Cryopreservation enhances embryogenic capacity of *Gentiana cruciata* (L.) suspension culture and maintains (epi)genetic uniformity of regenerants

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Abstract The embryogenic cell suspension culture of *Gentiana cruciata*, cryopreserved by the encapsulation/dehydration method, survived both short- (48 h) and long-term (1.5 years) cryostorage with more than 80% viability. To assess the influence of cryotreatments on the embryogenic potential, a proembryogenic mass was encapsulated and exposed to the following treatments: (1) osmotic dehydration (OD), (2) OD + air desiccation (AD) and (3) OD + AD + cryostorage (LN). The somatic embryogenesis efficiency increased ten times after osmotic dehydration. The AD and LN cryotreatments did not cause any significant alterations in somatic embryo production. We monitored the (epi)genetic stability of 288 regenerants derived from: non-cryotreated, short-term, and long-term cryostored tissue using metAFLP markers and ten primer combinations. Changes in the sequence and DNA methylation levels were studied by subjecting the DNA to digestion with two pairs of isoschisomer restriction enzymes (*Kpn*I/*Mse*I and *Acc65*I/*Mse*I). Two new AFLP unique DNA fragments at the DNA sequence level, with no differences at the methylation level, were found between regenerants derived from cryopreserved tissue, compared with the non-cryotreated controls. The *Acc65*I/*Mse*I methylation levels for the three groups of regenerants were not significantly different. Cluster analysis was capable of identifying a number of sub-clusters. Only one of the sub-clusters comprises almost all regenerants derived from non-cryotreated and short-term cryostored tissue. Plantlets derived from long-term cryostored tissue were grouped into separate clusters. The observed AFLP alterations did not appear to be associated with the use of cryopreservation, but were probably related to the process of in vitro culture.

Keywords Encapsulation/dehydration · metAFLP · Osmotic dehydration · Short- and long-term cryostorage · Somatic embryogenesis

Introduction

Cryo-conservation protocols have been established for many vegetatively propagated species and introduced into the world genebanks (Engelmann 2004). The ultralow temperature of liquid nitrogen efficiently stops the metabolic processes. Plant material is then capable to be preserved for a theoretically unlimited period of time (Walters et al. 2004). However, the cryotreatment, cryostorage or post-thaw manipulations can cause stress-induced factors and such factors may lead to morphogenic or genetic alterations.

The majority of studies prove that cryopreservation does not affect embryogenic potential of calli or suspension culture, which has been documented for example for a majority of 39 independent embryogenic lines of *Hevea brasiliensis* (Lardet et al. 2007). The increase in the embryogenic capacity after cryopreservation becomes visible for the first time in *Citrus deliciosa* (Aguilar et al. 1993). Usually, the regeneration enhancement is considered as the final result of the cryopreservation procedure. So far, the phenomenon was ascertained for embryogenic tissue of: *Festuca arundinacea*, *Lolium* spp. (Wang et al. 1994), *Hevea brasiliensis* (Engelmann et al. 1997), *Vitis vinifera*...
(Wang et al. 2002) and Cyclamen persicum (Winkelmann et al. 2004). Investigations are needed to determine whether the regeneration enhancement effect is caused by pretreatment, freezing, or by a combination of the two.

The assessment of plant genetic uniformity to validate newly established cryopreservation protocols has been a subject of increasing interest (Harding 2004). So far, only some evidence of genetic alterations after cryopreservation treatment of in vitro-derived plant material was shown. Changes at the DNA sequence level seem to appear sporadically, or accidentally, and most frequently are carried on by single specimens and characterized by one (Ahuja et al. 2002; Martín and González-Benito 2005; Dixit et al. 2003; Moukadiri et al. 1999; Sánchez et al. 2008) or two new markers (Martín and González-Benito 2009; Urbanová et al. 2006). However, experiments conducted on 6 different genotypes of Carica papaya showed varying levels of genomic DNA modifications (up to 10.07%) covering 14 changes (Kaity et al. 2008). Genetic stability of regenerants recovered from cryopreserved plant material is also associated with DNA methylation changes which were ascertained in all experiments conducted up till now. Studies of Fragaria gracilis (Hao et al. 2002a), Citrus sinensis (Hao et al. 2002b), Malus pumila (Hao et al. 2001), Ribes (Johnston et al. 2009), Carica papaya (Kaity et al. 2008, 2009), Humulus lupulus (Peredo et al. 2008), Prunus dulcis (Channuntapipat et al. 2003) and Arabidopsis thaliana (Wang and He 2009) showed detectable differences. These differences demonstrate the complexity of the processes assisting cryopreservation and influencing cell culture variation.

