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Protoplast isolation and plant regeneration of different doubled haploid lines of cauliflower (Brassica oleracea var. botrytis)

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Abstract Protoplasts were isolated from hypocotyls of 5-day-old seedlings of five doubled-haploid (DH) lines of cauliflower after enzymatic digestion in 0.1% macerozyme R-10 and 1.0% cellulose R-10. Shoot regeneration was achieved in four of the five DH lines. Protoplast yield and frequency of cell division and shoot regeneration varied among experiments and DH lines. Sequence-related amplified polymorphism (SRAP) analysis on all the regenerated plants of each DH line indicated that their genetic compositions were homologous with their mother plants, but few regenerated plants of DH lines no. 3 and 5 were detected with changes in ploidy levels.

Keywords Cauliflower · Protoplast · Plant regeneration · Ploidy level - SRAP analysis

Abbreviations

Introduction

Brassica oleracea is an extremely diverse species which includes several important vegetable crops including broccoli, brussels sprouts, cauliflower, various types of cabbage, kohlrabi and kale (Hansen et al. [1999](#page-6-0)). Worldwide cauliflower production is estimated at 5.3 million tonnes making it an economically important vegetable (Hodgkin [1995\)](#page-6-0). Traditional breeding of this crop has led to significant improvements in crop quality and productivity. However, plant breeding programmes are limited in terms of availability of suitable genes in the germplasm base. The application of new technologies, such as somatic hybridization and genetic transformation, to cauliflower varieties would pave the way for the generation of new cultivars possessing more agriculturally and economically viable genetic traits. An efficient plant regeneration system from protoplast has proved a very useful technique for crop genetic improvement programs, a prerequisite for somatic hybridization and genetic transformation by direct DNA uptake.

Techniques for protoplast isolation and culture in Brassica oleracea var. botrytis have been reported (Vatsya and Bhaskaran [1982](#page-7-0); Jourdan et al. [1990](#page-6-0); Walters and Earle [1990;](#page-7-0) Veera et al. [2009\)](#page-7-0) and somatic hybridizations between cauliflower and other species have also been studied but only for a few genotypes of minor or no commercial importance (Jourdan et al. [1989](#page-6-0); Yarrow et al. [1990](#page-7-0); Walters et al. [1992](#page-7-0); Navrátilová et al. [1997;](#page-6-0) Sheng et al. [2008](#page-7-0); Scholze et al. [2010\)](#page-7-0). Moreover, these investigators used mainly F_1 cultivars and such material is genetically uniform, but it is often difficult to analyze the progeny owing to heterozygosity in the background. An alternative is doubled haploid (DH) lines produced by microspore culture, which are also uniform and show complete homozygosity in their background (Tsukazaki et al. [2002](#page-7-0)). Therefore, the improved DH lines can be directly used as parents for the production of F_1 hybrids. For breeding, it is more necessary and significative to regenerate plants from different genotypes of DH lines.

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Table 1 Composition of protoplast culture media

The objectives of this study were to analyze protoplast yields derived from in vitro-grown hypocotyls of five DH lines of cauliflower with high breeding value, establish conditions for protoplast culture and shoot regeneration, and determine genetic fidelity of regenerated plants by SRAP analysis.

Materials and methods

Plant material

For protoplast isolation, a total of five DH lines of cauliflower (DH1-5) produced by microspore culture in our previous study (Gu et al. [2007](#page-6-0)) were selected, which have desirable attributes but need to be further improved.

