



Cardenolide and glucosinolate accumulation in shoot cultures of *Erysimum crepidifolium* Rchb.

Elisa Horn¹ · Yvonne Kemmler¹ · Wolfgang Kreis¹ · Jennifer Munkert¹

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Abstract

Erysimum crepidifolium Rchb. is one of the few Brassicaceae species accumulating glucosinolates as well as cardenolides. This is possibly providing a selective advantage in evolution as both compounds are part of a chemical defense system. In order to study the biosynthesis of these compounds, a regeneration protocol for *E. crepidifolium* using *in vitro* shoot cultures derived from seeds has been developed. Murashige and Skoog (MS) culture medium supplemented with various combinations of cytokinins and auxins was used. MS medium containing NAA (naphthaleneacetic acid, 0.04 mg mL⁻¹) and BAP (6-benzylaminopurine, 0.2 · 10⁻² mg mL⁻¹) proved to be optimal for root formation. Plantlets developed well on modified MS medium without the use of phytohormones. About 80% of the plantlets rooted *in vitro* developed into intact plants after transfer to the greenhouse. Cardenolides (1.75 mg g⁻¹ dry weight (DW)) were detected in cultured shoots on solid DDV media while glucosinolates mainly accumulated in roots where 0.025 mg g⁻¹ FW were detected in shoots cultured on the same medium (DDV). The expression of two *progesterone 5β-reductase* and three *Δ⁵-3β-hydroxysteroid dehydrogenase* genes were measured in shoot cultures since the encoded enzymes are supposed to be involved in cardenolide biosynthesis. *E. crepidifolium* shoot cultures propagated on solid media meet the necessary requirements, *i.e.*, clonal homogeneity, product accumulation, and gene expression, for a suitable model to study cardenolide but not glucosinolate biosynthesis.

Keywords *Erysimum crepidifolium* Rchb. · Regeneration · Glucosinolates · Cardenolides

Introduction

Arabidopsis thaliana (L.) Heynh., a member of the Brassicaceae family, is widely used as a model organism in genetic studies. The genus *Erysimum* L. belongs to the same family and comprises more than 200 species (Bailey *et al.* 2006; Polatschek 2013a, b). The production of cardenolides by species in the genus *Erysimum* is one of the best-studied examples of an evolutionarily recent gain of a novel chemical defense (*e.g.*, Jaretsky and Wilcke 1932; Makarevich *et al.* 1994; Züst *et al.* 2020). Most Brassicaceae species accumulate glucosinolates which are activated by myrosinases to produce repellent mustard oils (Fig. 1; Fahey *et al.* 2001). However, several herbivores have evolved to overcome this defense

mechanism (Winde and Wittstock 2011). Cardenolide biosynthesis may have evolved further since the ‘mustard oil bomb’ (Matile 1980) to provide a new type of repellent. In fact, several glucosinolate-adapted beetles refused to feed on dietary cardenolides at levels found in *Erysimum* species (Nielsen 1978a, b).

Cardenolides and cardenolide-containing plant extracts including *Erysimum* species (Zhu *et al.* 1989; Burger and Wachter 1998) have been used for centuries in the therapy of heart insufficiency in humans. The biosynthesis of cardenolides has mainly been studied in *Digitalis* (Kreis and Müller-Uri 2010, 2013; Kreis 2017). The likely existence of a very similar cardenolide pathway in *Erysimum* has not been demonstrated. Munkert *et al.* (2011, 2014) proposed *Erysimum* species as model plants for the study of cardenolide biosynthesis in Brassicaceae when investigating the enzymes involved and their respective genes. This idea was further elaborated by Züst *et al.* (2018).

