### PLANT TISSUE CULTURE





# Alkaloid synthesis is coupled to shoot morphogenesis in Argemone mexicana L. (Papaveraceae) in vitro cultures

Miriam Monforte-González<sup>1</sup> • J. Germán Serrano-Gamboa<sup>1</sup> • Cecilia Guízar-González<sup>2</sup> • Ma. Lourdes Miranda-Ham<sup>1</sup> • Felipe A. Vázquez-Flota<sup>1</sup> ®

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### Abstract

During the induction process of an *in vitro* callus culture of *Argemone mexicana* L. (Papaveraceae), the levels of two benzylisoquinoline alkaloids known as berberine and sanguinarine displayed opposing trends. While the berberine levels steadily decreased from the initial explant stage up to the early proliferation of unorganized parenchymatous cell masses, the sanguinarine content increased. Once the callus culture was established, sanguinarine was the primary alkaloid present and berberine could no longer be detected. However, upon shoot regeneration, the berberine accumulation recovered, but sanguinarine was found in the newly formed leafy tissue. After root formation, sanguinarine was relocated to this organ, whereas berberine was evenly distributed between both tissues. Explants from stem internodes did not form callus, and berberine—plus sanguinarine—containing axillary shoots emerged from lateral buds in the induction medium. In contrast to callus-derived shoots, no root formation was observed. Therefore, alkaloid synthesis in A. mexicana in vitro cultures is related to the level of tissue organization in different ways, and while berberine accumulation seems to require the presence of differentiated organs, this is not the case for sanguinarine. Moreover, leafy parts of rootless shoots acquired the capacity to accumulate sanguinarine, which is usually absent in aerial tissues of mature plants. However, when these shoots were rooted, sanguinarine was mainly located in the newly formed roots, while berberine was detected in the shoots at similar levels found in the roots.

Keywords Argemone mexicana · Berberine · Sanguinarine · Shoot regeneration

## Introduction

Accumulation of the benzylisoquinoline alkaloids (BIA) berberine and sanguinarine follows a tissue and developmental pattern in prickly poppy (Argemone mexicana L., Papaveraceae; Xool-Tamayo et al. [2017a](#page-6-0), [b](#page-6-0)). Because both of these alkaloids have diverse industrial and medicinal applications (Rubio-Piña and Vázquez-Flota [2013](#page-5-0)), there is a renewed interest to produce them through cell culture technology. In a recent report, rootless shoot cultures of this plant were obtained. The presence of berberine, which is evenly distributed in aerial and underground tissues of A. mexicana, was observed in these shoot cultures. Interestingly, accumulation of sanguinarine, which is normally absent in the aerial tissues but abundant in roots, was also recorded (Xool-Tamayo et al. [2017a](#page-6-0); Vázquez-Flota et al. [2018](#page-6-0)). Likewise, the onset of sanguinarine accumulation has been described in A. mexicana leaf-derived cell cultures, concomitantly with loss of berberine (Trujillo-Villanueva et al. [2010;](#page-6-0) Xool-Tamayo et al. [2017a\)](#page-6-0). Modifications in the tissue-associated metabolite profiles are not uncommon upon introduction of in vitro cultures, as the morphological alterations that tissues undergo during the process also have an impact on their bio-synthetic capacity (Murthy et al. [2014](#page-5-0)). However, loss of berberine accumulation in callus cultures derived from A. mexicana leaves is noteworthy for two reasons. First, berberine-producing in vitro cultures have been obtained from other species, including Berberis buxifolia Lam. (Alvarez et al. [2009](#page-5-0)), Coptis japonica (Thunb.) Makino (Sato et al. [2001](#page-6-0)), and Thalictrum minus L. (Hara et al. [1994\)](#page-5-0). This



 $\boxtimes$  Felipe A. Vázquez-Flota [felipe@cicy.mx](mailto:felipe@cicy.mx)

<sup>1</sup> Unidad de Bioquímica y Biología Molecular de Plantas, Centro de Investigación Científica de Yucatán, Calle 42 No. 130 Chuburná, 97205 Mérida, Yucatán, Mexico

