Regular Article Detection of *Glutathione S-Transferase* gene (*GST2, GST3*) during induction of somatic embryogenesis in grape (Vitis vinifera L.)

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Glutathione S-transferases (GSTs) are an important group of multifunctional enzymes that belong to diverse multigene families. In plants these enzymes are involved in the detoxification of xenobiotic compounds, herbicide detoxification, and primary and secondary metaboplism and play an important role in plant growth and development. There are several reports that show that the expression of GST is linked to the developmental phases of somatic embryogenesis. This study highlights the detection of transcript abundances of glutathione S-transferase genes *GST2* and *GST3* in the process of somatic embryogenesis of *Vitis vinifera*, and the lack thereof in non-embryogenic tissue of leaf cultures of grape (control). These results indicate that the expression of *GST2* and *GST3* could be used as a molecular signal for the identification of embryogenic cultures during the early development of somatic embryos.

Key words: Dharwad, Gene expression, Karnataka, Somatic embryogenesis, Sonaka, Tas-e-Ganesh, Thompson, transcription factor, *Vitis vinifera*.

Glutathione S-transferases (GSTs) are classified as specialized multifunctional dimeric enzymes involved in enzymatic detoxification of endo and xenobiotics (Moons, 2005: Jiang *et al.* 2010). GSTs are found in nearly all forms of life, including bacteria, fungi, yeast, insects, mammals and higher plants. Plant GSTs are divided into six

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classes, namely, phi (F), tau (U), zeta (Z), theta (T), lamba (L), and dehydroascorbate reductases (Dixon et al. 2002; Moons, 2005; Jiang et al. 2010). Among them, in plant, tau (U) and *phi* (F), classes are the most represented whereas theta and zeta ones are found in animals (Jiang et al. 2010; Galland et al. 2007). The functions of GSTs in plants are diverse and might be due to the ability to conjugate glutathione (GSH) to various targets involved in biotic and abiotic stress (Jiang et al. 2010). In addition to this, plant GSTs are also involved in plant growth and development (Gong et al. 2005; Moons, 2005; Jiang et al. 2010). GSTs have been found to bind hormones such as auxin and cytokinin (Zettl et al. 1994; Gonneau et al. 1998), and can induced by а wide variety be of phytohormones, including ethylene, auxin, methyl jasmonate, salicylic acid, and abscisic acid (Wagner et al. 2002; Moons, 2003; Smith et al. 2003). All these hormones regulate many aspects of plant development which implies that plant GSTs may play vital roles in plant growth and development as well (Jiang et al. 2010). GST genes can be induced by various exogenous factors such as pathogen attack, heavy metals, heat shock, wounding or auxin (Singla et al. 2007; Jiang et al. 2010).

Somatic embryogenesis (SE) is the developmental phenomenon where somatic cells under in vitro conditions under the influence of external growth regulators are programmed towards the embryogenic pathway and reflects cellular totipotency in higher plants (Konar and Nataraja, 1965; Nataraja and Konar, 1970; Malabadi and van Staden, 2003; Malabadi et al. 2004, 2005, 2010; Malabadi and Nataraja, 2006a, 2007a, 2007c; Malabadi, 2006; Malabadi and van Staden, 2005a, 2005b, 2005c, 2006; Malabadi et al. 2009abc; Malabadi and Teixeira da Silva, 2011; Feher et al. 2003; Namasivayam, 2007; Aronen et al. 2007, 2008; Malabadi et al. 2011a-2011g; Malabadi et al. 2012ab; Mulgund et al. 2012). Acquisition of embryogenic competence largely relies on dedifferentiation information must be erased or altered to make the cells responsive for new signals (Dudits et al. 1991, 1995; Pasternak et al. 2002; Feher et al. 2003; Namasivayam, 2007; Chugh and Khurana, 2002; Singla et al. 2007; Malabadi and van Staden, 2003; Malabadi et al. 2004, 2005, 2010; Malabadi and Nataraja, 2006a, 2007a; Malabadi, 2006; Malabadi and van Staden, 2005a, 2005b, 2005c, 2006; Feher et al. 2003; Namasivayam, 2007; Aronen et al. 2007, 2008; Malabadi et al. 2011a-2011g; Malabadi et al. 2012ab; Mulgund et al. 2012). developmental The switch from а differentiated and resting cell state to a dedifferentiated, dividing, embryogenic state may involves the general reorganization of chromatin structure, overall reprogramming of gene expression, as well as cellular metabolism (Dudits et al. 1991, 1995; Pasternak et al. 2002; Feher et al. 2003; Namasivayam, 2007; Konar and Nataraja, 1965; Nataraja and Konar, 1970; Chugh and Khurana, 2002; Singla et al. 2007). Such programmes developmental involve activation of various signal cascades and differential gene expression, which confers the ability to manifest the embryogenic potential on somatic cells (Chugh and Khurana, 2002; Singla et al. 2007; Malabadi and Nataraja, 2007b; Malabadi et al. 2011a-2011g). There are many reports on profiling of expressed genes during the somatic embryogenesis pathway (Feher et al. 2003; Malabadi et al. 2011a; Malabadi and Nataraja, 2007b). Arrays of genes that are activated or somatic differentially expressed during embryogenesis process have been identified in many plant species (Zimmerman, 1993; Chugh and Khurana, 2002; Feher et al. 2003; Ikeda et al. 2006; Raghavan, 2006; Quiroz-Figueroa et al. 2006; Malabadi et al. 2011a; Malabadi et al. 2009b; Malabadi and Nataraja, 2007b; Park et al. 2009).