Several molecular markers were used to analyze genetic stability of cryopreserved plant material (Harding 2004). Currently, application of randomly amplified polymorphic DNA (RAPD) and amplified DNA methylation polymorphism (AFLP) is most frequently noticed. However, in terms of a study of structural and methylation DNA changes, more than one molecular marker is applied, e.g. RAF (randomly amplified DNA fingerprinting) and AMP (amplified DNA methylation polymorphism) for Carica papaya (Kaity et al. 2008, 2009), AFLP (amplified fragment length polymorphism) and MSAP for Malus pumila and Fragaria gracilis (Hao et al. 2001, 2002a), RAPD and MSAP for Citrus sinensis (Hao et al. 2002b) or RAPD, AFLP and MSAP for Humulus lupulus (Peredo et al. 2008). The sensitivity of detecting changes, in both the structure and methylation of DNA, was improved by subjecting the DNA to digestion separately with HpaII, Bsp143I (relatively methylation sensitive), Mspl and Mbol (methylation insensitive) restriction endonucleases before RAPD-PCR (Channuntapipat et al. 2003). This latter comparison of methylation (in)sensitive isoschisomers resulted in detectable differences due to cryopreservation and in vitro culture, at the sequence and DNA methylation levels. Recently, a new variant of the metAFLP approach was introduced that is dedicated to simultaneous quantification of sequence, and site DNA methylation changes within the sequence recognized by the Acc65I/Msels and KpnI/Msels pairs of endonucleases. Acc65I and KpnI are isoschizomers that differ in their sensitivity to site DNA methylation. Acc65I is insensitive to dcm methylation, although its activity is blocked by both dam and CpG methylation. KpnI is insensitive to all forms of methylation (Bednarek et al. 2007). In the present study, a new variant of metAFLP was employed to screen for changes in genomic DNA structure and DNA methylation patterns.

The gentian cell suspension cultures are an invaluable long-term source of numerous somatic embryos (Mikuła et al. 2005b, 2008; Fiuk and Rybczyński 2008), totipotent protoplasts (Fiuk and Rybczyński 2007), and embryogenic tissue for the production of interspecies somatic hybrids and transformed plants (Rybczyński et al. 2008). This type of culture had a considerably higher effectiveness of somatic embryogenesis (Mikuła et al. 2005b) than the agar culture (Mikuła and Rybczyński 2001). The somatic embryos could also be received regularly for more than 11 years for G. tibetica (Mikuła et al. 2008), and over 3 years for G. cruciata (Mikuła et al. 2005b). Nonetheless, the reduction in embryogenic competence is clearly visible with time (Mikuła et al. 2008). After a 12.5-year maintenance, the somatic embryo productivity of the G. tibetica suspension culture decreased by about 53 times. Our previous studies showed that cryopreservation is a reliable approach to preserve the viability and recovery of gentian cell suspension cultures. Our previous studies also showed that the encapsulation/dehydration protocol is the most useful among the three studied methods of cryo-conservation (Mikuła 2006; Mikuła et al. 2008). The aims of the current studies were (1) to assess the influence of the subsequent steps of cryopreservation by encapsulation/dehydration on the G. cruciata somatic embryo production and (2) to evaluate the uniformity of regenerants with the help of metAFLP analysis.