<sup>2</sup> Centro de Investigación y Asistencia en Tecnología y Diseño del Estado de Jalisco, Unidad Zapopán, Camino el Arenero No. 1227 El Bajío del Arenal, 45019 Zapopan, Jalisco, Mexico

suggests that tissue organization is not required for the synthesis and accumulation of this alkaloid. Secondly, in contrast to A. mexicana, the presence of sanguinarine has neither been reported in the above-mentioned cultures, nor in the plant species from which they were obtained. Accumulation of sanguinarine in rootless shoot cultures is also remarkable because significant amounts of transcripts that correspond to key biosynthetic genes accumulate in aerial tissues in which the alkaloid is absent, such as stems of mature plants (Xool-Tamayo et al. [2017a,](#page-6-0) [b](#page-6-0); Vázquez-Flota et al. [2018\)](#page-6-0). Therefore, although A. mexicana aerial tissues seem to have the capacity to produce and accumulate sanguinarine, in whole plants, it is primarily restricted to their underground parts.

Although berberine and sanguinarine belong to different BIA groups (protoberberine and benzophenanthridine, respectively), they share the early biosynthetic reactions, up to the formation of S-scoulerine, and then diverge into specific branches (see figures; Hagel and Facchini [2012](#page-5-0)). In this context, it is interesting to find possible interactions among the biosynthetic processes of berberine and sanguinarine in A. mexicana. In the present study, the modification in distribution patterns of each alkaloid in association with well-defined morphogenetic events was analyzed. Alkaloid distribution and expression of biosynthetic genes during callus formation from leaf explants to shoot regeneration and rooting were followed. Alkaloid and transcript distributions were also followed along axillary shoot formation from stem internodes, which did not form callus and did not respond to rooting treatments.

## Materials and Methods

Plant materials and induction of in vitro cultures All chemical used for *in vitro* cultures were tissue culture grade from Sigma-Aldrich® (St. Louis, MO), unless specified. Seeds were germinated in vitro in Magenta™ GA-7 vessels boxes (Sigma-Aldrich®) on half-strength Murashige and Skoog (MS, Murashige and Skoog [1962\)](#page-5-0) basal medium, without sucrose or growth regulators. Seeds were disinfected by successive immersions in 80%  $(v/v)$  ethanol, diluted NaClO solution  $[3\% (v/v)$  active sodium hypochlorite], and sterile water for 1, 3, and 10 min, respectively, prior to being imbibed overnight in 5 μM gibberellic acid to promote germination. Once the seedlings were about 8 cm high and displayed between four and six pairs of leaves, both leaves and stem internodes were exscinded as explants. To induce callus or shoot formation, explants were placed on semisolid MS media contained in Magenta<sup>™</sup> vessels supplemented with 20 g  $L^{-1}$ sucrose; 0.045, 0.225, 2.225, 4.45, or 22.25 μM thidiazuron (TDZ); and  $0.2\%$  (w/v) Gelrite<sup>TM</sup>. All media were sterilized by autoclaving 30 min at 121°C (1.4 kg cm<sup>-2</sup>), after adjusting the pH to 5.8 with 0.1 M HCl.



Six explants per vessel were used, and a total of six boxes per treatment were evaluated. Culture vessels were kept under continuous illumination with a photon density of 50 µmol m<sup>-2</sup> s<sup>-1</sup>, provided by a combination of 39-W fluorescent and 60-W incandescent lamps (Phillips, Alto Collection and Studio Collections, respectively; Philips de México, Cd de México, México) at a constant 25°C and 70% relative humidity. Shoots arose either from previously formed callus or directly from explants, depending on the initial tissue used. Newly formed shoots were placed on MS medium supplemented with 5 μM indole-3-butyric acid (IBA) for rooting. In parallel, callus that did not produce leafy structures was exscinded from leaf explants and maintained on MS media with 2 or 8  $\mu$ M 6-benzylaminopurine (BAP), and 1  $\mu$ M 1-naphthaleneacetic acid (NAA). Samples of callus, shoots, and roots were collected for alkaloid and transcript analysis in triplicate. The experiment was performed in two independent repetitions. Each portion was analyzed separately in those cases in which cultured tissues displayed sections with different organization levels in the same unit (callus and shoots portions in the same cluster).