The transcripts of GST were detected in abundance during auxin induction and in somatic embryos (Galland *et al.* 2007; Singla *et* al. 2007). A GST gene (CHI-GST1) was specifically expressed in leaf tissues of the chicory embryogenic responsive genotype "474" from the third day of the induction phase, when the first reactivated cells were observed but not in a non-embryogenic responsive genotype (Galland et al. 2007). GST transcripts have been shown to accumulate in Chicorium (Galland et al. 2001), Medicago sativa (Thibaud-Nissen et al. 2003), Triticum aestivum (Singla et al. 2007), Cyclamen persicum (Rensing et al. 2005; Winkelmann et al. 2006; Hoenemann et al. 2012), cotton (Zeng et al. 2006), and Crocus sativus (Sharifi et al. 2012) somatic embryos and GSTs appears to be a major regulator of the interacting genes sequenced in the present case in response to auxin (Galland et al. 2007). Some GSTs are induced by auxin (Nagata et al. 1994), H₂O₂ (Levine et al. 1994), and might target transcription factors like WRKY (Du and Chen, 2000; Singla et al. 2007), and transport certain gene products produced during oxidative stress to the vacuole (Marrs 1996; Edwards et al. 2000; Singla et al. 2007). Reactive oxygen species (ROS) have been shown to act as second messenger during auxin and stress-induced embryogenesis (Nagata et al. 1994; Maraschin et al. 2005; Singla et al. 2007). These facts suggest that the GST genes could be active to attain the embryogenic competence in different plant system (Ikeda et al. 2006; Giroux and Pauls, 1997; Chugh and Khurana, 2002; Singla et al. 2007). This study aims to detect the transcript abundances of Vitis vinifera glutathione Stransferase genes GST2, GST3 in the process of somatic embryogenesis of grape during the crucial step of transfer of embryogenic cultures on to maturation medium which triggers the realization of embryo development.

Materials and methods Induction of embryogenic tissue

Leaf explants were harvested from 6-7- month- old mother plants of three *V*. vinifera L. cvs: Thompson, Sonaka and Tas-e-Ganesh. These were carefully washed in double distilled water (DDW) (Malabadi et al. 2010). They were surface decontaminated sequentially with 0.1% streptomycin (1 min), 70% (v/v) ethanol (5 min) and 0.1% (w/v) HgCl₂ (2 min) (Sigma-Aldrich, St. Louis, USA), and thoroughly rinsed with sterilized double distilled water. Leaf sections were cultured on Nitsch and Nitsch (1969) NN basal medium with 3.0% sucrose, 0.7% agar, *myo*-inositol, 1.0 gl⁻¹ casein 0.5 gl-1 hydrosylate, 0.5 gl⁻¹ L-glutamine, 250 mgl⁻¹ peptone, 0.2 gl-1 p-aminobenzoic acid, and 0.1 gl-1 biotin, all purchased from Sigma (Malabadi et al. 2010). The medium was supplemented with 4.54 µM thidiazuron (TDZ) and 2, 4-dichlorophenoxy acetic acid (2, 4-D) at a concentration of 4.52 μ M singly without any other growth hormones (Malabadi et al. 2010). The cultures were raised in 25 mm × 145 mm glass culture tubes (Borosil, Mumbai, India) containing 15 ml of the above basal medium under cool white fluorescent light (Mysore lamps, India) at 100 μ mol m⁻² s⁻¹ and 25 ± 3°C with a relative humidity of 55-60% (Malabadi et al. 2010). The pH of the media was adjusted to 5.8 with 1 N NaOH or HCl before agar was added. Media without TDZ and 2, 4-D served as the control. The media were then sterilized by autoclaving at 121°C at 1.04 Kg cm-2 for 15 min. L-glutamine, biotin, and *p*-aminobenzoic acid were filter sterilized (Whatman filter paper, pore size = $0.45 \mu m$; diameter of paper = 25 mm) and added to the media after autoclaving when the medium had cooled to below 50°C (Malabadi et al. 2010).

