



Evaluation of somatic embryo production during embryogenic tissue proliferation stage using morphology, maternal genotype, proliferation rate and tissue age of *Pinus thunbergii* Parl

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Abstract To determine the optimal embryogenic capacity (somatic embryo production) of the selected elite nematode-resistant genotypes of *Pinus thunbergii*, variables such as embryogenic tissue (ET) morphology, maternal genotype, proliferation rate and tissue age were analyzed. ET morphology and histological evaluation of the proliferation stage showed a decrease in filamentous clump and protuberant surfaces and a decline in the acetocarmine-staining area, which indicates a decrease in somatic embryo production (SEP). Variations in cell physiology during the proliferation stage showed that SEP was positively correlated with soluble sugars and proteins, but negatively correlated with

starch, peroxidase, and superoxidase. In addition, SEP was significantly ($p < 0.001$) affected by maternal genotype, tissue age and proliferation rate. Moreover, SEP was positively correlated with proliferation rate ($r = 0.98$, $p < 0.001$), but negatively correlated with tissue age ($r = -0.95$, $p < 0.001$). In general, the results suggest that SEP could be assessed in ET proliferation stages by the apparent cell morphology, histology, proliferation rate and tissue age, which provides novel insights for evaluating the ET maturation capacity (number of somatic embryos) during the proliferation stage of *P. thunbergii* somatic embryogenesis.

Keywords Somatic embryo production · Cell physiology · Cell morphology · *Pinus thunbergii*

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Introduction

Somatic embryogenesis may be used as a tool to increase the quality and productivity of elite genotypes. Many efforts have been made to optimize somatic embryogenesis systems (Jones and van Staden 2001; Finkelstein et al. 2008; Herve et al. 2013; Jain and Gupta 2018); however, for many *Pinus* species, maturation with high somatic embryo yields is an important limiting factor for practical applications (Yildirim et al. 2006; Klimaszewska et al. 2011). Similarly, somatic embryogenesis of *Pinus thunbergii* Parl. has been reported since 1999, but not implemented on a commercial scale (Maruyama and Hosoi 2016). One of the major limiting factors for commercial application of somatic embryogenesis is the poor maturation efficiency (quantity of cotyledonary somatic embryos per gram fresh mass [SEs g⁻¹ FW]) (Lelu-walter and Paques 2009; Song et al. 2018). In this study, we evaluated somatic embryo

production during the embryogenic tissue proliferation stage and established early-stage indicators to select high-yielding cell lines for practical applications. These indicators were beneficial in improving the maturation efficiency of elite trees, especially for pine wood nematode-resistant *P. thunbergii*. Maruyama et al. (2005) reported a protocol for somatic embryogenesis of *P. thunbergii*, and, our group established a somatic embryogenesis protocol for *P. thunbergii* resistant to pine wood nematode and selected a maturation medium that was optimal for most cell lines of pine wood nematode-resistant *P. thunbergii* (Sun et al. 2019a, b). However, our ability to predict the performance of selected elite embryogenic tissue (ET) or maturation capacity during early stages was still limited. Krajnakova et al. (2009) reported embryogenic mass proliferation was negatively correlated with somatic embryo production (SEP). However, a positive correlation between embryogenic mass growth and SEP was reported in stone pine (Carneros et al. 2017). In addition, correlations between ET morphology and SEP have been reported. For example, ET surfaces that have a “spiky” morphology (like the filamentous and protuberant surfaces of ETs in our study) are more productive than those having smooth surfaces (Ramarosandratana et al. 2001). Additionally, ET characterized by special morphology (embryonal-suspensor mass [ESM] structures) produced the greatest maturation yield in *Pinus* species (Breton et al. 2005; Vestman et al. 2010). These traits can be used to assess the ET maturation capacity before the maturation process of somatic embryogenesis. Bradai et al. (2016) considered genotype a key factor that determines the quantity of SEs present during the maturation stage. Gahan and George (2008) reported that although regulatory mechanisms controlling regeneration became inoperative during subcultures, the genotype of embryonal cells did not essentially change. In contrast, Park et al. (2011) found that the embryogenic capacity and DNA content in the nucleus decreased during prolonged maintenance. In general, the genotype has to be carefully chosen because not all ETs grow well in culture (Montalban et al. 2012; Zhang et al. 2014; Malik et al. 2020). In addition, whether the maturation of SEs is correlated with morphological development and intracellular communication has also been considered (Godel-Jedrychowska et al. 2020). For practical applications, the selection of a suitable ET with high-yielding somatic embryos (SEs) during early stages is needed. Here, we studied the effects of ET morphology and physiology, maternal genotype, tissue age and proliferation rate on the SEP of *P. thunbergii*, and we established criteria to predict the ET maturation capacity before maturation of the somatic embryo. The aim was to select the ET having high-yielding SEs in the proliferation stage for practical applications.