In previous efforts, greenhouse plants originating from seeds were used (Munkert *et al.* 2014); however, the generation of reliable data is often hindered by the limited

✉ Jennifer Munkert
jennifer.munkert@fau.de

¹ Pharmaceutical Biology, Department of Biology,
Friedrich-Alexander-University, Staudtstr. 5,
D-91058 Erlangen, Germany

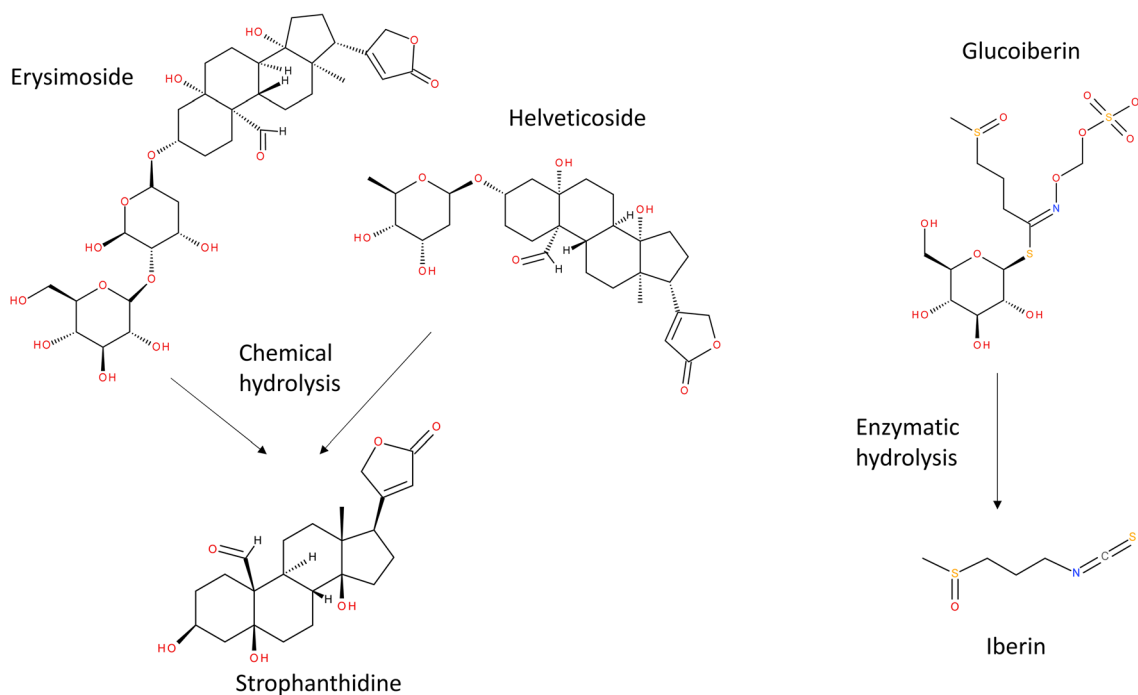


Figure 1. Structures of the major cardenolides (erysimoside, helveticoside) and the major glucosinolate (glucoiberin) present in *Erysimum crepidifolium* Rchb. shoot cultures and their respective aglycones released after hydrolysis.

availability of homogeneous plant material (Schöner *et al.* 1986; Kreis *et al.* 2015). Therefore, the main purpose of this work was to develop a rapid micropropagation protocol for genetically homogenous *E. crepidifolium* shoot cultures and plants. Plant growth, cardenolide content, and glucosinolate content were quantified during plant regeneration and until permanent shoot cultures were established (Fig. 1, 2).

Materials and methods

Chemicals Myrosinase was prepared from *Sinapis alba* (mustard) seeds as described by Meitingner and Kreis (2018). All cardenolides analyzed (erysimoside, helveticoside, k-strophantoside, β -methyl digitoxin; Fig. 1) were sourced from our reference collection or from Merck KGaA (Darmstadt, Germany). All other chemicals including MS medium, growth regulators, or solvents were purchased from Carl Roth GmbH & Co. KG (Karlsruhe, Germany) or from Merck KGaA (Darmstadt, Germany).

