they also suggested that the mt-DNA pattern is correlated with the phenotypic abnormality of these somatic hybrids and that nuclear-cytoplasmic incompatibility may be the cause of dieback. In the present study, such phenotypic abnormality was also observed in the somatic hybrids between round kumquat and 'Morita navel' orange, which had both the cp- and mt-DNA of round kumquat. In our preliminary studies, we carried out hybridization between the monoembryonic cultivar 'Kiyomi' tangor (Citrus unshiu  $\times$  C. sinensis) and several Fortunella species. 'Kiyomi' tangor yielded a lot of hybrid seed in crosses with Changshou kumquat (Fortunella obovata) and Malayan kumquat (Fortunella polyandra) (data not shown). However, when 30 flowers of 'Kiyomi' tangor were crossed with Meiwa kumquat and oval kumquat (Fortunella margarita), only two hybrid seeds were obtained in each case. The two seeds obtained in each cross germinated but grew very slowly. Furthermore, we could not obtain normal seeds from hybridization between 'Kiyomi' tangor and round kumquat. These results suggest that the reason for this phenotypic abnormality may not be nuclear-cytoplasmic incompatibility, but nuclear-nuclear incompatibility between some Citrus species and round kumquat.

In conclusion, we obtained intergeneric somatic hybrids between round kumquat and 'Morita navel' orange with cp- and mt-DNA of round kumquat. This is the first report of successful somatic hybridization of kumquat and navel orange with male and female sterility.

To date, three normal strains have already been grafted onto trifoliate orange and planted in our experimental field. These amphidiploid somatic hybrids could be useful parents for producing triploid seedless kumquat cultivars with flavor of navel orange. These plants have already been propagated and are under cultivation for evaluating male or female fertility and fruit characteristics.

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