The cultures were examined for the presence of different developmental stages of somatic embryos by morphological and cytological observations of callus (Malabadi *et al.* 2010). The cultures showing different cell divisions were identified and subcultured on the initiation medium for further 6 weeks for the better development of early stages of somatic embryogenesis (Malabadi *et al.* 2010).

The full strength inorganic salts NN (Nitsch Nitsch, and 1969) basal medium supplemented with 4.52 µM 2, 4-D and 4.54 μ M TDZ (induction medium) was used as an effective induction medium for producing the embryogenic tissue (Malabadi et al. 2010). Embryogenic tissue showing different cell divisions such as 2 to 8 celled stages was identified using microscopic observation (Malabadi et al. 2010). On the other hand the callus without pro-embryonic cell divisions was considered as non-embryogenic. Nonembryogenic tissue was separated immediately from the rest of the tissue to avoid the overgrowth of the tissue (Malabadi et al. 2010). The efficiency of plant growth regulators and their concentrations were analyzed on the basis of visual observation (callusing percentage, percentage of explants forming embryogenic tissue, callus growth callus necrosis). and The ineffective treatments were discontinued as previously reported (Malabadi et al. 2010).

Maintenance of embryogenic tissue

The embryogenic tissue of three of V. vinifera L. cvs: Thompson, sonaka and Tas-e-Ganesh showing various developmental stages of somatic embryos was maintained on full strength inorganic salts NN (Nitsch and Nitsch, 1969) basal medium supplemented with 4.52 μ M 2, 4-D and 4.54 μ M TDZ for the proliferation of callus (maintenance medium) (Malabadi et al. 2010). The embryogenic tissue was subcultured for every 4 weeks. All the cultures were maintained under a cool white fluorescent light (100µmol m-2 s-1) at 25±3°C with a relative humidity of 55-60%. The percentage of cultures showing somatic embryogenesis was recorded (Malabadi et al. 2010).

RNA preparation and cDNA synthesis

Frozen samples of embryogenic tissue derived from leaf explants of three of *V*. *vinifera* L. cvs: Thompson, Sonaka and Tase-Ganesh and control (non-embryogenic tissue)

were ground in a mortar and pestle with liquid nitrogen. Total RNA was isolated according to the modified method of Chang et al. (1993). To remove residual genomic DNA, 25 µg of RNA was treated with TURBO-DNase[™] (Ambion, Austin, TX, USA) (Malabadi et al. 2011; Park et al. 2009). cDNA was generated from 1 µg of DNase-treated RNA using the Superscript II RT system (Invitrogen, CA, USA) according to the manufacturer's protocol. Each reaction was run in duplicate, generating two independent cDNA samples for each RNA sample isolated from three V. vinifera L. cvs: Thompson, sonaka and Tas-e-Ganesh (Malabadi et al. 2011; Malabadi and Nataraja, 2007g).