Plant regeneration Seeds from wild plants of *E. crepidifolium* were surface-sterilized in 4% calcium hypochlorite, washed in autoclaved distilled water, and germinated aseptically on solid Murashige and Skoog (MS) medium (Murashige and Skoog 1962) containing 30 g glucose and 2.25 mg 6-benzylaminopurine (BAP) per liter (solid DDV medium). After 6 to 8 wk, developing shoots were transferred to liquid MS medium containing 30 g glucose and 2.25 mg BAP per

liter (liquid DDV medium). After 6 wk of adaption to the liquid media, the shoots were transferred to a liquid MS medium containing 30 g glucose and varying concentrations of naphthaleneacetic acid (NAA; 0 to 80 mg) and BAP (0 to 2.25 mg) per liter to initiate rooting. The optimal composition was established at 40 mg NAA and 0.02 mg BAP per liter (liquid EDDV medium) (Fig. 2c). After 40 d in rooting medium, plantlets with well-developed roots were transferred to solid OH medium that is based on DDV medium with increased amounts of KH_2PO_4 (340 mg L^{-1}), glycine (4 mg L^{-1}), and glucose (33 g L^{-1}) but without additional phytohormones. After 2 mo, the plantlets were transferred to autoclaved soil and allowed to acclimatize. They were kept in seeding boxes for about 3 wk and were sprayed with water twice a day during the first week to guarantee a humid atmosphere. In order to reduce the humidity during the acclimatization period, spraying with water was reduced gradually (Arikat *et al.* 2004). Following this procedure the plants were cultivated at standard greenhouse conditions (Fig. 2d).

Permanent shoot cultures Shoots developed different phenotypes depending on whether they were kept in liquid DDV or on solid DDV medium (Fig. 2a, b). This resulted in two types of permanent shoot cultures, and the cardenolide and glucosinolate contents (see below) have been determined for both. All stock cultures were kept under permanent white light (fluorescent lightning, approximately $40 \mu\text{mol photons m}^{-2} \text{ s}^{-1}$) and at 21 to 24°C. Permanent cultures cultivated in liquid media were kept on MULTITRON gyratory shakers (Infors

Figure 2. *Erysimum crepidifolium* Rchb. germinated *in vitro*. (a) Permanent cultivation in liquid DDV medium. (b) Rooting *in vitro* in liquid medium. (c) Rooted cultivation on solid DDV medium. (d) Regenerated plants grown in the greenhouse.



GmbH, Sulzemoos, Germany) at 55 rpm. Stress treatment with either 2-cyclohexene-1-one or methylvinyl ketone (MVK) was applied as described by Munkert *et al.* 2014.

RNA extraction and cDNA synthesis RNA was isolated from the individual tissue samples. These samples were frozen with liquid nitrogen and then ground to a fine powder using mortar and pestle. Total RNA was extracted using the illustra™ RNAspinMini RNA Isolation Kit (GE Healthcare, Munich, Germany). cDNA was synthesized employing the SuperScript™ III First-Strand Synthesis for reverse transcription PCR (RT-PCR) Kit (Invitrogen, Karlsruhe, Germany).

Quantitative real-time analysis (qPCR) Real-time PCR was carried out in the StepOnePlus Real-Time PCR System (Life Technologies, Erlangen, Germany) using the SYBR Green Mastermix Kit (Applied Biosystems, Erlangen, Germany) according to the manual provided. The relative gene expression levels were calculated using the $2^{-\Delta\Delta C_t}$ -method (Livak and Schmittgen 2001) with actin as the reference gene. Primers for actin are based on *A. thaliana* actin (At3g18780). Primer pairs used for qPCR are listed in Table 1 and are based on following mRNA sequences: *EcP5βR1*, GU354236 (Munkert *et al.* 2011); *EcP5βR2*, KF234078 (Munkert *et al.* 2015); *Ec-*

3βHSD1, HM366939; *Ec-3βHSD2*, JF274706; *Ec-3βHSD3*, MT978165 (Munkert *et al.* 2014).