Materials and methods

Plant material and cryopreservation procedure

Experiments were carried out on proembryogenic mass (PEM) derived from the 2- and 3.5-year-old suspension culture of G. cruciata L. The induction and maintenance were elsewhere described (Mikuła et al. 2005a). The cell suspension was subcultured every 7 days. For cryopreservation by encapsulation/dehydration, the cell aggregates were encapsulated in 3% (w/v) sodium alginate (Sigma)
Osmotic dehydration (OD) was conducted by increasing the concentration of sucrose from 0.3, 0.5, 0.75 (48 h for each) to 1.0 M (1 day). Alginate beads were then harvested and surface dried in a laminar flow chamber at room temperature for 5 h (AD) and loaded into 2-ml cryovials (15 beads per vial). Cryovials were immersed directly into LN where they remained for 48 h or 1.5 year. The scheme of the cryopreservation procedure, including thawing and recovery of suspension cultures, is shown in Fig. 1.

Cell survival, after short- (48 h) and long-term (1.5 year) cryostorage, was determined in two independent experiments (9 replicates of 10 beads in each). The 2,3,5-triphenyltetrazoliumchloride (TTC) reduction test, as previously described for gentian cultures by Mikuła et al. (2006), was used. Encapsulated cells were stained immediately after embedding (as a control) and 2 days after thawing. The biomass growth of PEM after 4 weeks of post-thawing culture (the first 2 weeks in agar and later in liquid culture) (Mikuła et al. 2008) was assessed (1 replicate = 10 beads). Analyses were performed in two independent experiments (of 3 repetitions). The results were estimated using a single factor analysis of variance (ANOVA) and a Fisher’s least significant difference test (LSD) using Statgraphics Plus software. Significance is stated at \( P < 0.05 \).

Experimental design for embryogenic productivity assessment

To assess the influence of the cryotreatments on the embryogenic potential, the PEM derived from the 3.5-year-old suspension culture was encapsulated and exposed to the following treatments: (1) osmotic dehydration (OD), and (2) OD + air desiccation (AD) and (3) OD + AD + 48 h storage in liquid nitrogen (LN) (Fig. 1). After each cryotreatment step, the recovery of cultures was performed. After 6 weeks, the PEM derived from recovered suspension cultures was implanted onto the agar regeneration medium supplemented with 0.5 mg/L gibberellic acid, 1.0 mg/L kinetin and 80.0 mg/L adenine sulfate (Mikuła et al. 2005a). Cell aggregates from non-encapsulated, non-treated and non-frozen suspension culture were treated as the control.

The efficiency of somatic embryogenesis, expressed as a number of cotyledonary stage embryos per 50 mg of PEM, was determined in two independent experiments (12 Petri dishes for each). Data were recorded in the sixth week, after the transfer of PEM onto regeneration medium. The results were estimated using a single-factor analysis of variance (ANOVA) and a Fisher’s least significant difference test (LSD) using Statgraphics Plus software. Significance is stated at \( P < 0.05 \).

Plant material and experimental design for molecular analysis of regenerants

For molecular analysis, the somatic embryo-derived regenerants obtained from: control tissue i.e. non-cryotreated and non-cryostored PEM (NC), short-term (48 h) (SC) and long-term (1.5 year) cryostored PEM (LC) were employed (Fig. 2). NC and SC regenerants derived from 3.5-year-old cell suspension, LC regenerants derived from 2-year-old cell suspension. A 3.5-year-old suspension culture of
G. cruciata was 80 times more subcultured than the 2-year-old one. The regenerants were cultured on the half-strength MS basal medium. A 9-month-old plantlets, having at least 10 leaves, were subjected for molecular analysis. Regenerants originated from two independent PEM implantations on agar regeneration medium, within a 2-week interval (after 4 and 6 subcultures of recovered suspension cultures). Each implanting repetition included 48 plantlets (a total of 288 regenerants for NC, SC and LC).

Modified analysis of amplified fragment length polymorphism (metAFLP)

Total genomic DNA was isolated from leaves using the Dneasy Plant Mini Kit (Qiagen, GmbH, Hilden, Germany) according to the manufacturer’s instructions, following the methylation-sensitive AFLP previously described (Bednarek et al. 2007). Each DNA sample, representing an individual regenerant, was briefly placed in two independent vessels. One part of the sample was digested with KpnI/MseI, while the other with Acc65I/MseI, following by adaptor ligation and pre-selective amplification steps (Table 1). The products of the non-selective PCR underwent a selective amplification step using the following primer combinations: CpG-GAC/M-GC, CpXP-GAGA/M-CTG, CpXP-AGC/M-CGT, CpG-GGC/M-CAA, CpG-GAG/M-CGA, CpXP-AGG/M-CAC, CpG-GCA/M-CCG, CpG-GGT/M-CAT, CpXP-AGT/M-CCC, CpXP-GGT/M-CAG. Basic primer sequences were 5'-CAT GCG TAC AGT ACC-3' for CpG- and 5'-GAT GAG TCC TGA GTA A-3' for M-. The CpG and CpXP primers were P32 labeled at their 5'-ends.