Materials and methods

Plant material

Immature cones were collected from the pine wood nematode-resistant *P. thunbergii* in an experimental field at Nanjing Forestry University (Jiangsu, China). The mother trees were open-pollinated in a seed orchard. Tissue cultures were started in 2015 and 2016 (Sun et al. 2015; Wang et al. 2019). The immature cones were stored at 4 °C for 1 week, then female gametophytes were selected. The female gametophytes contained immature zygotic embryos when the explants were transplanted onto the initiation medium. The initiated ETs (cell lines) were transferred onto maintenance medium and numbered. The cell line number comprised the ET initiation time, seed family and clone, with the first two numbers representing the ET induction time (year after 2000), the last two numbers representing the seed family, and the number after the dash representing the clone. In this experiment, 12 established cell lines were used in total. Five selected ETs or cell lines (a clump of ET about 1 cm in diameter) were transferred onto maintenance medium in each Petri dish. The ETs were subcultured on maintenance medium every 2 weeks. The maintenance medium was DCR basic medium (Gupta and Durzan 1985), supplemented with 250 mg L⁻¹ 2-(4-morpholino) ethanesulfonic acid (MES), 15 g L⁻¹ maltose, 1 g L⁻¹ *myo*-inositol, 500 mg L⁻¹ casein acid hydrolysate, 500 mg L⁻¹ l-glutamine (filter-sterilized), 5.4 g L⁻¹ agar and plant growth regulators 2, 4-dichlorophenoxyacetic acid at 1 mg L⁻¹ and kinetin at 0.1 mg L⁻¹. Before sterilization, the pH of all media was adjusted to 5.8 and autoclaved (121 °C, 20 min).

Maturation of somatic embryos

The suspension culture before the maturation process was prepared according to previous reports (Sun et al. 2019b). Briefly, the suspension cultures were initiated by transferring 1 g ET to 30 mL of liquid proliferation medium in 100 mL Erlenmeyer flasks that were subsequently placed on an orbital shaker at 90 rpm. After 7 days of subculturing, the suspension cultures (2 mL) were spread onto the top of a stack of five pieces of sterilized filter paper. After the liquid medium was absorbed, the filter paper containing the cultures was placed on the surface of the solidified medium for embryo maturation. Approximately 500 mg of ET was transferred to each Petri dish. The previously described maturation medium is able to support most cell lines of *P. thunbergii* that are resistant to the pine wood nematode (Sun et al. 2019a). Briefly, the maturation medium was an LP basic medium (Gupta and Durzan, 1987), supplemented with 2 g L⁻¹ activated charcoal, 1 g L⁻¹ *myo*-inositol, 1.5 g L⁻¹ l-glutamine (filter-sterilized), 3.5 g L⁻¹ phytagel (Sigma-Aldrich,

St. Louis, USA), 30 g L⁻¹ maltose, 10 mg L⁻¹ abscisic acid and 125 g L⁻¹ PEG8000. The ET was maintained in the dark at 25 ± 1 °C for 10 weeks. After 10 weeks of culturing (without interval), the mature SEs (cotyledon embryo stage) were moved from maturation to germination medium, and the number of mature SEs was recorded. Then, the number of mature SEs per gram fresh mass (SEP) was calculated. The SEP was used as the criterion for evaluating the maturation capacity of the ET.

Maturation experiment 1: morphology and histological development

Morphology characteristics

Samples (10 replicates per tissue age) were taken from 3, 6, 12 and 24-month-old ETs of cell line 1537-1, 10 days after subculturing on fresh maintenance medium, and then made into cell suspensions. Briefly, 20 µL suspension was transferred to a slide, stained with 1% w/v acetocarmine for 30 s, then with Evans blue (Gaff and Okong'o-Ogola 1971). A cover glass was placed on the slide and the slide washed. The samples were observed with a stereomicroscope (Stemi 508, Germany) for stereological analysis.

Determination of intracellular substances

ETs (2 g) from cell line 1537-1 were ground at 4 °C in 1.8 mL of 100 mM cool phosphate buffer (pH 7.4), supplemented with 1 mM EDTA and 2% w/v polyvinylpyrrolidone polymer (PVPP). The mixtures were then centrifuged at 10,000 × g for 10 min. The supernatant was used to estimate activity of superoxidase (SOD), peroxidase (POD) and catalase (CAT). Three replicates were prepared to assay each enzyme.

SOD activity was assayed using 200 µL of the extract in 4 mL of reagent containing 50 mM phosphate buffer (pH 7.4), 0.1 mM EDTA, 10 µM nitroblue-tetrazolium (NBT), 10 mM L-methionine, and 0.001 mM riboflavin. After 0.5 min, absorbance was measured at 560 nm for 10 min. One SOD enzyme unit was defined as the quantity of SOD required to produce a 50% inhibition in the reduction of NBT. SOD activity was expressed as units per milligram protein per gram FW. SOD activity was assayed as described by Beauchamp and Fridovich (1971).

POD activity was measured as described by Britton and Maehly (1955). The 200 µL extract added in 3 mL of the reaction solution which containing guaiacol (50 mM), 50 mM phosphate buffer (pH 6.8), H₂O₂ (2%). After 0.5 min, absorbance at 470 nm was measured for 4 min. One unit of enzyme was expressed as µmoles per milliliter of H₂O₂ reduced per minute.

CAT activity was measured as described by Bergmeyer (1970). The reaction mixture (3 mL) contained 200 µL extract, 50 mM phosphate buffer (pH 7.0) with 0.001 M EDTA and H₂O₂ (3% w/v). After 0.5 min, absorbance at 240 nm was recorded for 4 min to determine the amount of H₂O₂ reduced per minute. One unit of CAT activity was defined as one µmole H₂O₂ destroyed per minute.