Cardenolide extraction and determination Cardenolides were extracted from plant and shoot samples as described by Wiegbe and Wichtl (1993). The extracts were analyzed by HPLC according to Wichtl *et al.* (1982) with slight modifications (Schebitz *et al.* 2010). For HPLC analysis, a Waters 1525 Binary HPLC Pump system with a Waters 2487 Dual

Table 1. Primer pairs used for qPCR

Name	Sequence in 5'-3' direction
qEcP5βR1 for:	GTCCACACGCGCCTAATCT
qEcP5βR1 rev:	AGTAAAAGGCGGATCGTGCC
qEcP5βR2 for:	CAATGGCTCGTGGAAGGT
qEcP5βR2 rev:	ACCGACCAACAACGGAAC
qEc3βHSD1 for:	ATG TCT GGA AAA AGA TTG GAT GG
qEc3βHSD1 rev:	ATC CAC CAG AAG CTG TTT TGA T
qEc3βHSD2 for:	GGG CTA GAG AAA GCA ACG TTT T
qEc3βHSD2 rev:	GCT CGT GGT ACA TAC GAT TGA C
qEc3βHSD3 for:	CGA TGT TAC GAA CGA GAC AGA G
qEc3βHSD3 rev:	CTC AGA CGT AAC GCT GGT TGT A

k Absorbance Detector was employed (Waters GmbH, Eschborn, Germany). The separation was carried out on a ReproSil® column (C18, 5 μm , 4.6×250 mm), eluting with a step gradient composed of double-distilled water (solvent A) and acetonitrile (solvent B): start (20%B), 20 min (32%B), 7 min (42%B), 13 min (60%B), 19 min (100%B), and 6 min (20%B). Erysimoside and helveticoside were detected and quantified at 220 nm using a UV detector. Erysimoside and helveticoside standards were purchased from Merck KGaA (Darmstadt, Germany) and β -methyl digitoxin served as the internal standard for quantification. In several experiments, cardenolides were hydrolyzed prior to quantification (“total cardenolides”) following the protocol of Mannich and Siewert (1942) with slight modifications only.

Glucosinolates Glucosinolates were extracted as described by Meitinger and Kreis (2018). For extraction, 1 mg ascorbic acid as well as 250 μL of a sinigrin solution was added to 0.35 g of fresh plant material until a final concentration of 1.8 mM was reached. In order to ensure complete hydrolysis of endogenous glucosinolates, myrosinase prepared from mustard seeds (Meitinger and Kreis 2018) was added.

After 2 h of incubation at 23°C and 10 min of centrifugation at 10,000g, 3 μg μL^{-1} benzylisothiocyanate was added as an internal standard and the supernatant extracted with dichloromethane. After separation, the organic phase was analyzed by GC-MS using the following conditions: DB-5 ms (Agilent Technologies Deutschland GmbH, Waldbronn, Germany) column (30 m \times 0.25 mm, film 0.25 μm); column oven temperature: 60°C; injector temperature: 230°C; injection volume: 1 μL ; split ratio: 1:100; ion source temperature: 200°C; interface temperature: 210°C; and MS in scan mode: 35–200 m/z. The temperature gradient with helium as the carrier gas was implemented at a flow rate of 46.3 cm s^{-1} . The temperature was kept constant for 3 min at 60°C before it was increased to 210°C at a rate of 10°C per min.

Results and discussion

Micropropagation protocol of *Erysimum crepidifolium* Rchb. Surface-sterilized single seeds were placed on a modified MS solid medium containing 2 mg L^{-1} BAP (DDV). In previous studies, BAP proved to be the most suitable phytohormone to initiate shoot multiplication without root formation (Zimmerman and Scorza 1992).

Alternatively, rooting was induced on solid media supplemented with NAA. Murashige and Skoog (1962) previously reported that NAA is a more potent auxin than natural occurring indoleacetic acid (IAA). We here tested 20 different combinations of BAP and NAA in order to initiate root formation. Optimal results were achieved when liquid MS medium containing 2 mg L^{-1} BAP and 40 mg L^{-1} NAA was used and root

formation began after 2 wk. George and Sherrington (1984) also reported that rooting can be induced with a nutrient medium containing high auxin and low cytokinin concentration. Even higher concentrations of BAP and NAA stimulated callus formation as had been reported before for other plants (Hussey and Stacey 1981; Iriando 1990a, 1990b).