Isolation and purification of protoplasts

Protoplasts were aseptically isolated from hypocotyls of 5-day-old seedlings grown in the dark at 25° C on hormone free Murashige and Skoog basal medium (MS medium, Murashige and Skoog [1962\)](#page-6-0). Hypocotyls (500 mg) were cut into small segments and incubated for 2 h with 5 ml of a preplasmolytic solution (54.6 g/l Sorbitol, 7.4 g/l $CaCl₂·2H₂O$, pH 5.6–5.8) followed by incubation in 5 ml of a cell wall degrading enzyme solution (Table 1) containing 0.1% macerozyme R-10 and 1.0% cellulose R-10. Following overnight incubation (15 h) at 24° C in the dark, the solution of protoplast-enzyme mixtures was filtered through a 200 lm nylon mesh into a centrifuge tube. The filtrate was then over-layered with 2.5 ml washing solution $(18.4 \text{ g/l } CaCl_2 \cdot 2H_2O, 9.0 \text{ g/l } NaCl, 1.0 \text{ g/l } Glucose, 0.3 \text{ g/l }$ KCl, pH 5.6–5.8). Following centrifugation (1,200 rpm for 10 min), intact protoplasts, floating at the interface of the two solutions, were collected and washed twice by re-suspending in the washing solution (same solution as described above), and then centrifuged for 4 min at 800 rpm. Protoplast viability was measured with fluorescein diacetate (Widholm [1972\)](#page-7-0).

Protoplast culture

Culture in liquid medium

Isolated protoplasts were cultured at a density of 5×10^5 protoplasts ml^{-1} in liquid medium 1/2 (Table 1) containing 0.2 mg l^{-1} naphthaleneacetic acid (NAA), 0.2 mg l^{-1} benzyladenine (BA) and 0.5 mg 1^{-1} 2, 4-Dichlorophenoxyacetic acid (2, 4-D) in a total volume of 2 ml in 3 cm Petri dishes without agitation. The plates were sealed with parafilm and cultured in darkness at 25°C. When normal cell divisions were observed, 400 μ l of 2/2 dilution medium (Table 1) was added to each dish every 3 days. Suspensions with colonies consisting of more than 20 cells were transferred to solidified 3/10 medium (Table 1) for callus growth under dim lighting (2.99 lmol $m^{-2} s^{-1}$). Calli that reached a size of 2–3 mm were transferred to shoot inducing medium and subcultured once every 2 weeks.

Nurse culture

Protoplasts of tuber mustard were employed as the nurse cells in the present study, which had been reported with positive effect in the protoplasts culture of Brassica species (Chen et al. [2004](#page-6-0)). Purified protoplasts of cauliflower were resuspended in 1/2 medium at a density of 2×10^5 ml⁻¹ and 1 ml of this mixture was quickly mixed with an equal volume of the 1/2 agarose medium (1/2 medium with 2 g/l agarose) to give evenly dispersed protoplasts. Then, 2 ml liquid medium (MS medium contained 0.5 mg l^{-1} BA, $1 \text{ mg } 1^{-1}$ 2,4-D, $0.5 \text{ mg} 1^{-1}$ NAA, 0.4 M mannitol) with protoplasts of tuber mustard at a density of 1×10^5 ml⁻¹ was added onto the agarose medium containing cauliflower protoplasts. After 2 weeks of culture in darkness at 25° C, the liquid medium was removed and the plates were transferred under dim lighting (2.99 lmol m^{-2} s⁻¹) for callus growth.

Shoot regeneration

Calli were transferred to shoot regeneration medium when they reached 2–3 mm in size. For shoot regeneration,

various media (based on MS-medium, with 30 g 1^{-1} sucrose and 8.0 g 1^{-1} plant agar) including different plant growth regulator (PGR) combinations and concentrations were evaluated. Calli were cultured at 24° C in the light (16 h photoperiod, 25–30 μ mol m⁻² s⁻¹). Shoot regeneration was evaluated after 10 weeks of culture by recording the number of calli with shoots.

Acclimatization

Regenerated plants from each DH line were transferred to the greenhouse to monitor their growth. Plants were transferred to soil (peat substrate) at $22-24$ °C and kept at 100% humidity for 2 weeks. After 4–6 weeks, they were transferred to 9 cm pots, and grown under normal greenhouse conditions.

Flow cytometry

Healthy young leaves, 0.5 cm^2 , were chopped with a razor blade in 0.6 ml ice-cold buffer (100 ml deionized water, 0.1 mol/l citric acid, 0.5 g Tween 20) and 2.4 ml propidium iodide (PI) solution was then added. The suspension was filtered through a mesh, and after 5 min, these were analyzed using a BD flow cytometer (FACSCalibur, BD Biosciences, USA). All regenerated plants were tested for ploidy changes by flow cytometry with three replications.

