



# Synergistic effect of coronatine and sorbitol on artemisinin production in cell suspension culture of *Artemisia annua* L. cv. Anamed

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## Abstract

Artemisinin is an efficient anti-malarial drug and it possesses biological activity against a wide range of cancers. The combined application of two different elicitors can be an efficient way to increase the production of secondary metabolite in plant cell cultures. The results of coronatine (Cor) pretreatment and three concentrations of sorbitol were assessed on the growth, biochemical traits, expression of artemisinin biosynthetic genes, and artemisinin production in *Artemisia annua* cell suspension culture (CSC). After pretreating CSC with 0.05  $\mu\text{M}$  Cor [on the 14th day (three days before the stationary phase) for 48 h], liquid medium in the culture flasks was decanted and replaced with fresh medium (containing 30 g/L sucrose) plus or minus sorbitol at selected concentrations (0, 20, 30, and 40 g/L) on day 16th (one day before the stationary phase). The sorbitol treatment enhanced the contents of malondialdehyde (MDA) and hydrogen peroxide ( $\text{H}_2\text{O}_2$ ) and resulted in oxidative stress. Cor-pretreatment increased the activity of antioxidant enzymes and consequently it reduced  $\text{H}_2\text{O}_2$  content and oxidative stress which resulted in decreased MDA content and better growth. The application of Cor plus sorbitol resulted in a dramatic enhancement in the expression of artemisinin biosynthetic genes and artemisinin production at all concentrations. The expression levels of artemisinin biosynthetic genes (about 7.66, 8.67, 8.67, and 8.33-fold in *ADS*, *CYP71AV1*, *ALDH1*, and *DBR2* genes, respectively at 4 h after sorbitol treatment) and artemisinin production (9.33 mg/L, 8-fold) peaked at 30 g/L sorbitol plus Cor and decreased at 40 g/L sorbitol, probably because of higher oxidative stress.

## Keymessage

The simultaneous application of Cor and sorbitol resulted in a dramatic enhancement in the expression of artemisinin biosynthetic genes and artemisinin production owing to a synergistic or potentiating result.

**Keywords** Antioxidant enzyme · Osmotic stress · Relative gene expression · Secondary metabolite

## Abbreviations

AA	Artemisinic aldehyde	CSC	Cell suspension culture
ADS	Amorpha-4, 11-diene synthase	CYP71AV1	Amorphadiene-12-hydroxylase
ALDH1	Aldehyde dehydrogenase 1	DBR2	Artemisinic aldehyde $\Delta 11(13)$ reductase
APX	Ascorbate peroxidase	DCW	Dry cell weight
Cor	Coronatine	DHAA	Dihydroartemisinic aldehyde
		EDTA	Ethylenediaminetetracetic acid
		GA3	Gibberellic acid
		GR	Glutathione reductase
		$\text{H}_2\text{O}_2$	Hydrogen peroxide
		HPLC	High-performance liquid chromatography
		Kin	Kinetin
		LSD	Least significant difference
		MDA	Malondialdehyde
		MeJA	Methyl jasmonate
		NAA	1-Naphthaleneacetic acid
		PVP-40	Polyvinylpyrrolidone
		RED1	Dihydroartemisinic aldehyde reductase

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ROS	Reactive oxygen species
SE	Standard error
SM	Secondary metabolite
SOD	Superoxide dismutase
TBA	Thiobarbituric acid
TCA	Trichloroacetic acid

## Introduction

Artemisinin is an efficient anti-malarial drug and it possesses biological activity against a wide range of cancers (Efferth 2009, 2017). Some *Artemisia* species produce artemisinin which is a sesquiterpene lactone (Ranjbar et al. 2015; Hamidi et al. 2018; Salehi et al. 2018a, b). The main source of artemisinin is *Artemisia annua* where the artemisinin production is subjected to environmental and seasonal changes. Plant tissue and cell cultures are the viable renewable resource of plant natural products, but a very low amount of secondary metabolites (SMs) is synthesized in cultures compared with those in intact plants (Oksman-Caldentey and Inzé 2004; Naik and Al-Khayri 2016; Salehi et al. 2017, 2018c, 2019a, b). Phyton Biotech (<https://phytonbiotech.com/>) has received \$400,000 in 2017 to find out if it can manufacture artemisinin directly by fermenting cells from *A. annua* itself. Plants synthesize SMs from primary metabolites for the defense aim (Naik and Al-Khayri 2016). Elicitor is a stress agent that increases the SM production in a particular tissue, organs, and cells (Naik and Al-Khayri 2016). Elicitor induces stress which results in the activation of several defense-related genes or inactivation of non-defense-related genes, transient phosphorylation/dephosphorylation of proteins, and enzyme expression (Narayani and Srivastava 2017). The depth science of the metabolic response to elicitation in plant cells, as well as a good comprehension of the mechanisms responsible for these results, can result in the cost-effective and sustainable commercial production of plant SMs (Ramirez-Estrada et al. 2016). Even overexpressing the biosynthetic pathway key genes in plant cells still require elicitation for high production of a related SM (Naik and Al-Khayri 2016; Ramirez-Estrada et al. 2016). Hence, choosing the most efficient elicitor for enhancing SM biosynthesis in plant cell cultures is important. In the exponential growth phase of plant cell cultures, the primary metabolite precursors are needed for biomass formation and therefore, many metabolites are produced at low contents, or not all. The induction of SM production from primary compounds is more efficient in the stationary growth phase (Cusido et al. 2002; Malik et al. 2011; Ramirez-Estrada et al. 2016). The artemisinin production has been outstandingly increased in *A. annua* cell suspension cultures (CSC) by treatment with methyl jasmonate (Baldi and Dixit 2008), potassium nitrate (KNO<sub>3</sub>, Keng et al. 2010),  $\beta$ -cyclodextrins