PCR products were denatured in an 80% formamide loading buffer, and separated in 6% denaturing polyacrylamide gel following by overnight exposition to X-ray films at −70°C.

Molecular data analysis

The banding patterns, identified in both KpnI/MseI and Acc65I/MseI AFLP platforms, were juxtaposed based on the common bands and treated as a single profile. Thus, bands missing in one platform and present in the other were also juxtaposed. This juxtaposition resulted in a profile with the same number of markers in both platforms (Bednarek et al. 2007). The AFLP patterns were recorded as 0-1 binary matrices, where “1” indicates the presence and “0” the absence of a given fragment following by statistical analysis. The methylation restriction points for Acc65I/MseI were calculated as a number of methylated loci (bands present on the KpnI/MseI platform and absent on Acc65I/MseI platform) divided by the total number of bands expressed by percentage. GenAlex 6.2 software was used to calculate the number of AFLP fragments. Measures of genetic uniformity among regenerant samples were determined using the Jaccard dissimilarity coefficient. Genetic XLStat v.2009.04 excel add-in software was used to draw the dendrogram (UPGMA, Jaccard) and to perform principle coordinate analysis (PCoA).

### Table 1 Adapters and metAFLP oligomer sequences

| Adapters/primers | Sequence 5’ → 3’ |
|------------------|------------------|
| Acc65I adapter 1 | CTC GTA GCA TGC GTA CA |
| Acc65I adapter 2 | GTA CTG TAC GCA TGC TAC |
| KpnI adapter 1   | CTC GTA GCA TGC GTA CAG TAC |
| KpnI adapter 2   | TGT ACG CAT GCT AC |
| MseI adapter 1   | TAC TCA GGA CTC ATC |
| MseI adapter 2   | GAC GAT GAG TCC TGA G |
| Acc65I/KpnI pre-selective primer | GCA TGC GTA CAG TAC C |
| MseI pre-selective primer | GAT GAG TCC TGA GTA AC |
Results

PEM survival and suspension culture recovery after short-term and long-term cryostorage

Determined by TTC test, the survival of PEM stored in LN for 48 h and for 1.5 years reached 82% (Table 2). Cryotreatments and liquid nitrogen exposure did not change the ability of the tissue to culture recovery. The biomass growth, after the 48 h and 1.5 years of PEM cryostorage, was maintained on the level assessed for control suspension culture (Table 2). After 4 weeks of agar/liquid (2 weeks for each), post-thawing culture about 8 g fresh weight per 10 beads for each treatment were achieved.

Embryogenic capacity of 3.5-year-old cell suspension culture

Six-week-old suspension cultures, recovered after the (OD; OD + AD; OD + AD + LN) cryotreatments, were used for somatic embryo production. The productivity of non-cryotreated 3.5-year-old suspension culture was 9 somatic embryos from 50 mg PEM (Fig. 3). OD treatment increased the somatic embryogenesis process resulting in 90 somatic embryos per 50 mg PEM. Air desiccation and LN treatment did not further affect the embryogenic efficiency.

AFLP analysis of the regenerants

In total, both KpnI/MseI and Acc65I/MseI platforms generated 560 clearly visible AFLP markers shared among the sets. However, when calculations were performed using either the KpnI/MseI or the Acc65I/MseI AFLP platform, there were from 533 to 554 bands shared among the NC, SC and LC sets of regenerants (Table 3). The KpnI/MseI-based platform allowed for the identification of 499 bands appearing with a frequency greater than 5% among regenerants of the NC set, 528 among SC, and 515 among LC. In the case of Acc65I/MseI digests, 531 AFLP bands were amplified for both NC and LC regenerants, and 529 for SC set. Two new DNA fragments distinguished 98 SC regenerants from the 196 NC and LC regenerants (Table 3). Between plantlets, no changes in the methylation level were found. The percentage of methylated loci varied between 16.61% for NC and LC regenerants, and 16.88% for SC (Table 4).