Plantlets were transferred onto solid, hormone-free MS medium after 6 wk on rooting medium. After an additional 8 wk, the plantlets were ready to be potted (Fig. 2a–d). Plantlets were transferred to the greenhouse and kept initially on a sterilized mixture of sand and soil to avoid microbial damage. Shoot cultures grown *in vitro* fail to form sufficient layers of epicuticular wax (Grout and Aston 1977; Sutter and Langhans 1979). In order to still guarantee a humid atmosphere, the plantlets were kept in seeding boxes for 3 more wk. Two months after their transfer to the greenhouse, 77.5% of the plants had survived and became well developed (Fig. 2d). This protocol provides genetically homogenous plants that can be used in further studies; however, vegetative propagation by scions and runners is also feasible (unpublished observations).

Glucosinolates and cardenolides in permanent cultures and during regeneration Developing shoots from stock cultures on solid medium were transferred into liquid MS medium for further proliferation. Cultivation in liquid medium facilitates the delivery of biosynthetic precursors or chemical stressors and makes it easier to monitor an increase in biomass (Fig. 2b). Permanent shoot cultures are suitable for studying the regulation of cardenolide biosynthesis or for detailed analyses of the influence of light and chloroplast differentiation which has been demonstrated in somatic embryos of *D. lanata* (Scheibner *et al.* 1989). After transfer into liquid medium, *E. crepidifolium* shoots developed an abnormal, clinched phenotype (Fig. 2a). The chlorophyll content was 0.3 mg to 0.9 mg g^{-1} fresh weight (FW) and the biomass only doubled within 3 wk. A 30 mg mL^{-1} glucose medium was provided after 30 d of culture, and the partially submerged shoot cultures accumulated neither glucosinolates nor cardenolides (see below). On the solid DDV medium, shoot tufts developed successfully (Fig. 2c) and could be propagated easily. Shoot cultures kept at these conditions contained cardenolides but no glucosinolates (see below).

Glucosinolates The glucosinolate content was calculated from iberin released from glucoiberin by the action of myrosinase (Fig. 1). Myrosinase from mustard seeds was added to ensure the complete conversion of glucosinolates to their respective isothiocyanates since the endogenous myrosinase activity was very low from shoots cultured in liquid medium (data not shown). Hydrolysis of glucosinolates usually occurs when plant tissue is damaged and glucosinolates come in contact with endogenous myrosinases (Rask *et al.* 2000; Vaughn

and Berhow 2005; Loebers *et al.* 2014). After prolonged cultivation on a solid hormone-free medium, *E. crepidifolium* shoot cultures contained 0.025 ± 0.01 mg g⁻¹ FW ($n = 3$) glucosinolates. Glucosinolates did not accumulate in shoots cultured in a liquid medium (Table 2).

Only low levels of glucosinolates were detected in leaves of greenhouse plants (Table 2). The glucosinolate content of roots was considerably higher and reached 4.55 mg iberin g⁻¹ FW. Concentrations as high as 5% of the dry seed mass were reported for individual glucosinolates in flowers and seeds (Bennett *et al.* 2004).

Cardenolides *E. crepidifolium* accumulates approximately 1% cardenolides in the dry mass of its above ground organs, most of them bearing k-strophanthidin (str) as their aglycone. Digitoxose (dox) and glucose (glc) are the sugar components of the main glycosides erysimoside (str-dox-glc) and helveticoside (str-dox), representing about 90% of the cardenolide mixture (Fig. 1; Gmelin and Bredenberg 1966). Since only traces of the minor cardenolides such as erycordin and glucostrophalloside (Makarevich *et al.* 1974) were detected, we only quantified erysimoside and helveticoside either individually or as their degradation product, k-strophanthidin, released by chemical hydrolysis of the cardenolide extract (Fig. 1). Approximately 80 µg g⁻¹ FW cardenolides were detected in shoots cultivated on the DDV medium even after prolonged subculture for more than 2 y. The concentrations decreased dramatically when stock shoot cultures were transferred from the solid into the liquid DDV medium where they developed an abnormal phenotype (Fig. 2a). While minor amounts of cardenolides were still present after 3 wk in liquid medium (about 10 µg g⁻¹ FW), they had vanished completely after two mo. Cardenolide formation also ceased when shoots were kept on the solid DDV medium but cultivated in permanent darkness. After 1 wk, the cardenolide content dropped down to 10% of that seen in control shoots grown at standard

conditions. This was paralleled by the degradation of chlorophyll which dropped from 0.3 to 0.03 mg g⁻¹ FW within 12 d.