and methyl jasmonate (MeJA, Durante et al. 2011). Moreover, increased artemisinin production has been gotten in *A. annua* hairy root cultures by treating with GA3 (Smith et al. 1997), *Colletotrichum mycelia* (Wang et al. 2001), fungal cerebroside and nitric oxide (Wang et al. 2009), oligogalacturonides (Zhang et al. 2010), and elicitor derived from *Verticillium dahlia* (Wang et al. 2000). The studies showed that osmotic stress is effective to stimulate the production of the diverse SMs in the plant cell cultures (Do and Cormier 1990; Zhang et al. 1995; Kim et al. 2001; Wu et al. 2005; Shi et al. 2007; Liu and Cheng 2008; Naik and Al-Khayri 2016; Hussein and Aqlan 2011; Sarmadi et al. 2018). The osmotic stress can be induced by osmotic agents such as sucrose, glucose and polyols (sugar alcohols) in plant cell cultures to increase SM production (Kim et al. 2001; Shi et al. 2007; Hussein and Aqlan 2011; Sarmadi et al. 2018). Sorbitol (C<sub>6</sub>H<sub>14</sub>O<sub>6</sub>, non-metabolic sugar) raise osmotic potential in the medium only and were not applied as nutrient sources in plant cell cultures, therefore it was applied as an osmoticum for increasing medium osmolality and developing osmotic stress in the plant CSCs. Reduction in tissues water absorption lead to growth decrement and physiological changes (Akula and Ravishankar 2011). Coronatine (Cor) functions as a molecular imitator of the isoleucine-conjugated form of jasmonic acid (JA-Ile, Katsir et al. 2008), while it is more stable and less toxic than MeJA (Onrubia et al. 2013), and as a result, possesses a similar action mechanism to the elicitor MeJA. Interestingly, cultures treated with Cor are usually more productive when compared with higher concentrations of MeJA in the same culture conditions (Ramirez-Estrada et al. 2016). Cor usually activates the production of SMs in the plant cell cultures at concentrations lower compared with that of MeJA (Onrubia et al. 2013; Ramirez-Estrada et al. 2016). In addition to increasing the SM production, Cor can alleviate biotic and abiotic stress in plants (Zare Dehabadi et al. 2013; Ceylan et al. 2013; Zhou et al. 2015; Ahmad et al. 2016). The simultaneous application of two different elicitors has been shown to be more efficient owing to a synergistic or potentiating result such as stimulation of artemisinin production by combined cerebroside and nitric oxide elicitation in *A. annua* hairy roots (Wang et al. 2009), significant increase of artemisinin yields in *A. annua* CSC elicited by combination of  $\beta$ -cyclodextrin and methyl jasmonate (MeJA, Durante et al. 2011), the highest artemisinin content in *A. annua* hairy root culture elicited by combination of MeJA and cell homogenate of *Piriformospora indica* (Ahlawat et al. 2014), effective enhancement of total tanshinone by combined use of osmotic stress (sorbitol) and yeast elicitor in *Salvia miltiorrhiza* Bunge hairy-root cultures (Shi et al. 2007); the highest levels of taxol in *Taxus media* and *Taxus globosa* CSC by combined use of Cor and  $\beta$ -cyclodextrin (Ramirez-Estrada et al. 2015), dramatic enhancement of taxol production in *Corylus avellana* CSC

elicited by salicylic acid pretreatment and ultrasound stress (Rezaei et al. 2011), and significant effect of combined use of salicylic acid and osmotic stress on production of taxane in the callus culture of *Taxus baccata* (Sarmadi et al. 2018). To the best of our knowledge, there are no published studies on the influence of Cor, osmotic stress, and the synergistic result of Cor and osmotic stress on artemisinin production in *A. annua* CSC. The current study was carried out to assess the results of Cor pretreatment and different concentrations of sorbitol on the growth, biochemical traits, expression of artemisinin biosynthetic genes, and artemisinin production in *A. annua* CSC for the first time.

## Materials and methods

### Cell suspension cultures

*Artemisia annua* cv. Anamed which is regarded as a high artemisinin cultivar (Salehi et al. 2018a, b) was used for the establishment of CSC, employing the procedure described by Baldi and Dixit (2008) with slight modifications. Callus was developed from aseptically germinated seedlings and maintained on MS medium, supplemented with NAA and Kin (0.5 mg/L of each) as growth regulators and solidified with 8 g/L agar agar. The CSC was developed with cultivating 5 g fresh callus into 250 mL flasks, containing 100 mL of liquid MS media with 30 g/L sucrose and 0.1 mg/L of each of NAA and Kin. The flasks were incubated on a rotary shaker (125 rpm) at  $25 \pm 2$  °C with a photoperiod of 16/8 h light/dark cycle. The CSCs were then subcultured until the cells reached homogeneity.

### Elicitation treatments

About  $1 \pm 0.1$  g of cells (fresh mass) was placed to 100 mL flasks, containing 30 mL of the cell culture medium. At the preliminary experiment, three concentrations (0.01, 0.05, and 0.5  $\mu$ M) of Cor elicitor were added to the CSC on the 14th day (three days before the stationary phase). High artemisinin content and no effect on biomass formation was detected on 0.05  $\mu$ M Cor at 48 h after elicitation and therefore, this concentration was selected as Cor pretreatment. Experimental treatments included control, Cor, sorbitol, and Cor  $\times$  sorbitol treatments. The CSC pretreated with 0.05  $\mu$ M of Cor on the 14th day (three days before the stationary phase) for 48 h. For osmotic stress treatment, the liquid medium in the culture flasks was decanted and replaced with fresh medium, containing 30 g/L sucrose plus or minus sorbitol at selected concentrations (0, 20, 30, and 40 g/L) on the 16th day (one day before the stationary phase). The cell growth, hydrogen peroxide ( $H_2O_2$ ) and malondialdehyde (MDA) contents, enzyme activities, and artemisinin content

were measured on the 19th day (according to the preliminary test, the highest amount of artemisinin was obtained on the 19th day). The expression of artemisinin biosynthesis genes was determined at 30 min, 4 h, and 24 h after treating with sorbitol.