Starch and endogenous soluble sugars were measured using a phenol-sulfuric acid method (Dubois et al. 1956). Samples of 2 g (with three replicates) were ground in deionized water, then centrifuged at 10,000 × g for 15 min at 4 °C. The pellet was dried, weighed and heated for 10 min. The supernatants were treated with 5% v/v phenol and 98% v/v sulfuric acid, and absorbance at 485 nm was recorded.

Total soluble proteins were measured using the Bradford (1976) method; 2 g callus (three replicates) was ground with 1.5 mL phosphate buffer (pH 7.0), then centrifuged at 10,000 × g for 10 min at 4 °C. The supernatant was discarded, and the pellet was washed with acetone, then dissolved in 1.0 mL of 0.1 M NaOH. Absorbance was then measured at 595 nm.

Maturation experiment 2: genotype, tissue age and proliferation rate

The effects of genotype, tissue age and proliferation rate on the production of SEs were evaluated using cell lines from seed families 36, 37 and 39 of pine wood nematode-resistant *P. thunbergii* and analyzing the mature SEs produced on the maturation medium.

The effects of genotypes on SEP were analyzed using 2 mL of a cell suspension (approximately 500 mg FW) generated from approximately 6-month-old ETs of the respective cell lines and transferred to the maturation medium. Four replicates were prepared per genotype.

For determining proliferation rates, two cell lines (1537-1 and 1637-2) with the highest SEP initiated in 2015 and 2016, respectively, were mainly experimented. Fresh ET (1 g FW on maintenance medium) from 3-, 6-, 12-, 15-, 18-, 21- and 24-month-old ETs was transferred onto the maintenance medium, with four replicates per culture age. The ETs were subcultured on maintenance medium every 2 weeks. The proliferation rate was determined after 2 weeks.

For analyzing the effect of tissue age (the total culture time of ETs on maintenance medium) on SEP, the two cell lines (1537-1 and 1637-2) with the highest SEP initiated in 2015 and 2016, respectively, were mainly used. A 2-mL cell suspension of each cell line was generated from 3-, 6-, 12-, 15-, 18-, 21- and 24-month-old ETs and transferred to the maturation medium. Four replicates were prepared per culture age. The number of mature SEs was recorded for each line after 10 weeks.

Statistical analyses

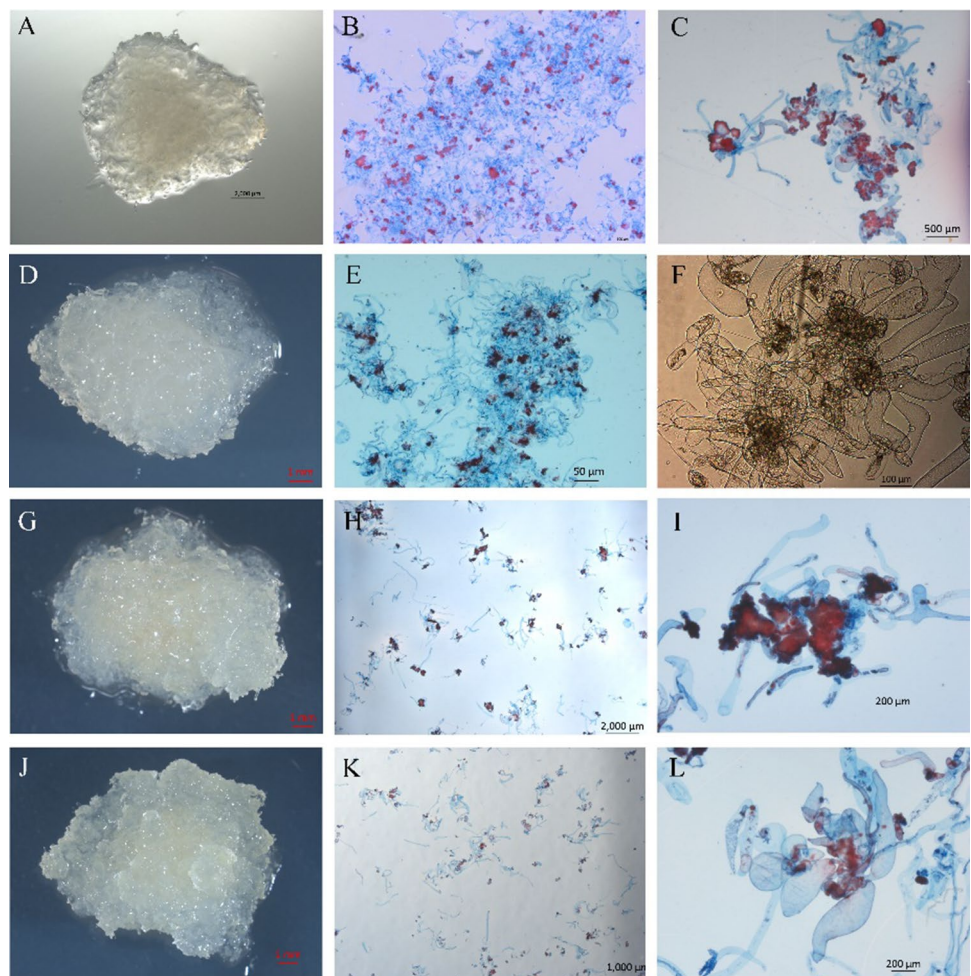
SEP after the various treatments was evaluated using an ANOVA and Pearson's correlation coefficients. Means were compared using Duncan's multiple range test with significance set at $p < 0.05$. All analyses were done using SPSS version 19 (IBM, Armonk, NY, USA) and R version 3.6.0 with the R stats package (R Core Team, 2019). The proliferation rate (fold-change) was defined as (FW at the end of treatment – FW at the start of treatment)/FW at the start of treatment. The SEP was defined as the number of mature SEs on maturation medium /the FW of ETs transferred from the proliferation to maturation medium. The FW of ETs transferred from the proliferation to maturation medium was calculated as the mass of the filter containing the tissues (2 mL suspension culture) – the mass of the filter (2 mL liquid medium). The data from cell line 1537–1, which had the greatest production, was used to calculate Pearson's correlation coefficients for tissue ages (3-, 6-, 12- and 24-month-old), soluble sugar, starch, soluble protein, enzymatic activity (SOD, POD and CAT), proliferation rate and SEP.