Alkaloid extraction and analysis Solvents were from JT Baker (Phillipsburg, NJ). Alkaloids were extracted from 20 mg of freeze-dried powdered tissues (8 to 10 pieces of the different tissues) in 15 mL of methanol with continuous shaking at 50 rpm for 2 h at room temperature (27°C). After incubation, the debris was separated by centrifugation at  $1400 \times g$ , and 1 mL of the supernatant was taken, centrifuged again at  $1400 \times g$  to eliminate any remaining tissue, and the supernatant was reduced to dryness in vacuo. The residue was dissolved in 100 μL of methanol, and 1 μL was loaded on  $10 \times 20$  cm (length  $\times$  width) thin layer chromatography (TLC) silica gel 60 plate  $F_{254}$  (Merck KGaA; Darmstadt, Germany) for separation. Alkaloids were separated with mixtures of nbutanol:water:NH<sub>4</sub>OH  $(8:1:1 \ v:v:v)$  and benzene:ethanol (9.4:0.6 *v*:*v*), as mobile phases to resolve berberine ( $R_f$  0.32) and sanguinarine  $(R_f 0.36)$ , respectively (Monforte-González et al. [2012](#page-5-0)). Samples were loaded 1 cm above the lower edge of the plate and chromatographed to up to 1 cm below the upper edge. After separation, alkaloids present in the extracts were identified by matching Rf values to commercial standards (Sigma-Aldrich®), and autofluorescence under exposure to  $\lambda_{365}$  radiation provided by a hand-held UV lamp (Spectroline ENF-240C; Westbury, NY). Berberine and sanguinarine displayed bright blue-green and orange-reddish tones, respectively. The identity of the alkaloids and chromatographic coincidences on the plates after separation had been previously confirmed by mass spectroscopy-liquid chromatography ( $m/z$  370.16 for berberine and  $m/z$  332.00 for sanguinarine; Monforte-González et al. [2012;](#page-5-0) Vázquez-Flota et al. [2018\)](#page-6-0). After separation, alkaloids were quantified by in situ fluorescence in a Camag TLC Scanner 4 (Camag,

Muttenz, Switzerland), controlled by the WinCATS 1.4.10 planar chromatography manager (Camag). The excitation wavelength was set at 330 nm, with a 400-nm K400 Camag cut off filter (Monforte-González et al. [2012](#page-5-0)). Three independent analyses were performed for each tissue, and differences were assessed by Tukey's test at  $p < 0.05$ .

Nucleic acid analysis Extraction of total RNA from the collected tissues and additional analysis by Real-Time Quantitative Reverse Transcription Polymerase Chain Reaction (qRT-PCR) was performed as described by (Xool-Tamayo et al. ([2017a,](#page-6-0) [b\)](#page-6-0). Selected transcripts were analyzed to monitor common reactions (norcoclaurine synthase; NCS), and the berberine- (S-scoulerine O-methyltransferase; SOMT) and sanguinarine- (s-cheilanthifoline synthase; CheSyn) committed biosynthetic branches, respectively (Xool-Tamayo et al. [2017b\)](#page-6-0). The corresponding primer sequences were (5′ to 3′; forward and reverse) CATCGCTA ATTACGTTCTCAAGAATCA and ATAGTAGTACATGG AATTACCTGGATGGGA for NCS; ATCCTATCCATGTC TACGAGGGCTATT and CCAGTACCACCACCAACATC TAACA for SOMT; and AGGTCTTCAAGGTGTTGCCC and TCTTTTCCCGCCCGTAACAT for CheSyn. Primer design was based on accessions EU 88189, KT984756, and EF451152 for A. mexicana NCS, SOMT, and CheSyn (CYP719A14), respectively (Xool-Tamayo et al. [2017a,](#page-6-0) [b\)](#page-6-0). For RT-qPCR, first strand cDNAwas synthetized from 1 μg of total RNA that was mixed with 500 ng of oligo (dT) 18 primer, 1 mM of dNTPs, and 200 U of M-MLV Reverse Transcriptase (Invitrogen; Carlsbad, CA), in a total volume of 20 μL, following the manufacturer's instructions. RNA was previously treated with 1 U of DNAse I (Invitrogen) for 30 min at 25°C. The cDNA produced was adjusted to 5 ng and subjected to PCR in a final volume of  $25 \mu L$ , containing 10 μL of SYBR™ Master Mix (Applied Biosystems™, Warrington, Cheshire, UK), with 10 μM of each primer and 2 U of Taq Pol (Invitrogen). Polymerase chain reactions were performed using Eco™ Real-Time PCR System (Illumina, San Diego, CA), following a program of 35 cycles of 60, 30, and 30 s at 95, 62, and 72°C for DNA denaturation, primer alignment, and amplification, respectively. Previously, the reaction mix was heated for 3 min at 95°C for enzyme activation. Gene transcripts were quantified using the  $\Delta\Delta Cq$  method (Simon [2003](#page-6-0)), which compared amounts of the targeted transcript with those of actin, as the reference (Simon [2003](#page-6-0)). Actin primer sequences were (forward and reverse 5′ to 3′) CACIACTACTGCTAAACGGGAAA and ACATCTGC TGGAAGGTGCTG, respectively). The cycle threshold value (Ct) for each PCR was calculated using the EcoStudy software (Illumina). Samples were analyzed in triplicate and the differences were assessed by Tukey's test at  $p < 0.05$ .