The UPGMA analysis (Fig. 4) was capable of identifying a number of sub-clusters, but only two of the them encompassed most of the regenerants. One of the sub-clusters was comprised of almost all the plantlets derived from non-cryotreated and short-term cryostored tissue, while the other sub-cluster included most of the LC regenerants, independent of the AFLP platform used. The Jaccard’s dissimilarity genetic distance between the sub-clusters was 0.27 (KpnI/MseI) and 0.34 (Acc65I/MseI). Although agglomeration analyses failed to separate NC and SC sets, principle coordinate analysis (PCoA) grouped all the regenerants according to the way they were derived, and grouped the regenerants independently of the AFLP.

### Table 2 PEM survival and fresh weight growth after 4 weeks of post-thawing culture for 48 h and 1.5-year PEM cryostorage

|                     | Control (encapsulated PEM) | PEM after LN storage for 48 h | 1.5 years |
|---------------------|-----------------------------|------------------------------|-----------|
| PEM survival (%)    | –                           | 81.50 ± 9.14a                | 82.9 ± 12.01a |
| Fresh weight growth (g) | 8.14 ± 1.20a               | 8.27 ± 1.05a                | 8.03 ± 1.09a |

Values marked by the same letter are not significantly different at $P < 0.05$ (LSD test). Data were recorded from two independent experiments with nine or three replicates (10 beads in each) for assessment of PEM survival or fresh weight growth, respectively.

### Table 3 Arrangement of band pattern in KpnI/MseI and Acc65I/MseI AFLP platforms for the three sets of *G. cruciata* regenerants derived from: non-cryotreated PEM (NC), short- (SC) and long-time (LC) cryostored PEM

| AFLP fragments | Group of regenerants |
|----------------|----------------------|
|                | KpnI/MseI | Acc65I/MseI |
| No. AFLP fragments | NC | SC | LC | NC | SC | LC |
| No. AFLP fragments with a frequency ≥ 5% | 499 | 528 | 515 | 531 | 529 | 531 |
| No. new AFLP fragments | 0 | 2 | 0 | 2 | 0 |
platform used for the calculations (Fig. 5). However, this grouping was more clearly visible on the **Acc65I/MseI** (Fig. 5b) than on the **KpnI/MseI** (Fig. 5a) platform.

**Discussion**

We developed a reliable encapsulation/dehydration protocol for cryopreservation of gentian cell suspensions (Mikuła et al. 2008), which offers more than 80% survival of PEM. The protocol also provides very fast, simple, and reproducible recovery of suspension cultures during a 4-week period. Our studies showed that liquid nitrogen did not affect the re-growth of suspension cultures and did not influence the genome size in PEM and in regenerants (Mikuła et al. 2008). At present, the influence of subsequent steps of the encapsulation/dehydration procedure on the embryogenic capacity of *G. cruciata* cell suspension culture and metAFLP analysis of regenerants derived from short-term and long-term cryostored PEM were undertaken.

**Influence of cryopreservation on the embryogenic potential**

The results presented here show that the *G. cruciata* suspension culture, recovered after cryopreservation by encapsulation/dehydration, regenerated a higher number of somatic embryos than unfrozen culture. There are only a few other references to the same problem and they support our findings. Wang et al. (1994) obtained a 28% regeneration frequency from the cell clusters of *Festuca arundinacea*, while the corresponding re-established suspension culture acquired an 82% regeneration capacity after cryopreservation. Moreover, more than a twofold higher plant regeneration frequency was observed for protoplasts isolated from suspension culture recovered after cryopreservation than from non-cryopreserved one. Cryopreservation promoted embryogenesis and subsequent plant germination of *Vitis vinifera* (Wang et al. 2002). The number of embryos increased from 106 before cryopreservation to 785 after freezing. Non-frozen and frozen tissues of *Cyclamen persicum* were sources of an average of 120–300, and 470 somatic embryos per gram, respectively (Winkelmann et al. 2004).