Regenerated plants analyzed by PCR-SRAP

Total genomic DNA was isolated from leaf tissue collected in bulk from regenerated plants as described using the cetyltrimethylammonium bromide (CTAB) method (Paterson et al. [1993\)](#page-6-0). DNA concentration was quantified by visual comparison to λ DNA standards on ethidium bromide-stained agarose gels.

Among the 50 pairs of primers tested, 10 primer combinations (Table 2) showed clear and reproducible bands were selected for amplification of the template DNA to detect variation. PCR amplification was performed following Sheng et al. [\(2008](#page-7-0)). The PCR products were electrophoresed in 6% polyacrylamide denatured gels, ran at 100 W for 2 h, and the banding patterns were visualized using silver staining as described by Panaud et al. [\(1996](#page-6-0)). The gel was photographed after being dried at room temperature.

Results

Protoplast isolation and culture

Protoplasts were successfully isolated from hypocotyls of all five DH lines of cauliflower. Protoplast yield varied between 0.3–5.2 \times 10⁶ protoplasts per gram fresh weight in the presence of different enzyme combinations (Table [3](#page-3-0)). The protoplast yield significantly increased with increasing enzyme concentration, but protoplast viability was significantly diminished when beyond ER2. Therefore, the enzyme combinations ER2 was the optimum for protoplast production with the highest protoplast yield and viability.

Effect of genotype and culture system on cell division

Five DH lines were compared to evaluate efficiency of protoplast culture under two culture conditions, liquid culture and nurse culture. For all genotypes, protoplastderived cells began to divide after 3–4 days. Overall, significant differences between the two culture systems were observed in the five DH lines and nurse culture resulted in much higher frequency of cell division after 6 days of culture (Fig. [1\)](#page-3-0), which is $5-63\%$ in liquid medium but 32–92% for them employed by nurse culture. DH line 5 showed the highest frequency of cell division (92%) in nurse culture. However, DH line 1 cultured in liquid medium showed the lowest frequency of cell division (5%) but in nurse culture, the frequency improved to be 32%.

Table 2 The primer sequences of SRAP used in this experiment

Forward primers $(5'–3')$		Reverse primers $(5'–3')$	
Me1	TGAGTCCAAACCGGATA	Em1	GACTGCGTACGAATTAAT
Me ₃	TGAGTCCAAACCGGAAT	Em ₆	GACTGCGTACGAATTGCA
Me ₅	TGAGTCCAAACCGGAAG	Em8	GACTGCGTACGAATTCTG
Me9	GACTGCGTACGAATTCGA	Em10	GACTGCGTACGAATTCAG
Me11	CTGGCGAACTCCGGATG	Em12	GACTGCGTACGAATTGTC
		Em13	GACTGCGTACGAATTGGT

The selected 10 primer combinations were: $Me1 + Em6$, $Me1 + Em12$, $Me1 + Em13$, $Me3 + Em1$, $Me3 + Em10$, $Me5 + Em6$, $Me5 + Em10$, $Me9 + Em8$, $Me11 + Em1$, $Me11 + Em13$

and yield of protoplasts

Means within a column

significantly different at

Fig. 1 Frequency of cell division of different DH lines protoplasts with liquid or nurse culture after 6 days (means of two experiments with three replications)

In general, frequency of cell division for all five DH lines varied not only between genotypes, but also between different culture systems of protoplasts of the same genotype.