### Results

Morphogenetic response of A. mexicana tissues Most treatments resulted in callus formation on the leaf explants. Callus proliferation was observed over the edges of the tissues after 2 wk in culture (Fig. [1](#page-3-0)a) and leafy structures formed on callus, turned into well-formed shoots after two additional weeks (Fig. [1](#page-3-0)b). The organogenetic response differed among TDZ treatments, ranging from 0.3 to 1.8 shoots per explant  $(p < 0.05$ ; one-way ANOVA, Fig. [1](#page-3-0)b). The highest value was registered with 22.25 μM TDZ, and shoots formed on this treatment were used for additional characterization. After remaining one extra week in the induction media (a total of 5 wk), shoots were transplanted to the rooting media, and root development was detected after 1 wk (Fig. [1](#page-3-0)c). These roots continued to extend for the rest of the experimental period for two additional wk.

Internode explants developed axillary shoots from lateral buds in less than 10 d in culture, without any callus formation (Fig. [1](#page-3-0)d). Using the assay conditions, one single shoot formed from each bud, with average responses ranging between 0.5 and 2 shoots per explant in the different treatments (each internode presented two opposite buds). The maximum response was detected with 22.25 μM TDZ. Individual axillary shoots were exscinded from explants and cultured on rooting media, and there was no root formation recorded after 2 wk.

Two sets of shoot cultures, which differed both in origin and in their ability to form roots, were obtained, which were regenerated shoots, which displayed rooting capacity (Fig. [1](#page-3-0)c), and axillary shoots that were non-responsive to rooting treatments (Fig.  $1d$  $1d$ ).

Alkaloid content in in vitro A. mexicana regenerated and axillary shoots Berberine and sanguinarine were quantified on both sets of A. mexicana shoots at different phases of the in vitro culture process (Fig. [2\)](#page-3-0). Berberine was the only alkaloid detected in the initial explants of both leaves and stem internodes. However, disorganization of leaf explants into callus tissue resulted in the onset of sanguinarine accumulation, which was simultaneous with the loss of berberine (Fig.  $2a$  $2a$ ). It should be noted that the leaf portion in explants kept its original alkaloid profile, which consisted in the presence of berberine, but not sanguinarine. Shoot development from callus tissue restored berberine accumulation and maintained levels of sanguinarine. However, upon rooting the regenerated shoots, the sanguinarine accumulation was switched from the leafy parts to the roots (see the Shoot/Roots bar cluster in Fig. [2](#page-3-0)a). In contrast, berberine levels in leafy tissues of the regenerated shoots remained similar to those of the roots, once this organ was formed (Fig.  $2a$  $2a$ ).