Results

Morphological development and physiological variations in *P. thunbergii* embryogenic cells

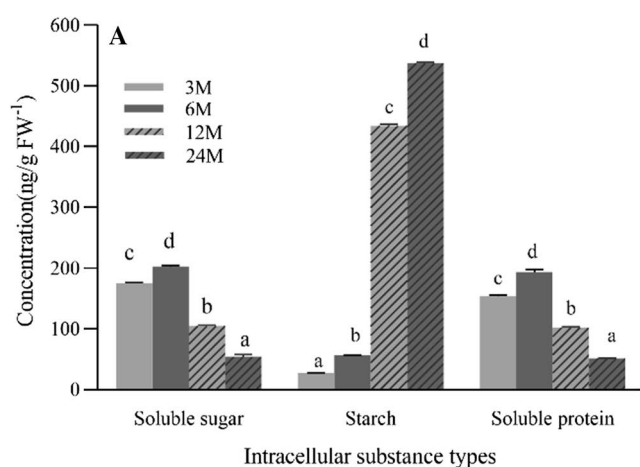
During 3 months of subculturing on maintenance medium, the filamentous and protuberant surfaces of the clumps were visible, and the ET surface was transparent and glossy (Fig. 1a). After double staining by Evans blue and acetocarmine, we found most cell types had early somatic embryonic features (embryonal suspensor mass, ESM) (Fig. 1b, c). After 6 months of subculturing, the ET transparency decreased, and the filamentous and protuberant surfaces of the clumps were reduced; however, the ET surface was still glossy (Fig. 1d). The ET still showed early somatic embryo-associated features (Fig. 1e, f). At 12 months, the filamentous and protuberant surfaces of the clumps rarely appeared, and the ET became non-transparent (Fig. 1g). In addition, a gradual decrease was observed in the red area when using acetocarmine staining, and abnormal cells were present (Fig. 1h, i). By 24 months, the filamentous and protuberant surfaces of

Fig. 1 Morphology and histological features of embryogenic cells with tissue age of *Pinus thunbergii*. **a–i**: Embryonic tissue (ET) subcultures at, **a–c** 3 months; **d–f** at 6 months; **g–i** at 12 months; **j–l** at 24 months



the clumps had disappeared (Fig. 1j), and the proportions of red areas and acetocarmine staining were significantly reduced (Fig. 1k); Further observations revealed that the consistency in cell morphology was low, as was the proportion of cells having early SE structures. Moreover, the number of abnormal cells, such as cylindrical and oval cell types, increased (Fig. 1l). The emergence of abnormal cells indicated that the embryogenic capacity of the ET had declined. In general, the ETs were young and vigorous for up to 6 months. However, by 12 months, the filamentous and protuberant surfaces of the clumps had disappeared, and the acetocarmine-stained areas and the proportion of ESMs had decreased.

In a further study of cellular physiology, the intracellular substances (soluble sugar, starch and soluble protein) ($p < 0.001$) and enzymatic activities (SOD, POD and CAT) ($p < 0.001$) differed significantly among the tissue ages (Fig. 2). The trends in soluble sugar and soluble protein were similar, first increasing and then decreasing with tissue age, and both peaked at 6 months. However, starch content increased from 27.1 ng g⁻¹ FW at 3 months to 537.1 ng g⁻¹ FW at 24 months (Fig. 2a). Among the enzymatic activities, CAT activity increased from 27.6 U g⁻¹ FW at 3 months to 47.7 U g⁻¹ FW at 12 months, then dropped to 36.7 U g⁻¹ FW at 24 months. In contrast, the POD activity showed a decreasing trend from 70.2 U g⁻¹ FW at 3 months to 50.3 U g⁻¹ FW at 6 months, then increased with tissue age. SOD activity did not differ significantly between 3 and 6 months, but it had increased significantly by 24 months (Fig. 2b). We found that cell morphology responded to variations in intracellular substances and enzyme activities as the tissue aged.