### Measurement of cell growth

The cell growth was determined by measuring the dry cell weight (DCW). In brief, separation of biomass from medium was done by filtration. Removing the residual medium was done by washing with distilled water. Then biomass was lyophilized to constant weight, using a vacuum-freeze drier.

### $H_2O_2$ determination

$H_2O_2$  content was assessed employing the procedure described by Alexieva et al. (2001). In brief, addition of 500  $\mu$ L of 0.1% trichloroacetic acid (TCA) freeze-dried cells extract supernatant was done to 500  $\mu$ L of phosphate buffer (100 mM, pH 7.0) and 2000  $\mu$ L of KI (1 M). The reaction was made for 1 h in darkness and absorbance recorded at 390 nm. The  $H_2O_2$  concentration was measured, using a  $H_2O_2$  standard curve.

### Lipid peroxidation assay

The lipid peroxidation level was measured using determination of MDA amount generated by the thiobarbituric acid (TBA) reaction (Heath and Packer 1968). Homogenization of 0.3 g freeze-dried cells was done in 1500  $\mu$ L of 0.1% TCA and then centrifuged at  $10,000 \times g$  for 5 min. Addition of 4 mL of 0.5% TBA in 20% TCA was done to 1000  $\mu$ L of supernatant. Heating of mixture at 95 °C for 30 min, cooling rapidly and centrifuging (at  $10,000 \times g$  for 15 min) were done. The absorbance at 600 nm and 532 nm was subtracted and then MDA content was computed employing the extinction coefficient of 155/mM cm.

### Determination of enzyme activities

Antioxidant enzymes was extracted by homogenizing 0.5 g of freeze-dried cells in 2.5 mL of sodium phosphate buffer (50 mM, pH 7), and then centrifuging at  $10,000 \times g$  and 4 °C for 15 min (Gapińska et al. 2008). Sodium phosphate buffer contained 1 mM Ethylenediaminetetracetic acid (EDTA) and 1% Polyvinylpyrrolidone (PVP-40).

### Ascorbate peroxidase (EC 1.11.1.11)

Ascorbate peroxidase (APX) activity was assayed employing the procedure described by Nakano and Asada (1981). The reaction mixture contained 50 mM K-phosphate buffer (pH

7), 0.5 mM ascorbate, 0.1 mM EDTA Na<sub>2</sub>, 0.1 mM H<sub>2</sub>O<sub>2</sub> and 0.1 mL of enzyme extract in a final assay volume of 1 mL. The reduction in absorbance at 290 nm was recorded and enzyme activity was computed using an extinction coefficient of 2.8/mM cm.

### Glutathione reductase (EC 1.6.4.2)

Glutathione reductase (GR) activity was assayed according to the method of Foyer and Halliwell (1976). The assay medium contained 25 mM Na-phosphate buffer (pH 7.8), 0.5 mM oxidized form of glutathione (GSSG), and 0.12 mM NADPH and 0.1 mL enzyme extract in a final assay volume of 1 mL. NADPH oxidation was followed at 340 nm. Activity was computed, using the extinction coefficient of NADPH (6.2/mM cm). One unit of GR was defined as 1 mmol/mL GSSG reduced per min.

### Superoxide dismutase (EC 1.15.1.1)

The activity of superoxide dismutase (SOD) was assessed using the procedure depicted by Giannopolitis and Ries (1977). The reaction was constructed using the mixture of potassium phosphate buffer (50 mM, pH 7.8), EDTA (0.1 mM), 75  $\mu$ L nitroblue-tetrazolium (NBT), methionine (13 mM), 2  $\mu$ M riboflavin and 100  $\mu$ L of enzyme extract and it was stirred and put 50 cm below a light bank (eight fluorescent lamps 15 W) for 10 min. Finally, absorbance was recorded at 560 nm. One unit of SOD activity was described as enzyme content that inhibited 50% of NBT photoreduction.

### Quantification of artemisinin

After cell separation from CSC, the cell samples were frozen in liquid nitrogen and lyophilized overnight. 50 mg lyophilized cell powder were extracted with 4 ml *n*-hexane for 16 h under magnetic stirring, then for 15 min in an ultrasonic water bath. The extract was centrifuged at 4000  $\times$ g for 10 min, and the supernatant was removed and placed in new tubes. The pellet was resuspended in 1 ml *n*-hexane and centrifuged again for 10 min at 4000  $\times$ g. This second supernatant was added to the first, and the pellet discarded (Caretto et al. 2011). Extracts were then dried under vacuum. Within 24 h, samples were reconstituted in 1 ml of acetonitrile, filtered through pre-wetted 0.2  $\mu$ m (25 mm) nylon Millex-GN filters (Millipore Corporation, Bedford, MA), connected to disposable 3-ml syringes (Salehi et al. 2018a, b). The artemisinin content of the extracts was determined by an HPLC (high-performance liquid chromatography) system (Waters, USA), equipped with a C18 column (NUCLEODUR 100-5 C18 ec, 250 mm  $\times$  4.6 mm, China) and detection was conducted at 210 nm wavelength. The acetonitrile:water 65:35%

(v/v) was used as a mobile phase with 1 mL/min flow rate (Lapkin et al. 2009; Salehi et al. 2018a, b). Artemisinin was identified by comparison with artemisinin standard. Artemisinin content was determined, using calibration curve of artemisinin standard (Sigma).