The capacity of sanguinarine accumulation, along with berberine, was also observed in the axillary shoots, which developed from the lateral internode buds. Interestingly, the stem

<span id="page-3-0"></span>Figure 1. In vitro cultures of Argemone mexicana tissues. Callus induced from leaves (a). Shoot regeneration from leafcallus (b). Rooting of regenerated shoots (c). Shoot development from axial stem internode  $(d)$ .



portion of these explants that did not suffer any morphological alteration showed a decreased berberine level but did not acquire the capacity to accumulate sanguinarine (Fig.  $2b$ ). This suggests that these effects were not related to the culture time but rather to the morphogenetic events, as was also recorded for the leaf explants.



Figure 2. Alkaloid distribution in Argemone mexicana tissues submitted to in vitro culture. Cultures induced from leaves  $(a)$  and stem internode explants (b). Sets of bars above labels interrupted by a backslash (/) symbol correspond to masses composed of two different tissues (either leaf or stem explants vs callus and shoots newly formed). Each type of tissue was independently analyzed. Bars represent the mean values of three repetitions with standard deviation, whereas asterisks (\*) represent not detected. Bars with the same literal in boldface (berberine) or italics (sanguinarine) presented non-significant differences (Tukey's test at  $p < 0.05$ ).



<span id="page-4-0"></span>Tissue distribution of transcripts involved in alkaloid biosynthesis in the A. mexicana regenerated and axillary shoots The distribution of selected transcripts involved in alkaloid biosynthesis was followed through the different morphogenetic stages for both sets of regenerated A. mexicana shoots to analyze their biosynthetic capacities. Markers for the early common biosynthetic reactions (NCS), and for those specifically committed to either sanguinarine (CheSyn), or berberine (SOMT), were chosen (Fig. 3). In concordance with the alkaloid profiles of the initial leaf (Fig.  $3a$ ) and stem internode (Fig. 3b) explants, which only presented berberine levels, RT-qPCR revealed high transcript levels for NCS and SOMT, which differed from those corresponding to CheSyn. However, upon leaf differentiation into callus, the CheSyn transcripts markedly increased, whereas those for SOMT dropped significantly ( $p < 0.05$  Tukey's test; Fig. 3*a*). This trend in CheSyn and SOMT transcript levels was always observed in calluses that were either still attached to the original leaf explants, isolated as an independent culture, or during the development of new shoots (Fig.  $3a$ ). Transcripts for *SOMT* were recovered in the regenerated rootless shoots, whereas there were similar transcript levels for CheSyn (see the Callus/Shoot bar cluster in Fig. 3a). However, when roots were formed, CheSyn transcripts were mainly detected in the tissue with shoots that displayed levels just above the detection limit, unlike those observed for SOMT transcript distribution (Fig.  $3a$ ). Therefore, the distribution of both *CheSyn* and SOMT transcripts matched those of their corresponding sanguinarine and berberine alkaloids. On the other hand, although some variations were observed, NCS transcripts remained in fair amounts, regardless of the tissue organization observed (Fig.  $3a$ ).

Similar to the regenerated shoots, rootless axillary shoots showed a high CheSyn expression level, while the SOMT level was constant (Fig. 3b), coincidental to the alkaloid distribution (Fig. [2](#page-3-0)b). In addition, stem sections that did not present morphological alterations retained both transcript and alkaloid profiles as the original explants (Fig. 3b).

## **Discussion**

This research was conducted to establish a better understanding of morphogenesis-related alkaloid biosynthesis in A. mexicana tissues. This was accomplished by developing an organogenesis regenerating protocol, which started with leaf explants on medium containing TDZ. Shoots that emerged from leaf-derived callus were then rooted on IBA (Fig.  $1a 1a$ c). Although axillary shoots were also obtained from stem internode explants, these could not be rooted (Fig.  $1d$  $1d$ ). The lack of rooting has also been noticed in multiple adventitious shoots derived from A. mexicana hypocotyls, which did not develop roots when exposed to different treatments (Xool-Tamayo et al. [2017a\)](#page-6-0). Moreover, isolated roots displayed restricted growth in in vitro cultures (Xool-Tamayo et al. [2017a\)](#page-6-0). This suggests a limited in vitro rooting capacity for this plant. In fact, although shoot regeneration from Argemone callus was lower than in other Papaveraceae species, such as Papaver somniferum L. (Park and Facchini [2000](#page-5-0)), and Papaver bracteatum Lindl. (Rostampour et al. [2010](#page-5-0)), all of the shoots were successfully rooted.