Effects of maternal genotype, tissue age and proliferation rate on the SEP of *P. thunbergii*

SEP was significantly ($p < 0.001$) affected by genotype, tissue age and proliferation rate (Figs. 3, 4). In seed family 36, SEP varied from 58 to 182 mature SEs per gram FW, and in seed family 37, the SEP varied from 354 to 1983 mature SEs per gram FW. Thus, the maternal genotypes appeared to determine the SEP level, and the SEP varied among established cell lines having the same maternal genotype. Additionally, in tests of cell lines initiated in 2015 and 2016, the highest SEP was obtained in cell line 1537-1 and 1637-2, respectively (Fig. 3a, b). The SEP and proliferation rate trends with tissue age were similar, especially in the high-yielding cell lines (Fig. 4). Thus, in cell line 1537-1, both the SEP and proliferation rate increased from 3 to 6 months, and then, they decreased along with tissue age (up to 24 months) (Fig. 4a). However, in cell line 1637-2, SEP increased from 3 to 6 months, but the proliferation rate decreased (Fig. 4b). In addition, in the rest of the cell lines, the proliferation rates all decreased with tissue age. At 24-months, the proliferation rates maintained an approximately onefold increase. Although the SEP increased from 3 to 6 months in some cell lines, it decreased or lost the maturation capacity from 6 to 24 months (Electronic supplementary material Fig. S1). In general, the SEP and proliferation rate trends decreased as the tissue age increased.

In addition, the interactions among genotypes, tissue age, proliferation rate and SEP showed that a high SEP was obtained in cell lines 1539-1 and 1537-1 at 3–6 months, when the proliferation rate was \geq threefold. When the tissue was 9–12 months old, the proliferation rate dropped to 2–threefold, and the SEP ranged from 1,762 to 86 SEs per gram FW. Moreover, when the tissue was 12–24 months old,

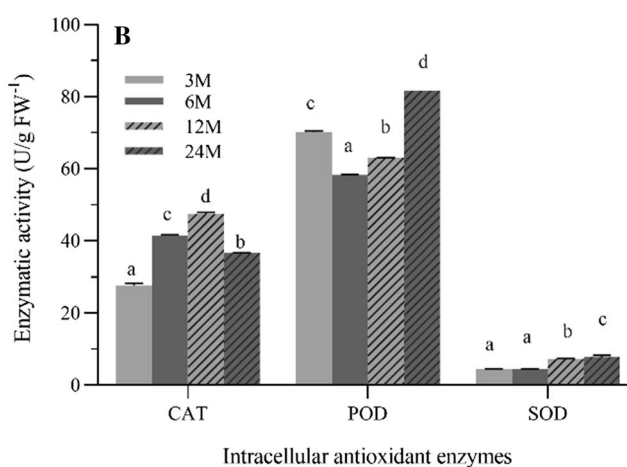


Fig. 2 Mean (\pm SE), **a** concentration of intracellular metabolites and **b** activity of enzymes from tissues of *Pinus thunbergii* at different age (3–24 months [M]). Different lowercase letters above the bars for an

indicator indicate a significant difference ($p < 0.001$) at the tissue ages using Duncan's test. CAT: catalase

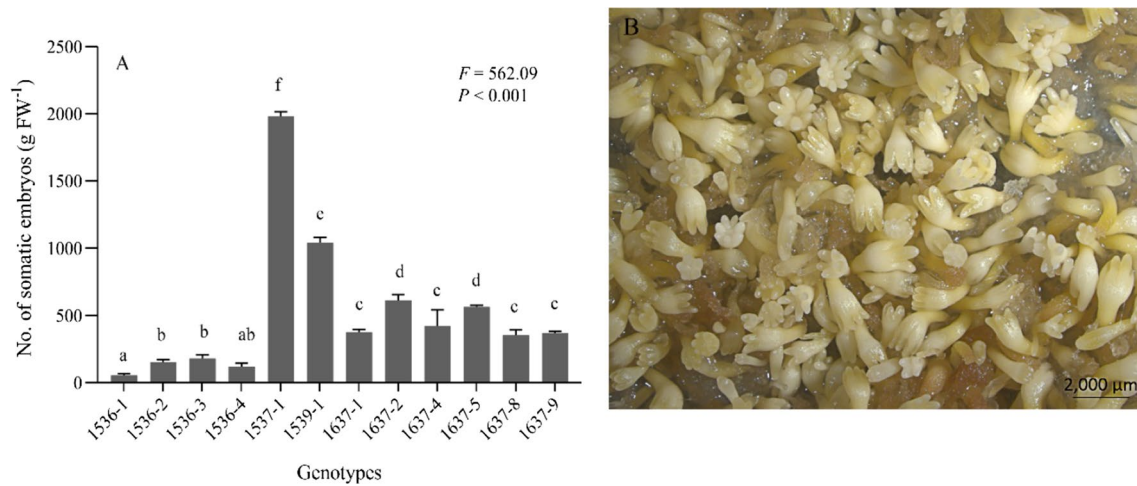


Fig. 3 Somatic embryo production by *Pinus thunbergii*. **a** Mean number (\pm SE) of somatic embryos produced by each genotype. **b** Somatic embryos produced by the most prolific cell line, 1537-1.