Cloning of grape (*Vitis vinifera*) glutathione S- transferase gene (*GST2*, *GST3*)

For cloning of grape (Vitis vinifera) glutathione S-transferase genes GST2, GST3, gene specific primers were designed using Primer 3.0 software based on homologous sequences publically available on NCBI databases (www.ncbi.com). The nucleotide sequences of Vitis vinifera glutathione Stransferase gene GST2 (Gene Acc No-AY156049), Vitis vinifera glutathione Stransferase gene GST3 (Gene Acc no-AY156050) were selected from the NCBI databases respectively. The gene specific primers (GST2: Forward: 5' AGC TCT TTG ACT 3'; Reverse: 3' TTA ACT CTT CTG CAT 5' and GST3; Forward: 5'GAT TGA GGA GAG GA 3'; Reverse: 5' TCA CTC CAA GAG GGG CCA T3') were custom synthesized (Bangalore Genei, Bangalore, Karnataka, India) and used for the PCR amplification of the GST2 and GST3 genes using cDNA from embryogenic tissue and control (nonembryogenic tissue) template. as the Amplified fragments from grape (Vitis vinifera) cDNA samples were excised from the agarose gel and purified with QIA quick gel extraction kit (Qiagen, USA). The eluted PCR product was cloned using the TOPO-TA cloning kit (Invitrogen, USA) (Malabadi et al.

2011; Park *et al.* 2009). Ligated products were then transformed into chemically competent TOPO 10 *Escherichia coli* cells using heat shock method and the transformants were selected on LB agar plates containing kanamycin (100 μ g ml⁻¹) (Malabadi *et al.* 2011; Park *et al.* 2009). The expression of *Vitis vinifera* glutathione S-transferase *GST2* and *GST3* genes during induction of somatic embryogenesis were confirmed by the DNA sequence analysis.

Somatic embryo maturation

The embryogenic tissue of three V. vinifera L. cvs: Thompson, Sonaka and Tas-e-Ganesh showing different developmental stages of somatic embryos, was transferred to maturation medium to induce cotyledonary embryo development (Malabadi et al. 2010). The full strength (inorganic salts) NN (Nitsch 1969) and Nitsch, basal medium supplemented with 3.0% sucrose, 5 µM ABA and 0.8% agar (maturation medium) was tested for this purpose (Malabadi et al. 2010). All the cultures were again maintained in the dark for 4 weeks. Microscopic observation was conducted to ensure the development of somatic embryos. The total number of somatic embryos produced after 8 weeks on maturation medium per one gram fresh weight of embryogenic tissue was recorded (Malabadi et al. 2010).

Germination and recovery of plantlets

After maturation, the cotyledonary somatic embryos were taken from the cultures for germination (Malabadi *et al.* 2010). The germination medium used was half strength (inorganic salts) NN (Nitsch and Nitsch, 1969) basal medium with 0.7% agar without any growth regulators (germination medium). Somatic embryos were considered germinated as soon as radical elongation occurred and conversion to plantlet was based on the presence of an epicotyl (Malabadi *et al.* 2010). After 4 weeks on germination medium, the plantlets were directly transferred to vermiculite. Plantlets were placed in a growth room under a 16 hr photoperiod (50μ mol m⁻² s⁻¹) for hardening. Somatic embryo proliferation in terms of root, shoot development, plant conversion was recorded (Malabadi *et al.* 2010).

Results and discussion

In the present study, leaf explants induced embryogenic tissue after 2-4 weeks of culture on the full strength inorganic salts NN (Nitsch and Nitsch, 1969) basal medium supplemented with 4.52 µM 2, 4-D and 4.54 μM TDZ (induction medium) in all the three varieties of V. vinifera (Malabadi et al. 2010). Embryogenic areas were clearly visible from the rest of the callus by their globular and glazy appearance and emerged as distinct white glossy structures. The embryogenic tissue was separated from non-embryogenic tissue for the gene expression studies. In a control study, the leaf explants did not promote callus formation (Malabadi et al. 2010). Leaf explants remained green for two weeks and, eventually turned brown and necrosed. Therefore, 4.52 µM 2, 4-D and 4.54 µM TDZ are the optimum concentrations for the induction of embryogenic tissue in all the 3 tested varieties of V. vinifera (Malabadi et al. 2010). Total RNA was isolated from both control (non-embryogenic tissue) and embryogenic tissue. Furthermore, cDNA preparation was done and used as the template for the identification of Vitis vinifera glutathione S-transferase genes GST2, GST3 expressed during somatic embryogenesis pathway. In our present study, we used the embryogenic cultures for the gene expression studies from the induction phase which showed the early sign of somatic embryogenesis. Microscopic observation of the callus revealed the abundance of two, four and eight celled stages often mixed with the globular and heart shaped embryos. Our results clearly demonstrated the GST transcript abundances in the embryogenic cultures showing clear evidence of GST role in inducing somatic embryogenesis in grapes.