In vitro morphogenetic events affected the accumulation of sanguinarine and berberine differently. Leaf explants presented abundant berberine amounts but lack of sanguinarine accumulation. During callus formation, these explants lost the former, while acquiring the latter (Fig.  $2a$  $2a$ ). This effect was directly related to tissue disorganization because explant portions not forming callus retained their original alkaloid profile (Fig. [2](#page-3-0)a). This was also observed in stem explants (Fig. [2](#page-3-0)b).

Interestingly, those callus tissues that were collected from the explants and kept on MS media with 2  $\mu$ M or 8  $\mu$ M BAP and 1 μM NAAwere able to proliferate and maintain the same





newly formed). Each type of tissue was independently analyzed. Bars represent the mean values of triplicates with standard deviation. Diagram at the left depicts the condensed biosynthetic sanguinarine and berberine pathways.



<span id="page-5-0"></span>alkaloid pattern as those initially collected from the leaf explants (loss of berberine, but presence of sanguinarine) (Guízar-González et al. 2012). Moreover, different induction treatments, such as the exposure to methyl jasmonate, salicylic acid, or yeast extract, did not modify this pattern, although sanguinarine accumulation increased (Trujillo-Villanueva et al. [2010;](#page-6-0) Guízar-González et al. 2012, 2016).

Although alkaloid biosynthetic capacity is frequently lost during tissue disorganization (Murthy *et al.* 2014), berberineproducing in vitro cultures have been reported for other species of the Papaveraceae family (Day et al. 1986; Facchini and Bird 1998; Zakaria et al. [2011\)](#page-6-0). This suggests that berberine biosynthesis does not require tissue integrity (Hara et al. 1994; Sato et al. [2001](#page-6-0); Alvarez et al. 2009). In contrast, the development of a novel sanguinarine accumulating capacity has been observed in hypocotyl-derived P. somniferum cell cultures, after fungal elicitation (Facchini et al. 1996). Interestingly, sanguinarine was also found in welldifferentiated leafy parts of callus-regenerated shoots, but it only occurred in the absence of roots (Fig.  $2a$  $2a$ ). Upon root formation, it was switched to this new location (Fig.  $2a$  $2a$ ). In contrast, berberine accumulation in these leafy tissues remained unaffected by rhizogenesis. It should be noted that the presence of both sanguinarine and berberine always coincided with CheSyn and SOMT expression, which are required in the corresponding biosynthetic pathways (Fig.  $3a$  $3a$ ), which points to a direct involvement of the tissue in this process.

The acquisition of sanguinarine accumulating capacity, while keeping that of berberine, which was observed in the newly formed leafy tissues of the regenerated shoots, was also noted in axillary shoots, formed from internode stem explants (Fig. [2](#page-3-0)b). Moreover, this trend has also been recorded in hypocotyl-derived A. mexicana rootless shoots (Xool-Tamayo et al. [2017a](#page-6-0)). Unfortunately, rooting has only been achieved in callus-regenerated shoots (Fig. [1.](#page-3-0) X). Rooting has not been reached in neither the axillary shoots obtained in this work nor in those hypocotyl-derived (Xool-Tamayo et al. [2017a\)](#page-6-0) in order to confirm the sanguinarine distribution pattern observed in regenerated shoots. However, during seedling development, sanguinarine was detected in newly unfolded leaflets up to the formation of secondary roots, when it markedly decreased in these tissues to be relocated to the roots. In contrast, berberine presence in aerial parts was unaffected by root development (Xool-Tamayo et al. [2017b](#page-6-0)), as it was in the regenerated rooted shoots in the present study (Figs. [2](#page-3-0)a and  $3a$  $3a$ ). Taken together, these data suggest that in A. mexicana, roots might play a role to define the final alkaloid distribution, rather than to confer biosynthetic capacities. Because synthesis of sanguinarine and berberine utilizes s-scoulerine as a common late intermediary (see Fig. [3](#page-4-0)), underlying molecular mechanisms seem to be related to the preferential expression of the committed biosynthetic pathway for each alkaloid (Fig.  $3a$  $3a$  and b).

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