This corroborates the findings of others (Luckner and Diettrich 1985; Eisenbeiß *et al.* 1999) who demonstrated that cardenolide formation depends on morphological differentiation and light. This is also consistent with observations concerning other natural compounds. For example, Coste *et al.* (2011) reported that hypericin biosynthesis is impaired after transfer from a solid into a liquid culture medium. Hugentobler and Renwick (1995) discussed possible reasons for differential formation of cardenolides and glucosinolates in *Erysimum* as a result of nutrient deficiency. Shoots transferred into rooting medium also ceased to produce cardenolides which can be explained by the altered medium composition (Table 2; 0.04 mg mL⁻¹ NAA) as was suggested by Patil *et al.* (2013). Plantlets grown on the hormone-free medium prior to their transfer into soil produced only minor quantities of cardenolides. However, cardenolides accumulated in a similar range in young leaves (0.38 mg g⁻¹ dry weight (DW)) of *Erysimum crepidifolium* plants as in liquid shoot cultures (0.47 mg g⁻¹ DW), but did not reach cardenolide levels of shoot cultures that were cultured on solid media (Table 2).

Expression of two *EcP5βR* genes and three *Ec3β-HSD* genes in *E. crepidifolium* shoot cultures Other than the cardenolide content, gene expression levels of *EcP5βRs* and *Ec3β-HSDs* were measured since the encoded enzymes, namely, progesterone 5β-reductase and Δ⁵-3β hydroxysteroid dehydrogenases, are reported to be involved in cardenolide biosynthesis in *Digitalis* (Kreis and Müller-Uri 2010, 2013). Two *EcP5βR* (*EcP5βR1*; *EcP5βR2*) genes have been reported to be expressed in *E. crepidifolium* greenhouse plants by Munkert *et al.* (2011, 2015) who also characterized recombinant forms of the encoded progesterone 5β-reductases. In permanent shoot cultures (solid medium), the expression of *EcP5βR1* was about six times higher than that of *EcP5βR2* (Fig. 3a). The expression of *EcP5βR2*, but not *EcP5βR1*, was stimulated by abiotic stress. When treated with 2-cyclohexen-1-one or methyl vinyl ketone (MVK), expression increased 6.8-fold and 3.4-fold, respectively, after 24 h, after 7 d, basic expression levels, similar to levels prior to abiotic stress treatment, were reached (Fig. 3b). Heterologous *P5βR1* of *D. purpurea* was also expressed constitutively in plants, whereas the expression *P5βR2* dramatically increased in stress conditions (Pérez-Bermúdez *et al.* 2010).

Munkert *et al.* (2014) identified three genes (*Ec3β-HSD1*; *Ec3β-HSD2*; *Ec3β-HSD3*) in *E. crepidifolium*. *Ec3β-HSD1* did not respond to various forms of stress, whereas *Ec3β-HSD2* and *Ec3β-HSD3* were upregulated when plants were treated with methyl jasmonate or methyl vinyl ketone. We observed that during rooting, the expression levels of *Ec3β-*

Table 2. Accumulation of glucosinolates (iberin released from glucoiberin) and cardenolide (k-strophanthidin released from erysimoside and helveticoside) in greenhouse *Erysimum crepidifolium* Rchb. plants and shoot cultures (mean ± STD, $n = 3$)

Plant part	Iberin (mg g ⁻¹ FW)	Cardenolides (mg g ⁻¹ DW)
Shoot culture (liquid)	n.d.	0.47 ± 0.27
Shoot culture (solid)	0.025 ± 0.01	1.75 ± 0.85
Young leaves	0.41 ± 0.02	0.38 ± 0.19
Mature leaves	0.11 ± 0.01	Traces
Main root	4.55 ± 1.37	Traces
Adventitious roots	0.52 ± 0.02	n.d.