### Real-time RT-PCR

Relative expression of artemisinin biosynthetic genes including *amorpha-4,11-diene synthase (ADS)*, *amorphadiene-12-hydroxylase (CYP)*, *aldehyde dehydrogenase 1 (ALDH1)*, artemisinic aldehyde  $\Delta$ 11(13) reductase (*DBR2*) and *dihydroartemisinic aldehyde reductase (RED1)*, Fig. 1) were assessed. Total RNA extraction, genomic DNA removal, cDNA synthesis, and qPCR were done; employing the procedure described by Salehi et al. 2018a, b.  $\beta$ -Actin was used as reference genes. The qPCR primers were obtained from Salehi et al. (2018a). Amplicon efficiencies of all primer pairs were computed with cDNA serial dilutions using this formula:  $E = 10^{-1/\text{slope}} - 1$ . Relative expression levels were calculated using the  $(1 + E)^{-\Delta\Delta\text{CT}}$  method (Pfaffi 2001). The transcription levels of five artemisinin biosynthesis genes under elicitation conditions including Cor, sorbitol and Cor + sorbitol conditions were compared relative to control condition, which was chosen as the reference condition.

### Statistical analysis

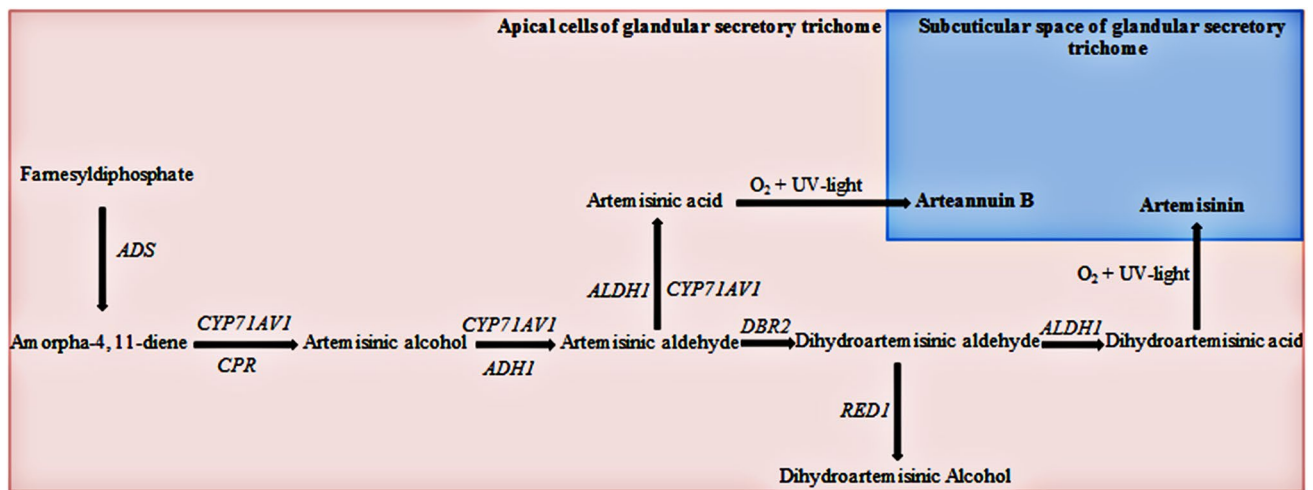
The study was conducted, using a factorial experiment based on a complete randomized block design (CRBD) with three replications. The data were tested for the normality and then analyses of variances were done, using PROC GLM of SAS (SAS Institute 2002). Mean comparisons were conducted using Fisher's least significant differences (LSDs) at 0.01 probability levels. Also, the standard error (SE) was calculated.

## Results and discussion

### Growth changes

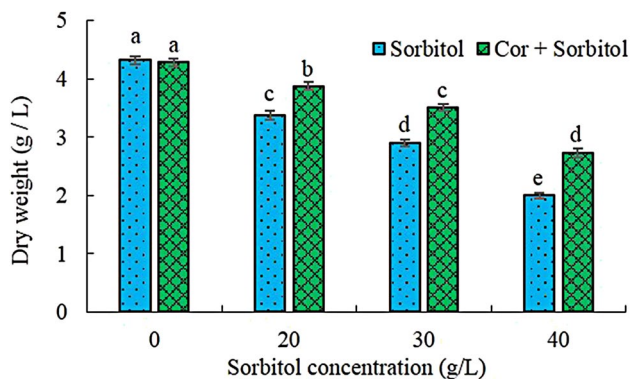
DCW (g) declined significantly ( $p < 0.01$ ) as the concentration of sorbitol increased, the least cell weight was observed in the medium containing 40 g/L sorbitol (1.99 g/L, Fig. 2). Adding sorbitol in the culture medium increased osmotic stress and caused cell degradation and consequently cell growth decrement. Also, other researchers reported that intracellular water and weight of plant calli decreased with increasing osmotic stress (Tholakalabavi et al. 1994; Akula and Ravishankar 2011; Sarmadi et al. 2018). Cor pretreatment had no significant effect on DCW on the sorbitol-free medium (Cor + 0 sorbitol, Fig. 2), while it increased DCW





**Fig. 1** Summary of artemisinin biosynthesis pathway. *Aa-ADS* amorpha-4,11-diene synthase, *Aa-CYP71AV1* amorpha-4,11-diene-12-hydroxylase, *Aa-CPR* cytochrome P450 reductase, *Aa-ADH1* alcohol

dehydrogenase 1, *Aa-ALDH1* aldehyde dehydrogenase 1, *Aa-DBR2* artemisinic aldehyde  $\Delta$ 11(13) reductase, *Aa-RED1* dihydroartemisinic aldehyde reductase (Salehi et al. 2018a)