Nearly all protoplast-derived cells, that underwent first divisions, developed further into micro-colonies of 2–3 mm in diameter 5–7 weeks following isolation. These were then transferred to differentiation media for shoot induction (Fig. 2). Significant differences in frequencies of shoot regeneration were observed among the different genotypes and PGR combinations and concentrations (Fig. [3](#page-4-0)). Shoot regeneration was achieved for four DH lines (except DH2). In the PGR combinations of $6-BA+ IAA$ and $ZT + IAA$, ZT was more effective for shoot induction in cauliflower (Fig. [3\)](#page-4-0). The highest frequency of shoot regeneration (83% for DH line 3) was obtained on medium containing 1.0 mg l^{-1} ZT and 0.2 mg 1^{-1} IAA (Figs. [3,](#page-4-0) [4](#page-4-0)) and this PGR combination also resulted in the highest frequency of shoot regeneration for DH line 1 and 5 (61 and 49%, respectively).

Fig. 2 Culture and regeneration of the protoplasts of DH1. a First cell division 3 days after culture; b microcalli formation after 21 days of culture; c, d shoot regeneration from calli

Fig. 3 Frequency of shoot regeneration of protoplast-derived calli of five DH lines of cauliflower on media with different plant growth regulator concentrations after 10 weeks of culture. Means of one experiment consisting of 3–5 replications with 8–12 calli/Petri dish

For DH line 4, shoots could only regenerated on the medium containing 1.0 mg l^{-1} ZT, but the frequency was low $(\langle 10\% \rangle)$.

Morphological, ploidy level and SRAP analysis

A total of 162 plants from different colonies originating from protoplasts of DH lines 1, 3, 4, 5 were successfully transferred to the greenhouse (Table 4). For DH lines 1 and 4, the ploidy levels of the regenerated plants were all diploid, which were consistent with their mother plants. However, two regenerated plants of DH line 3 and three plants originating from protoplasts of DH line 5 showed higher ploidy levels (Table 4; Fig. [5](#page-5-0)), which were generally stunted with shrunken leaves.

SRAP analysis was produced on all the regenerated plants of each DH line. Among the ten pairs of primers selected, each primer combination generated a unique set of amplification products ranging in size from 100 to 1,000 bp. The band patterns of the regenerated plants from protoplasts of each DH line were homologous with their mother plants, which indicated their genetic homogeneity. Figure [6](#page-5-0) showed two representative amplified band patterns produced by primer combinations of Me1 $+$ Em13 and Me9 $+$ Em8.

Table 4 Acclimatization and ploidy changes of plants regenerated from protoplasts

Fig. 4 Organogenic callus cultures with shoots of DH line 3 after 10 weeks of culture on media with different plant growth regulator combinations (6 cm Petri dishes). a 1.0 $ZT + 0.2$ IAA (mg/l); **b** 1.0 $ZT + 0.1$ IAA (mg/l); c 2.0 $BA + 0.2$ IAA (mg/l); d 2.0 $BA + 0.1$ IAA (mg/l); e 1.0 ZT (mg/l)

Fig. 5 Histograms of nuclear DNA contents obtained by flow cytometric analysis. a DH5 with diploid level; b a regenerated plant of DH5 with tetraploid level; c a regenerated plant of DH5 with hexaploid level. Ordinate: Number of nuclei; Abscissa: PI

Fig. 6 PCR-SRAP analysis of the regenerated plants. a Polymorphic bands of SRAP fragments generated by Me1 + Em13. M : DNA ladder, Lane 11: DH3, Lanes 1-10: Regenerated plants of DH3; **b** polymorphic bands generated by $Me9 + Em8$. *M*: DNA ladder, Lane 11: DH1, Lanes 1-10: Regenerated plants of DH1

Discussion

Although plantlets were successfully regenerated from protoplasts of cauliflower in previous studies (Vatsya and Bhaskaran [1982;](#page-7-0) Jourdan et al. [1990](#page-6-0); Walters and Earle [1990;](#page-7-0) Veera et al. [2009](#page-7-0)), their protocols varied greatly in either culture parameters or genotypes and mostly used leaves as explant material with low frequencies of shoot regeneration. In the present investigation, a new nurse culture method using protoplasts of tuber mustard as nurse cells was employed in the culture of cauliflower protoplasts for the first time and the highest frequency of cell division (92%) was obtained using this method. Moreover, plant regeneration from hypocotyls protoplasts of different DH lines of cauliflower was reproducibly achieved and the

fluorescence. Fluorescence intensity is proportional to nuclear DNA quantity and the position of the dominant peak (arrow indication) reflects the ploidy level

frequency of shoot regeneration is higher than previously reported for cauliflower, which gave a maximum regeneration rate of 72% (Veera et al. [2009](#page-7-0)).