n.d., not detectable

FW, fresh weight; DW, dry weight

Traces: less than 0.01 mg g⁻¹ DW

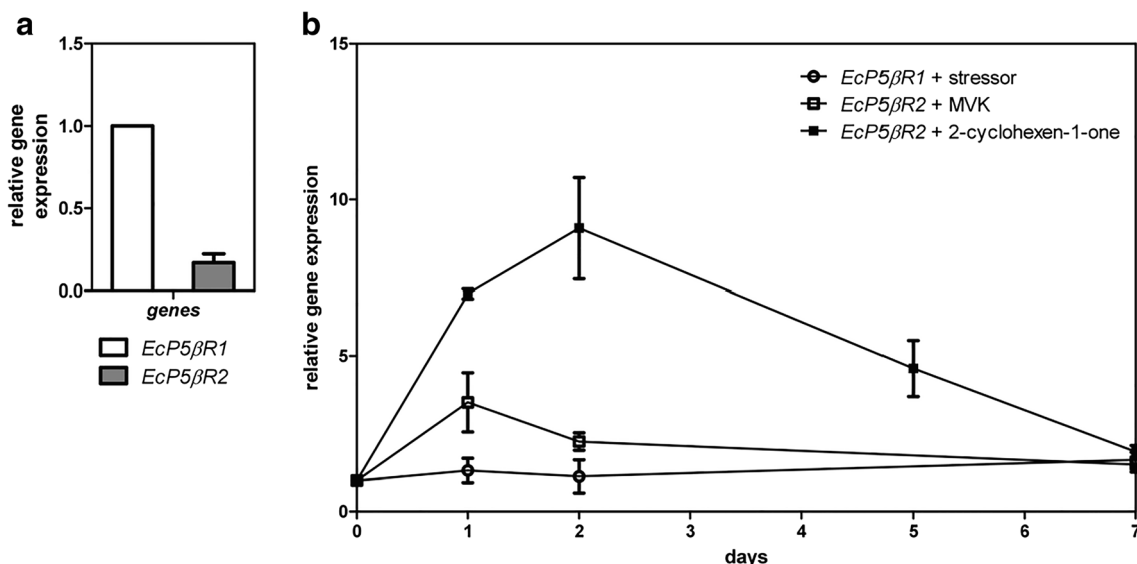


Figure 3. Relative gene expression of *Erysimum crepidifolium* Rchb. progesterone 5 β -reductase 1 and 2 (*EcP5 β R1*; *EcP5 β R2*). (a) Relative expression level of *EcP5 β R2* to *EcP5 β R1* normalized on actin. (b)

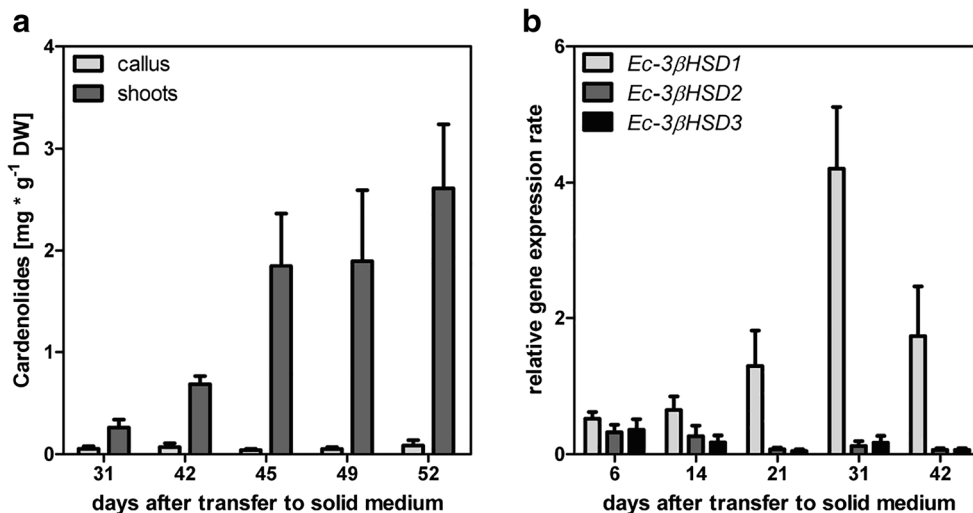
Relative gene expression of *EcP5 β R1* and *EcP5 β R2* normalized on actin, when shoots were treated with 2-cyclohexen-1-one or methyl vinyl ketone (MVK).