Furthermore, we can not pinpoint whether the GSTs gene expressions are stage specific or might be up-regulated or down regulated. A detailed study of suppression subtractive hybridization (SSH) library method should be used to study the role of different genes during different stages of somatic embryo development in grape, which could give a clear picture of the role of GST and other genes in down or up regulation.

A number of studies have been published, in which gene expression profiling includes GST has been used to analyse the process of somatic embryogenesis by [(Hoenemann et al. 2010 in . in Glycine max; (Thibaud-Nissen et al. 2003 in , Pinus roxburghii; (Malabadi et al. 2011a; Malabadi and Nataraja, 2007 in, Picea abies (Stasolla et al. 2004 in, Oryza sativa; (Su et al. 2006 in Zea mays (Che et al. 2006 in Gossypium hirsutum; (Zeng et al. 2006; Wu et al. 2009 in Cichorium intybus (Legrand et al. 2007 in Triticum aestivum; (Singla et al. 2007 in Elaeis guineensis (Lin et al. 2009). The possible involvement of GSTs in morphogenesis has been well documented on microspore and somatic embryogenesis (SE) (Gong et al. 2005). In barley, a GST homolog, ECGST, was not detected in freshly isolated microspores accumulated transcript but the in microspores undergoing early stages of embryogenesis (Vrinten et al. 1999; Gong et al. 2005). A GST gene CHIGST1 isolated from chicory has been shown to express in

cultured leaves of embryogenic cultivar forming somatic embryos but not in the cultured tissues of non-embryogenic cultivar (Galland *et al.* 2001; Gong *et al.* 2005). Furthermore, a GST homolog Dcarg-1 has been isolated in carrot during SE and its expression has been associated with somatic embryo formation (Kitamiya *et al.* 2000; Gong *et al.* 2005). Although these findings indicate the possible role of GSTs in SE, it remains to be substantiated (Gong *et al.* 2005). Results of this study provids direct evidence showing that GST is involved in shoot morphogenesis in vitro (Gong et al. 2005). This has been demonstrated in culture responses of transgenic plants, where down regulation of GST expression by RNAi in GST-DS1 markedly decreases the shoot regeneration capacity of the cultured tissues (Gong et al. 2005). In contrast, over expression of GST promotes regeneration of GST-S6 (Gong et al. 2005). Furthermore, regulatory role of GSTs in shoot regeneration and SE may be related to ethylene production (Gong et al. 2005). The regulatory role of GST in shoot regeneration may be attributed, in part, to stress because GST expression can be up regulated in response to stress as demonstrated in this and previous studies (Gong and Pua, 2004; Gong et al. 2005). GST genes are not only responsive to auxin but are induced by other hormones, e.g. ABA, mJA, and under various biotic and abiotic stresses, and may have a possible role in detoxifying excessive amounts of auxin, thus regulating the intracellular concentration or its inactive analogs (Singla et al. 2007). In our present study, the expression of GST in abundance during auxin induction and in somatic embryogenesis suggests that the GST genes could be active to attain the embryogenic competence in the grape. Thus, the presence of GST trascripts representing potent markers of somatic embryogenesis and involved in cross-talk between auxin and SE, clearly suggest their involvement in the initiation phase of somatic embryogenesis in the grape leaf base system. In one of the study reported by Galland et al. (2007), a GST gene (CHI-GST1) expression was linked to the early stages of somatic embryogenesis of *Cichorium*. Therefore, a GST gene (*CHI-GST1*) was up-regulated specifically in leaf tissues of the embryogenic responsive Cichorium '474' from the third day of culture (Galland et al. 2007). They have also mentioned that the localization of GST transcripts and proteins in Chicory leaf sections showed that during cell reactivation, i. e. from G_0 to the end of G_2 phases, GSTs were strictly associated with preparation embryogenic for mitosis (Galland *et al.* 2007). Therefore, GSHdependent developmental pathway was essential for initiation and maintenance of cell division. Further GST transcripts were not observed in a similar manner in developing somatic embryos of *Chicory* (Galland et al. 2007). The presence of GST transcripts and proteins in reactivated cell and multicelled embryos indicated that antioxidation mechanisms were active during precocious phases of SE in Chicory. Therefore, GST transcripts could be taken as one of the molecular marker during the induction of somatic embryogenesis in plants. This study provides the direct evidence that GSTs are involved in regulation of vegetative growth and somatic embryogenesis in plants.

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