Different lower case letters above the bars indicate a significant difference in number produced by genotypes ($p < 0.05$) in Duncan's test. Four replicates were prepared per cell lines

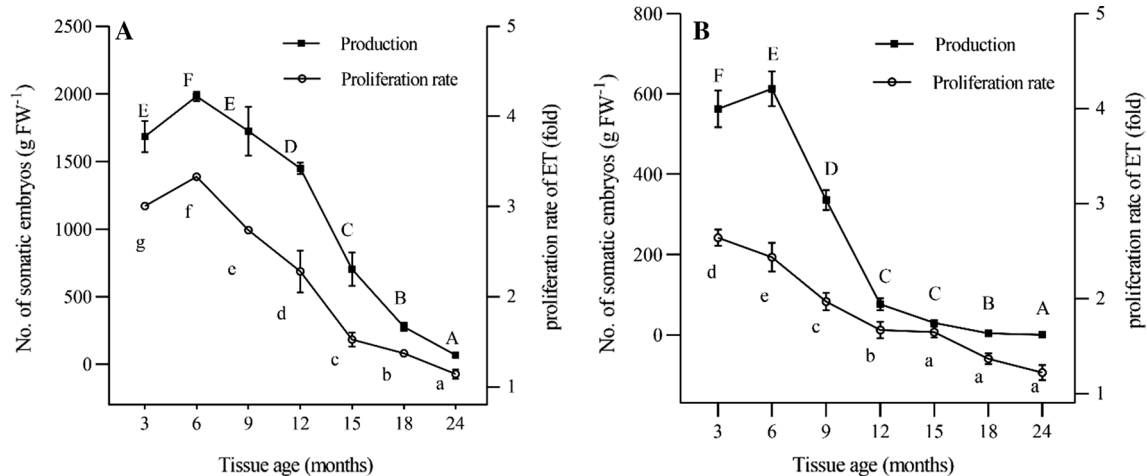


Fig. 4 Mean number (\pm SE) of somatic embryo produced and proliferation rate of embryonic tissue (ET) by cell lines, **a** 1537-1 and **b** 1637-2 at different tissue ages of *Pinus thunbergii*. Different lower case letters indicate a significant difference at a tissue age, different

upper case letters indicate a significant difference in production at a tissue age ($p < 0.001$; Duncan's test). Four replicates were prepared per cell lines

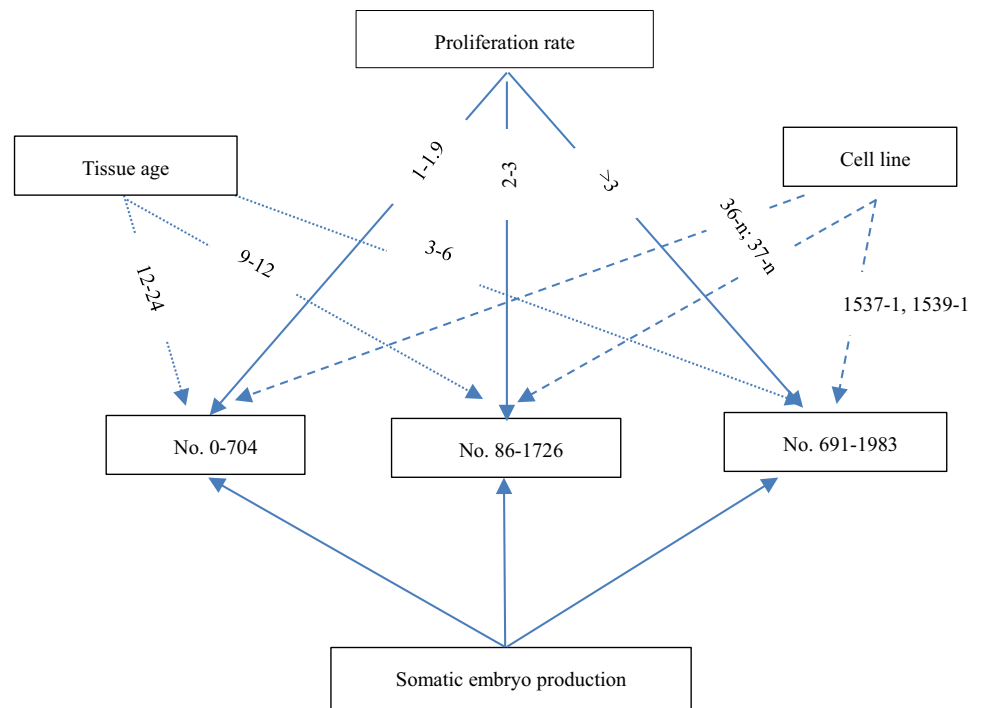
the proliferation rate dropped to 1–1.9 fold, and the SEP ranged from 704 to 0 SEs per gram FW (Fig. 5). At this time, some cell lines lost the capability to produce SEs. We found that although SEP differed among the genotypes, a higher SEP was easy to obtain from young ETs (3–6 months old), especially when the proliferation rate was \geq threefold.