HSD1 remained almost constant, whereas the expression level of *Ec3 β -HSD2* and *Ec3 β -HSD3* increased by almost eightfold immediately after transfer to the rooting medium which was interpreted as a stress response (data not shown). Exposing 4-month-old plants to various stress conditions, Munkert *et al.* (2014) found that cardenolide content paralleled gene expression of *Ec3 β -HSD2* and *Ec3 β -HSD3* in several but not all tested stress conditions. In our studies, shoots grown in liquid medium developed an abnormal phenotype and did not accumulate cardenolides although *Ec3 β -HSD2* and *Ec3 β -HSD3* were upregulated. This indicates that morphological differentiation plays a pivotal role in triggering cardenolide formation and that *Ec3 β -HSDs* may also be involved in other physiological processes.

After rooting and transfer to solid medium, the relative expression of *Ec3 β -HSD1* steadily increased from 0.51 to 4.19 and thus paralleled cardenolide accumulation, whereas the expression levels of *Ec3 β -HSD2* and *Ec3 β -HSD3* remained nearly unaffected. *Ec3 β -HSD1* expression seems to be related to plant development and differentiation, including cardenolide formation, whereas *Ec3 β -HSD2* and *Ec3 β -HSD3* are stress-sensitive (Fig. 4a, b).

All three 3 β -HSD isoforms investigated here were expressed in each phase of the regeneration protocol and in permanent shoot cultures in stages with or without cardenolide accumulation. Similar results were obtained for 3 β -HSD and *P5 β R1* of *Digitalis lanata* (Ernst *et al.* 2010); the latter is considered to be a key enzyme in cardenolide biosynthesis (Herl *et al.* 2008). Like the 3 β -HSD1 of *E. crepidifolium*,

Figure 4. (a) Cardenolide content (k-strophanthidin) after transfer of tuft-like structures to solid DDV medium. In developing *Erysimum crepidifolium* Rchb. shoots (dark gray) cardenolide formation increased. Tuft-like structures (callus) still present after 31 to 52 d continued to contain small amounts of cardenolides (light gray). Mean (\pm STD; $n = 3$). (b) Relative gene expression level of *Ec-3 β HSD1*, *Ec-3 β HSD2* to *Ec-3 β HSD3* normalized on actin.



the *P5βR1* of *D. purpurea* was expressed constitutively in plants. The expression *P5βR2* was mainly induced by stress (Pérez-Bermúdez *et al.* 2010).

Like the glucosinolates, cardenolides were only detected after shoot differentiation and during plant regeneration. Cardenolide formation returned after approximately 2 mo on solid medium (Fig. 4a) but did not correlate with the expression of the *3β-HSD* genes. Cardenolide content and pattern can change during plant growth and development as has been demonstrated widely for *Digitalis* species (Weiler and Zenk 1976; Wichtl and Freier 1978; Braga *et al.* 1997).

Conclusion

We conclude that *E. crepidifolium* shoot cultures propagated on solid and liquid media meet the essential requirements (clonal homogeneity, cardenolide production) of a suitable model to study cardenolide but not glucosinolate biosynthesis. The development of this micropropagation protocol additionally offers access to genetically identical mature plants for future studies.

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Authors' contributions All authors contributed to the study conception and design. WK and JM designed the study. Material preparation, data collection, and analysis were performed by EH and YK. The first draft of the manuscript was written by WK, EH, and JM. Further, all authors commented on previous versions of the manuscript. All authors read and approved the final manuscript.

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