**Fig. 2** Effects of Cor pretreatment and different concentrations of sorbitol treatments on cell growth of *A. annua* L. on 19th day of culture cycle. Error bars represent SE ( $n=3$ ). Means followed by the same letter are not significantly different according to LSD at 0.01 probability level

at all sorbitol concentration (Fig. 2). Cor can alleviate abiotic stress in plants (Zare Dehabadi et al. 2013; Ceylan et al. 2013; Zhou et al. 2015; Ahmad et al. 2016). A Cor pretreatment increased the relative growth rate of chickpea roots under PEG-induced osmotic stress, heat stress, and combined stresses (Ceylan et al. 2013) and fresh mass of *Ocimum basilicum* under arsenic treatment (Zare Dehabadi et al. 2013). More cell growth after Cor pretreatment suggest an increased tolerance to osmotic stress induced by sorbitol.

## H<sub>2</sub>O<sub>2</sub> and MDA content

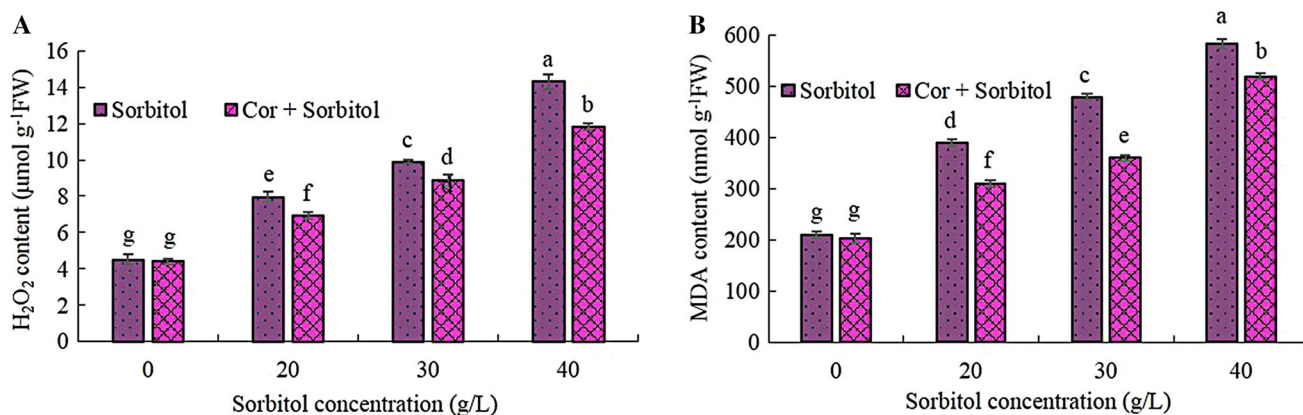
MDA and H<sub>2</sub>O<sub>2</sub> levels are indicators of ROS mediated destruction to cell membranes and oxidative stress, respectively under stressful and elicitor conditions (Hossain et al.

2015; You and Chan 2015; Sarmadi et al. 2018). In the current study, sorbitol significantly ( $p < 0.01$ ) enhanced the H<sub>2</sub>O<sub>2</sub> and MDA contents of the cells (Fig. 3). The highest contents of H<sub>2</sub>O<sub>2</sub> (14.32  $\mu$ mol/g FW) and MDA (583.33 nmol/g FW) were detected under 40 g/L sorbitol (Fig. 3). After the Cor pretreatment, less contents of H<sub>2</sub>O<sub>2</sub> and MDA were observed under all sorbitol concentrations, while Cor had no effect on H<sub>2</sub>O<sub>2</sub> and MDA contents in the sorbitol-free medium (Cor + 0 sorbitol, Fig. 3).

Environmental stresses and elicitors enhance the free radicals and reactive oxygen species (ROS) levels and cause lipid peroxidation and bio-membrane degradation (Wang et al. 2008; Hossain et al. 2015; You and Chan 2015). The level of membrane damage directly depends on the oxidative stress intensity and the ROS level. In the current study, H<sub>2</sub>O<sub>2</sub> level increased under sorbitol treatment and consequently MDA content increased (Fig. 3). Cor pretreatment reduced H<sub>2</sub>O<sub>2</sub> and MDA levels indicating Cor can stabilize and protect cell membranes. This result is consistent with those of Wang et al. (2008), Zare Dehabadi et al. (2013) and Ceylan et al. (2013) reporting a Cor-induced reduction of H<sub>2</sub>O<sub>2</sub> and MDA levels under drought stress, arsenic toxicity, and combined osmotic and heat stresses, respectively.