Correlation analysis of SEP in *P. thunbergii*

We determined the differences in SEP between maternal genotypes and cell lines, and we found that line 1537-1 had the highest SEP among the tested cell lines. The relationships

between the high-yielding SEs and cell physiological changes (soluble sugar, starch, soluble protein and enzyme activity) were studied. A correlation analysis showed that the SEP was positively correlated with soluble sugar and protein, but negatively correlated with starch. SOD and POD activities, however, had a significantly negative correlation with SEP, but CAT was not correlated with SEP. The relationships among enzyme activity levels showed the CAT was positively correlated with SOD, but negatively correlated with POD. Thus, the influence of CAT on SEP might be reduced by the counter-effect. Moreover, SEP was positively correlated with proliferation rate, but negatively correlated

Fig. 5 Relationship of proliferation rate, tissue age, genotype (cell line) and somatic embryo production for *Pinus thunbergii*. On the dashed line from Cell line, 36-n represents 1536-1, 1536-2, 1536-3, and 1536-4; 37-n represents 1537-1, 1637-1, 1637-2, 1637-4, 1637-5, 1637-8, and 1637-9. The number on the solid lines indicates the proliferation rate (fold). The number on the dotted lines from Tissue age is the subculturing time (months). The number on the dotted lines from Cell line is the code of the tested cell lines



with the tissue age. We found that starch was negatively correlated with soluble sugar, soluble protein and proliferation rate, but positively correlated with tissue age, while tissue

age was negatively correlated with SEP (Fig. 6). The starch interactions showed that starch accumulation during the proliferation stage was adversely related to the SEP production

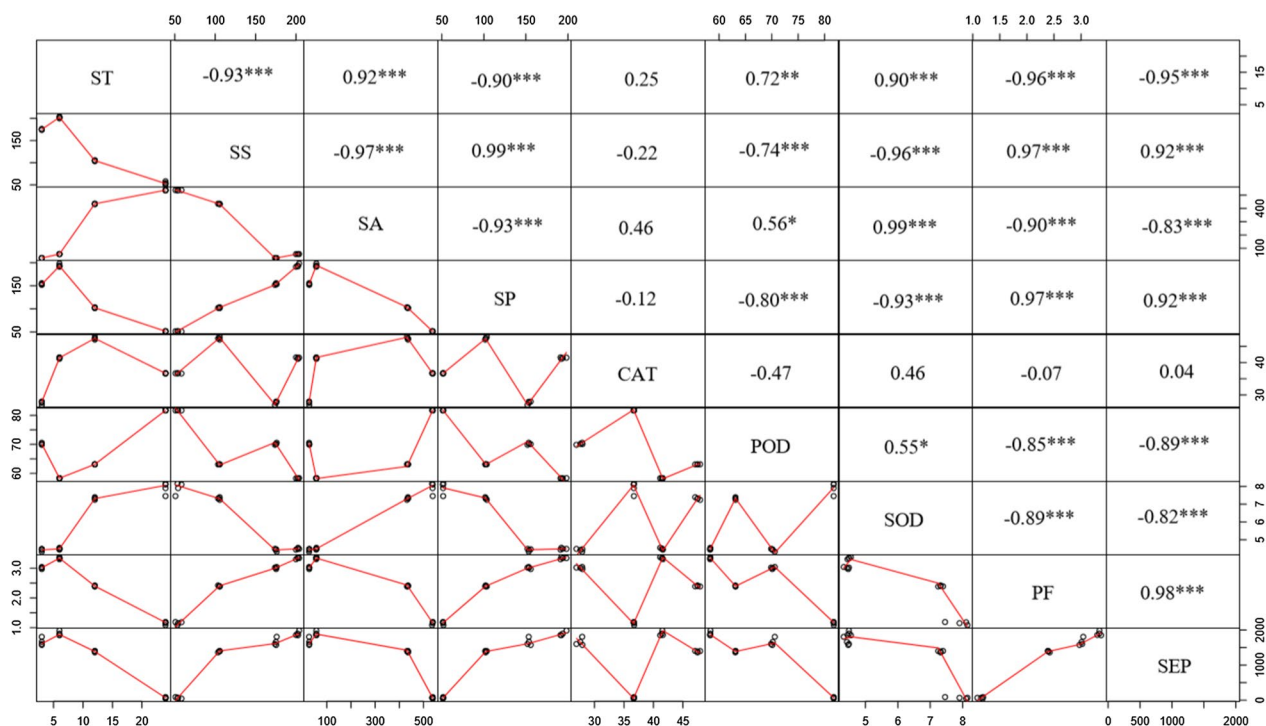


Fig. 6 Correlation analysis of somatic embryo production by the most prolific cell line, 1537-1. ST, tissue age; SS, soluble sugar; SA, starch; SP, soluble protein; CAT, catalase; POD, peroxidase;

SOD, superoxide dismutase; PF, propagation frequency; SEP, somatic embryo production. Significant differences for each variable: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$

of ETs. In addition, the decreases in soluble sugar, soluble protein and proliferation rate reflected the ET maturation capacity to some extent.