The fact that cryopreservation results in higher regeneration rates and is probably related to the selection of certain cell types. Selection seems to point towards embryogenic (totipotent) cells. These cells possess a more dense cytoplasm and contain less water (Häggman et al. 1998; Mikuła et al. 2005c). Consequently, embryogenic cells are more resistant to dehydration and cryopreservation. The phenomenon has been observed in proembryogenic masses of gentian suspension culture (Mikuła et al. 2005c). However, we found that the maximum growth of the embryogenic capacity of the *G. cruciata* cell suspension was reached following 7 days of osmotic dehydration of encapsulated PEM, in liquid medium supplemented with rising concentrations of sucrose (from 0.3, 0.5, 0.75 to 1.0 M). Somatic embryo productivity, from suspension culture recovered after this treatment, increased ten times. Similarly, dehydration treatment enabled encapsulated embryogenic cell suspensions of *Vitis vinifera* to increase the number of regenerated embryos more than fourfold (Wang et al. 2002). For this experiment, the two-step preculture (from 0.25 to 1.0 M sucrose) was extended to 5 days. The subsequent steps of the encapsulation/dehydration method, i.e. air desiccation and freezing, did not

**Table 4** Comparison of the methylation restriction points for **Acc65I/MseI** for the three sets of *G. cruciata* regenerants derived from: non-cryotreated PEM (NC), short- (SC) and long-time (LC) cryostored PEM (±SD)

| Group of regenerants | NC   | SC   | LC   |
|---------------------|------|------|------|
| Percentage of methylated loci | 16.61 ± 0.11 | 16.88 ± 0.86 | 16.61 ± 0.42 |

**Fig. 4** Agglomeration (UPGMA, Jaccard) analysis of the NC, SC and LC sets of regenerants based on digested by **KpnI/MseI** (a) and **Acc65I/MseI** (b) AFLP platforms. The number of regenerants is given in parentheses. NC, SC, LC regenerants derived from: non-cryotreated, short- and long-term cryostored PEM, respectively.

**Fig. 5** Agglomeration (UPGMA, Jaccard) analysis of the NC, SC and LC sets of regenerants based on digested by **KpnI/MseI** (a) and **Acc65I/MseI** (b) AFLP platforms. The number of regenerants is given in parentheses. NC, SC, LC regenerants derived from: non-cryotreated, short- and long-term cryostored PEM, respectively.
cause additional, significant alterations in embryogenic capacity of the *G. cruciata* suspension culture. In contrast, for *Vitis vinifera*, the embryogenic potential increased sevenfold after cryostorage (Wang et al. 2002).

It has been reported, for example, for *Daucus carota* or *Festuca rubra*, that somatic embryogenesis can be induced on primary explants (Kamada et al. 1993; Tanaka et al. 2009) or restored to long-term cultures (Zaghmout and Torello 1992), by the treatment with high levels of sucrose. In all the studies on cryopreservation, where elevation of embryogenic capacity was noticed, a prefreezing treatment with high concentrations of sucrose (Wang et al. 2002; Winkelmann et al. 2004; Engelmann et al. 1997; Aguilar et al. 1993) or sorbitol (Wang et al. 1994) was used. However, these treatments were employed only for a short time: from 1 h (Engelmann et al. 1997) to 3–4 days (Wang et al. 1994). The treatments at the most doubled the regeneration potential. In contrast, the encapsulation/dehydration method, which requires a 5–7-day-long sucrose treatment, had 4 or 9 times more elevated somatic embryo production for *Vitis vinifera* (Wang et al. 2002) and *G. cruciata*, respectively. The study by Kikuchi et al. (2006) confirmed that the response of stress-treated explants is relatively slow. It seems that not cryostorage, but primarily the prefreezing, slow dehydration treatment, is directly responsible for changes in embryogenic capacity.
(Epi)genetic changes in regenerated plantlets