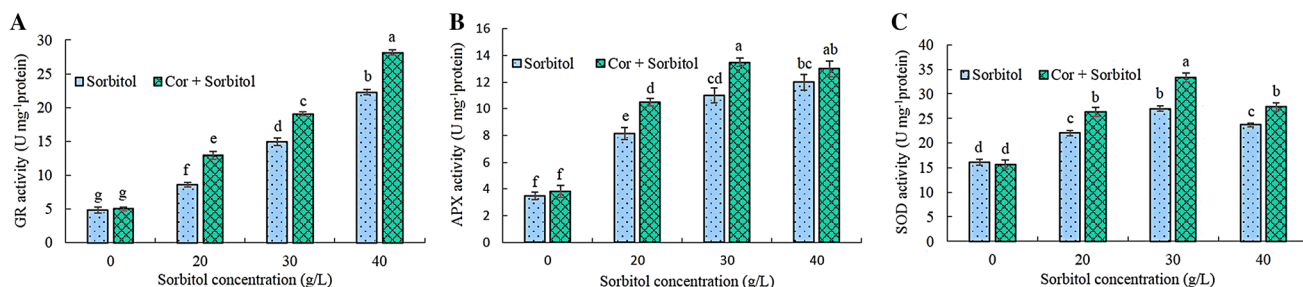
## Antioxidant enzyme activity

The activity of GR, APX, and SOD enzymes increased under all sorbitol treatments compared with those in control (Fig. 4). Increasing the antioxidant activities is a stress tolerance mechanism for preventing oxidative damage (Wang et al. 2008; Zare Dehabadi et al. 2013; Ceylan et al. 2013; Hossain et al. 2015; You and Chan 2015). The main mechanism for detoxifying the cells from free



**Fig. 3** Effects of Cor pretreatment and different concentrations of sorbitol treatments on H<sub>2</sub>O<sub>2</sub> (a) and MDA (b) content of *A. annua* cells on 19th day of culture cycle. Error bars represent SE (n=3).

Means followed by the same letter are not significantly different according to LSD at 0.01 probability level



**Fig. 4** Effects of Cor pretreatment and different concentrations of sorbitol treatments on the antioxidant enzyme activities of *A. annua* cells on 19th day of culture cycle. GR (a), APX (b), and SOD (c). Error bars represent SE (n=3).

Error bars represent SE (n=3). Means followed by the same letter are not significantly different according to LSD at 0.01 probability level

radicals is the enzymatic antioxidant system which functions for adapting and surviving the plant under stress conditions (Hossain et al. 2015; You and Chan 2015). The first and most important enzyme in the detoxifying the cells from ROS is SOD which converts radical superoxide (O<sub>2</sub>) to H<sub>2</sub>O<sub>2</sub> in the chloroplasts, mitochondria and cytosol and therefore it plays a fundamental role in the cell defense mechanisms against radical hydroxyl (OH) formation (You and Chan 2015). In the next step, enzymes such as GR and APX scavenge the produced H<sub>2</sub>O<sub>2</sub> (Hossain et al. 2015; You and Chan 2015). SOD enzyme had more activity under 30 g/L sorbitol treatment compared with 40 g/L sorbitol treatment (Fig. 4c). The activity of APX enzyme did not show a significant difference in 30 and 40 g/L sorbitol treatments (Fig. 4b). GR enzyme had higher activity under 40 g/L sorbitol treatment compared with 30 g/L sorbitol treatment (Fig. 4a). The higher activity of SOD antioxidant enzyme under 30 g/L sorbitol compared with that under 40 g/L sorbitol (Fig. 4c) indicates that its efficiency is higher in mild stress compared to severe stress. In severe stress, free radicals are generated more

quickly than they can be scavenged, and thus the destruction of enzymes will be more. The less activity of SOD enzyme under 40 g/L sorbitol compared with that under 30 g/L sorbitol (Fig. 4c) may be due to less tolerance of *A. annua* cells under high level of oxidative stress. Considering that SOD enzyme is the first enzyme in detoxification of cells from ROS and less activity of SOD under 40 g/L sorbitol compared with that under 30 g/L sorbitol (Fig. 4c), it seems that 40 g/L sorbitol is not proper elicitor for enhancing SM production.

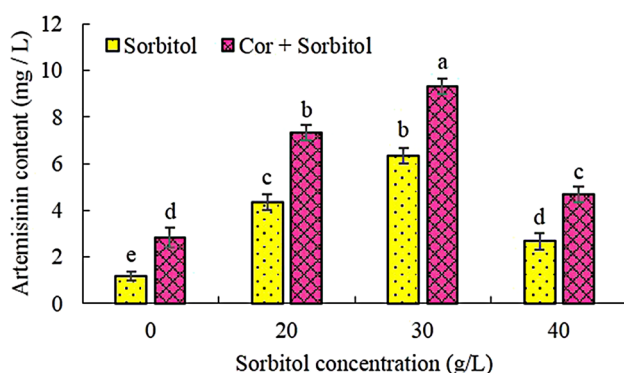
Cor pretreatment enhanced the activity of three studied antioxidant enzymes under different sorbitol concentration (Fig. 4). The antioxidant system neutralizes the impact of oxidative stress in plants and researchers showed that the exogenous use of Cor in stress conditions improves the antioxidant enzyme activities (Wang et al. 2008; Zare Dehabadi et al. 2013; Ceylan et al. 2013). Sarmadi et al. (2018) showed that salicylic acid pretreatment enhanced antioxidant enzyme activities under different glucose levels in a *T. baccata* callus culture.

## Artemisinin content

The artemisinin content was significantly higher in all sorbitol concentrations compared with that in the control, having a peak at 30 g/L sorbitol and declining at 40 g/L (Fig. 5). Previous studies reported that SM production increased in plant cell and tissue cultures elicited by sucrose, mannitol and sorbitol (Suzuki 1995; Zhang et al. 1996; Kim et al. 2001; Shi et al. 2007; Namdeo et al. 2007; Hussein and Aqlan 2011; Sarmadi et al. 2018). The SM production decreases under severe stress due to increased oxidative degradation (Sarmadi et al. 2018). The Cor pretreatment enhanced artemisinin content in CSC (Fig. 5). Cor increased both SM production and expression of SM biosynthetic genes in plant CSCs (Ramirez-Estrada et al. 2016). The application of Cor plus sorbitol resulted in a dramatic enhancement in artemisinin content. The highest content of artemisinin (9.33 mg/L, 8-fold, Fig. 5) was observed with the combination of Cor and 30 g/L sorbitol. In addition to increasing the SM gene expression, Cor alleviate abiotic stress (Fig. 3) and resulted in more cell growth (Fig. 2), which can lead to higher artemisinin production per liter. The simultaneous application of Cor and osmotic stress was more efficient to produce SM owing to a synergistic or potentiating result. Researchers reported that simultaneous application of two different elicitors is more efficient owing to a synergistic or potentiating result (Shi et al. 2007; Wang et al. 2009; Durante et al. 2011; Rezaei et al. 2011; Ahlawat et al. 2014; Ramirez-Estrada et al. 2015; Sarmadi et al. 2018).