Plant regeneration from protoplasts is influenced by many factors, including the methods of protoplast isolation, inoculation density, culture system, nurse cells, and medium compositions (Meyer et al. [2009](#page-6-0); Eeckhaut et al. [2009](#page-6-0); Castelblanque et al. [2010](#page-6-0); Prange et al. [2010](#page-7-0); Tiwari et al. [2010\)](#page-7-0). The use of feeder layer-culture systems with protoplasts of other species as nurse cells provided a simple and reliable technique for enhancing cell division and plating efficiency of isolated protoplasts (Walters and Earle [1990](#page-7-0); Karamian and Ebrahimzadeh [2001](#page-6-0)). Chen et al. [\(2004](#page-6-0)) reported the use of tuber mustard protoplasts as nurse cells, which was essential for sustained division and colony formation from red cabbage protoplasts. In the present study, nurse cells of tuber mustard protoplasts also had an enhancing effect on cauliflower protoplast division. By nurse culture, all five DH lines showed much higher cell division frequency compared to liquid culture, probably as a result of the release of growth factors by nurse cells into the surrounding medium, which stimulated the cell division of cauliflower. A similar stimulatory effect of nurse culture on cell division and plant regeneration from protoplasts has been reported for rice (Kyozuka et al. [1987](#page-6-0); Li and Murai [1990\)](#page-6-0) and maize (Petersen et al. [1992](#page-6-0)).

It was evident that there was no correlation between division efficiency and subsequent shoot differentiation in the present study (Figs. [1](#page-3-0), [3](#page-4-0)). A similar phenomenon had been observed in B. oleracea (Robertson et al. [1988\)](#page-7-0) and Petunia (Meyer et al. [2009](#page-6-0)). These results indicated that cell division and shoot differentiation are controlled by different developmental mechanisms.

The ploidy of protoplast-derived plants from each DH line were analyzed by flow cytometry, a well established and reliable method that is less time consuming than chromosome counting (Ulrich and Ulrich [1991](#page-7-0); Mallón et al. 2010; Stancheva et al. [2011;](#page-7-0) Yang et al. [2010](#page-7-0); Zhang et al. [2010](#page-7-0)). Changes in ploidy level were just observed in the regenerated plants of DH lines 3, 5 and the frequency were 3.5 and 2.8% respectively, which were much lower than the a previous study on protoplast culture of cauliflower with 15–25% tetraploidy in the regenerated plants (Hansen and Earle 1994). This may be attributed to the application of nurse culture, which inhibited the spontaneous fusion of diploid protoplasts.

SRAP is a new but very powerful molecular marker system, which was first reported by Li and Quiros (2001). Budak et al. (2004) compared the four marker systems in buffalograss and found the values of average discriminating power as: $SRAP > SSR > ISSN > RAPD$. Ferriol et al. (2003) reported that SRAP could provide more information and was more concordant with the genomic variations and the evolutionary history of the morphotypes than AFLP. In this study, SRAP amplification gave rise to a lot of polymorphic bands, leading to desirable discrimination ability. SRAP analysis indicated that plants regeneration from protoplast of cauliflower was homogeneous, although the morphology of few regenerated plantlets varied from mother plants. Variation of the morphology may be due to variation of the ploidy levels and the changing of the culture environments.

In summary, this study has demonstrated that protoplast isolation and culture up to the microcallus stage can be achieved for all the DH lines of cauliflower with breeding importance. At present, shoot regeneration is somewhat limiting, although this can be achieved for most DH lines used in this study. The protoplast regeneration system described herein will enable the pursuit of somatic hybridization and transformation by direct DNA uptake, which may result in the development and production of novel and popular cauliflower varieties.

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