In our work, the sensitivity of detecting changes in both structure and methylation of DNA was studied by subjecting DNA preparations to digestion with restriction endonucleases, before AFLP analysis. The pairs of isochisomers used were different in their sensitivity to methylation. We focused on monitoring the genetic uniformity of the 288 regenerants. We found only two new unique DNA fragments at the sequence DNA level after cryopreservation by encapsulation/dehydration. Our findings are consistent with earlier works which reported infrequent occurrences of genomic DNA alterations in plantlets recovered after cryopreservation. It was observed that the cryopreservation procedure did not generate additional variability to the initial diversity (Scocchi et al. 2004). The revealed genetic instability, among three cell lines of Oryza sativa (established from different mature embryos) and the corresponding re-established-after-freezing calli, appears to be related to tissue-culture-induced somaclonal variation (Moukadiri et al. 1999). In the majority of cases, the authors cite that these changes were evoked by in vitro culture procedures (Sales et al. 2001; Peredo et al. 2009; DeVerno et al. 1999; Moukadiri et al. 1999; Wang et al. 1994). DeVerno et al. (1999) found three unique RAPD fragments in two of six clones of embryogenic tissues re-established after cryopreservation, but not in the corresponding regenerants of Picea glauca. The genetic instability of cryopreserved embryogenic cell suspension culture of Festuca arundinacea and corresponding regenerated plants was confirmed by RAPD markers, using one of 19 different primers (Wang et al. 1994). Verification of PCR reactions ascertained that the sole difference observed is most likely due to the in vitro process. In other systems of plant regeneration, i.e. from already differentiated apical meristems, the number of new DNA fragments is also very limited. Martín and González-Benito (2005), using the encapsulation/dehydration method of cryopreservation, showed different RAPD band patterns in 1 out of the 46 regenerants of Dendranthema grandiflora derived from cryopreserved shoot apex, whereas all 21 plantlets obtained by vitrification were genetically stable. Based on the AFLP markers, three extra DNA fragments were found between the control somatic embryos of Quercus suber and cryopreserved samples (Fernandes et al. 2008). These differences were not supported by SSR analysis. The relatively high levels of DNA structure changes after cryopreservation (9–14 alterations; up to 10.07%) were described for two genotypes of Carica papaya (Kaity et al. 2008). For two other genotypes, the changes reached 1.5–2.84% (2 and 5 alterations), whereas the remaining two genotypes showed no changes.

Our studies show no differences between frozen- and non-frozen-PEM-derived regenerants on the DNA methylation level. Two new AFLP fragments detected in SC regenerants, whose DNA was digested with Acc65I/MseI, may be the reflection of DNA sequence changes. The methylation status of the genomic DNA was insignificantly different and reached 16.61–16.88%. The lack of alterations in sequences of DNA methylation patterns in G. cruciata regenerants found support in the work of Johnston et al. They showed that epigenetic changes after cryopreservation may be temporary (Johnston et al. 2009). Methylation changes were observed in plant material following exposure to section, pre-culture, PVS2, and LN treatments (Kaity et al. 2008). However, occurrence and level of methylation modifications seem to be genotype or individual-dependent (Kaity et al. 2008; Channuntapipat et al. 2003; Johnston et al. 2009; Peredo et al. 2008, 2009). Altered methylation of DNA sequences have been found among the DNA of almond regenerants derived from shoot tips that were cryostored for 3 days, when compared with those that were cryostored for 24 months (Channuntapipat et al. 2003). The change was most likely due to different sub-culturing than to the cryopreservation process, because the initial explants had been taken from shoots that had been in in vitro culture for different lengths of time.

It is necessary to take into consideration that G. cruciata NC and SC regenerants originated from PEM, which, with reference to LC, was exposed to 80 parallel subcultures during 1.5 years of culture. Almost all the NC and SC plantlets were grouped into one cluster by the UPGMA analysis. Despite of putting regenerants NC, SC and LC into separate clusters by PCoA (Fig. 5), approximate similarity between them may prove high genetic uniformity which was maintained during many years of suspension cultures, and high effectiveness of a 1.5-year-long cryostorage.

In conclusion, the present studies indicate the positive effect of cryotreatment on the embryogenic potential of G. cruciata suspension culture. Regenerants obtained from cryopreserved proembryogenic mass kept their genetic uniformity. Thus (1) simple and quick recovery of suspension cultures, (2) their high-embryogenic potential and (3) genetic uniformity of regenerants (irrespective of time cryostorage) are crucial factors to successful gentian cryopreservation.

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