## Gene expression

The qPCR method was used to understand the possible results of the Cor or/and sorbitol treatments on the expression of artemisinin biosynthetic genes (Fig. 1) starting from



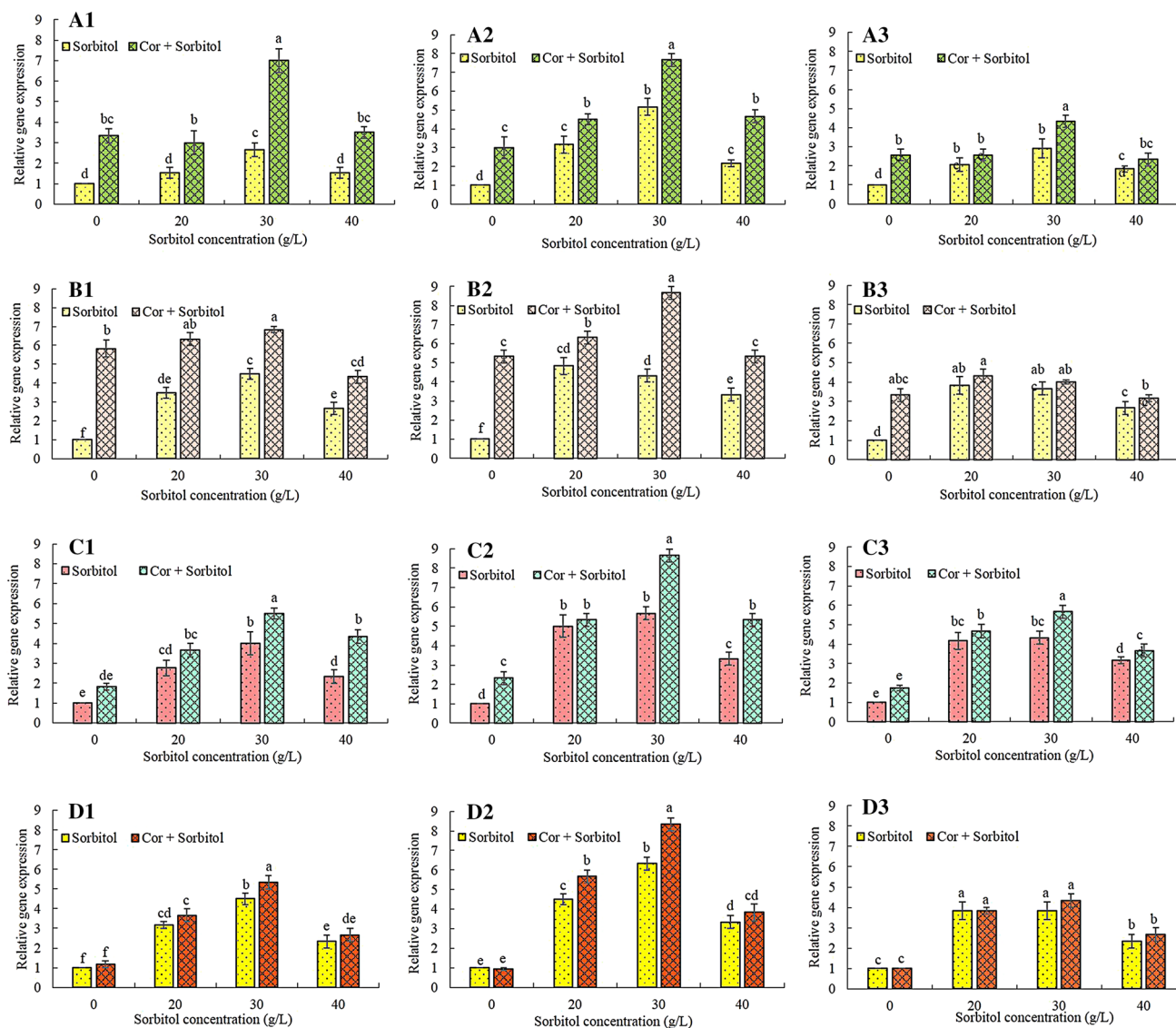
**Fig. 5** Effects of Cor pretreatment and different concentrations of sorbitol treatments on artemisinin production in *A. annua* L. cell suspension culture on 19th day of culture cycle. Error bars represent SE ( $n=3$ ). Means followed by the same letter are not significantly different according to LSD at 0.01 probability level

30 min up to 24 h. The expression of *RED1* gene was not significantly influenced by Cor or/and sorbitol treatments and therefore the results were not presented. The expression levels of *ADS*, *CYP71AV1*, *ALDH1* and *DBR2* genes increased under 20 and 30 g/L sorbitol treatments compared with those in control at all times (Fig. 6). The low content of  $H_2O_2$  as a signaling molecule could modify the expression of SM genes (Sarmadi et al. 2018). The osmotic stress induced by low concentrations of sorbitol triggered oxidative cascade which enhances expression of artemisinin biosynthetic genes (Fig. 6). The application of Cor plus sorbitol resulted in a dramatic enhancement in the expression of *ADS*, *CYP71AV1*, *ALDH1* and *DBR2* genes at all concentrations (Fig. 6). The expression of these genes peaked at 30 g/L sorbitol and decreased at 40 g/L sorbitol (Fig. 6), probably because of higher oxidative stress.