Discussion

Effects of cell morphology and physiology on SEP in *P. thunbergii*

Observations of tissue morphology indicated that subculturing for 6 months was a key factor. At this time, although the ET transparency decreased, and filamentous and protuberant surfaces of the clumps were reduced, further microscopic observations revealed that the proportion of the acetocarmine-stained red area changed little, and most of the cell types were still morphologically young (ESM). Meanwhile, SEP in *P. thunbergii* peaked. Similar results were reported in Scots pine (Abrahamsson et al. 2012). However, after 6 months of subculturing, the filamentous and protuberant surfaces of ET disappeared gradually, the proportion of the acetocarmine-stained red area was replaced with abnormal cells, and SEP decreased accordingly. We considered that the disappearance of filamentous and protuberant surfaces of ETs might indicate the beginning of ET aging. The degeneration of the ET morphology led to a decrease in SEP in *Picea abies* (Filonova 2000). We, therefore, hypothesized that the ET maturation capacity of *P. thunbergii* could be estimated by ET surface morphology, histological features, and the proportion of the acetocarmine-stained red areas during the proliferation stage. We found that the CAT activity and intracellular substance (soluble protein, soluble sugar and starch) levels increased when tissues were 6 months old, when SEP peaked. As for the degeneration of ET morphology at 6 months, as mentioned above, the decrease in ET transparency might be related to the increases in intracellular substances. In addition, although the filamentous and protuberant surfaces of the clumps were declining, the ETs still showed ESM structures and the highest proportions of acetocarmine-stained red areas. We speculated that the increase in SEP might be related to the increase in intracellular substance levels and enzymatic activities at this point in time. Grzyb et al. (2017) found that intracellular substances (especially soluble sugar) regulate the transport of phytohormones and the expression of phytohormone response pathways. Pacholczak et al. (2005) also found a correlation between soluble sugars and both endogenous indole-3-acetic acid and abscisic acid levels. In addition, our study found that ETs having lower levels of SOD and POD had good maturation capacities. Diehn et al. (2009) also demonstrated that totipotent cells contain lower levels of reactive oxygen. These factors may serve as suitable markers for assessing the ET maturation capacity during the proliferation stage under

favorable maturation conditions. This approach provides novel insights into early events that influence the ET maturation capacity in somatic embryogenesis of *P. thunbergii* resistant to the pine wood nematode.

Effects of maternal genotype, tissue age and proliferation rate on SEP in *P. thunbergii*

SEP was significantly ($p < 0.001$) affected by maternal genotype, tissue age and proliferation rate. The genotype-dependent SEs maturation and the considerable variations in SEP among cell lines had been reported previously (Montalban et al. 2010, 2012; Kim and Moon 2014; Salaj et al. 2019). Here, we determined that if the clone was productive, then the original maternal genotypes might have a greater potential to produce high-yielding SEs than other maternal genotypes. Thus, the cell lines from seed family 37 had higher SEP levels than those from seed family 36. This result further confirmed the decisive influence of maternal genotypes on SEP. The acquisition of ETs with high-yielding SEs may be correlated with the explant genotype (megagametophyte). Because the seed orchard was open-pollinated, the maternal genes were the same, but the paternal genes were different, which may explain the lack of ETs that produced high-yielding SEs in 2016. Mee et al. (2016) also found that genetic background had a great influence on SEP. The regular SEP level in maternal and clonal genotypes may be an important basis when selecting for the mass propagation of elite clonal genotypes.

In addition, SEP and proliferation rate tended to decrease as tissue age increased were identified. The degeneration of embryogenic potential with increasing tissue age is a common phenomenon in *Pinus* species (Wang et al. 2019; Yang et al. 2020). However, assessing SEP by tissue age alone is not accurate, because the degree of ET degeneration is different among genotypes and culture conditions. For example, cucumber (*Cucumis sativus* L.) maintains its embryogenic potential for more than 3.5 years in cytokinin-independent suspension cultures (Andryskova et al. 2009). Our results suggested the combination of ET surface morphology, histology (the proportion of the acetocarmine-stained red area) and variations in physiological characteristics (carbohydrates and enzymatic activities) could be used to precisely assess the ET maturation capacity. In general, young ETs were better choices for SEP than tissues used for long-term maintenance (von Aderkas et al. 2003; Breton et al. 2006; Lelu-Walter and Paques 2009; Park et al. 2011). Moreover, the interactions among maternal genotypes, tissue ages and proliferation rates of *P. thunbergii* showed that the high-yielding SEs occurred at a proliferation rate of more than threefold, when tissues were 6 months old for cell lines 1537–1 and 1539–1. We determined that specific maternal genotypes, young cultures (less than 6 months old) and high

proliferation rates (threefold) are necessary for high yields of SEs. Additionally, the SEP was genotype-dependent when the latter two conditions were met. Thus, there is increasing evidence that SEP is genotype-dependent.

Conclusions

Our results showed that SEP was significantly affected by maternal genotype ($p < 0.001$), tissue age ($p < 0.001$) and proliferation rate ($p < 0.001$). The maternal genotype determined the level of SEP, and the tissue age similarly influenced the proliferation rate and SEP, in which both were reduced as tissue age increased. In addition, a high proliferation rate ($>$ threefold) might be beneficial to the production of SEs. Moreover, during the proliferation stage, the ET's morphological evolution also showed that decreases in transparency and filamentous surfaces of ETs, as well as the proportion of acetocarmine-stained red areas, decreased along with SEP. The ET morphology and histology may reflect the SEP capability to some extent. Additionally, a correlation analysis showed that SEP was positive correlated with proliferation rate, soluble sugar and protein, but negatively correlated with tissue age, starch, SOD and POD. In general, the ET maturation capacity of pine wood nematode-resistant *P. thunbergii* may be estimated using maternal genotypes, proliferation rate, tissue age, ET surface morphology, histology, as the proportion of acetocarmine-stained red areas, and the variation among physiological characteristics (carbohydrates and enzymatic activities). This study provides new insights for assessing the ET maturation capacities accurately during the proliferation stage of *Pinus* species, especially pine wood nematode-resistant *P. thunbergii*.

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