Results indicated that expression of *ADS*, *CYP71AV1*, *ALDH1* and *DBR2* genes were enhanced about 7.66, 8.67, 8.67, and 8.33-fold, respectively at 4 h after Cor + 30 g/L sorbitol joint treatments (Fig. 6). Sorbitol up-regulated *DBR2* gene, while Cor had no effect on *DBR2* gene expression (Fig. 6d). Cor alleviated oxidative stress (Fig. 3) by enhancing antioxidant enzyme activities (Fig. 4) and possibly, other physiological mechanisms. Furthermore, Cor increased the expression of *ADS*, *CYP71AV1*, and *ALDH1* genes but did not affect the expression of the *DBR2* gene (Fig. 6). The oxidative stress had a positive result on the expression of *ADS*, *CYP71AV1*, *ALDH1*, and especially *DBR2* genes.

The relative turnover potential of the artemisinin biosynthetic enzymes can be computed, using the application of relative transcript levels  $[(1 + E)^{-\Delta\Delta CT}$  method, Pfaffi 2001] in combination with kinetic data  $[(1 + E)^{-\Delta\Delta CT} \times K_{cat} (S^{-1})$ , Olofsson et al. 2011; Salehi et al. 2018a]. For this estimate, it was presumed that the number of enzyme active sites was proportional to the transcription rate and that the enzymes were acting at substrate saturation with an optimal NADPH/NADP<sup>+</sup> ratio (Olofsson et al. 2011; Salehi et al. 2018a). In this condition, the  $k_{cat}$ -value is a well indicator of the conversion of substrate to product. The highest conversion potential of the *ADS* (0.14) and *ALDH1* [(34.17, on dihydroartemisinic aldehyde, DHAA) and (6.66, on artemisinic aldehyde, AA)] was observed in Cor + sorbitol treatment followed by Cor treatment and the highest conversion potential of *DBR2* (6.01) was observed in Cor + sorbitol treatment followed by sorbitol treatment (Table 1). In Cor + sorbitol, the highest relative turnover potential of *ADS*, *ALDH1*, and *DBR2* was observed. The Cor + sorbitol treatment was more efficient owing to this synergistic result. The relative turnover of *ALDH1* and *DBR2* are nearly equal in control, sorbitol, and Cor + sorbitol treatments, while the potential conversion capacity of *ALDH1* (3.49) was 2.9-fold than that of *DBR2* (1.21) in Cor treatment (Table 1). Aa-*DBR2* and Aa-*ALDH1* are acting on the





**Fig. 6** Effects of Cor pretreatment and different concentrations of sorbitol treatments on relative expression of *ADS* (a), *CYP71A1* (b), *ALDH1* (c), and *DBR2* (d) genes of *A. annua* cells at 30 min (a1, b1, c1, and d1), 4 h (a2, b2, c2, and d2), and 24 h (a3, b3, c3, and d3)

after sorbitol elicitation. Error bars represent SE (n=3). Means followed by the same letter are not significantly different according to LSD at 0.01 probability level

same pool of intermediate. Aa-DBR2 enzyme converts AA into DHAA which is a precursor of artemisinin, while Aa-ALDH1 catalyzed the transformation of AA into artemisinic acid which is a precursor of arteannuin B (Fig. 1). Since, the relative turnover of Aa-ALDH1 is higher than Aa-DBR2 under Cor treatment. Hence, it was deduced that artemisinic acid/arteannuin B was produced more compared with artemisinin under Cor treatment.

## Conclusion

The effects of Cor and sorbitol treatments on *A. annua* CSC are studied for the first time. Sorbitol treatment enhanced the membrane lipid peroxidation (Fig. 3b) through inducing oxidative stress and increasing  $H_2O_2$  level (Fig. 3a). The low oxidative stress (low content of



**Table 1** Estimation of relative turnover potential for four enzymes of artemisinin biosynthesis using the  $(1 + E)^{-\Delta\Delta CT}$  method

Enzyme	Substrate	$K_m$ ( $\mu\text{M}$ )	$K_{cat}$ ( $\text{S}^{-1}$ )	$\Delta\Delta CT$	Normalized transcript amount relative to REDI $(1 + E)^{-\Delta\Delta CT}$				Relative turnover				Reference for kinetic constants
					Con	Cor	Sor 30	Cor+Sor 30	Con	Cor	Sor 30	Cor+Sor 30	
ADS	FDP	2	0.004	-2	-4.59	24.02	14.62	34.06	0.02	0.1	0.06	0.14	Picaud et al. (2005)
DBR2	AA	19	2.600	2	1.1	0.47	1.12	2.31	0.65	1.21	2.91	6.01	Zhang et al. (2008)
ALDH1	DHAA	8.8	7.700	1	-1.22	0.5	2.33	4.44	3.85	17.94	15.4	34.17	Teoh et al. (2009)
	AA	2.6	1.500	1	-1.22	0.5	2.33	4.44	0.75	3.49	3	6.66	
RED1	DHAA	67	0.280	0	0	1	1	1	0.28	0.28	0.28	0.28	Rydén et al. (2010)

Con control, Cor coronatine, Sor 30 30 g/L sorbitol

$\text{H}_2\text{O}_2$  as a signaling molecule) increased the expression of *ADS*, *CYP71AV1*, *ALDH1*, and especially *DBR2* genes (Fig. 6). Also, artemisinin content (Fig. 5) and antioxidant enzyme activities (Fig. 4) increased under sorbitol treatment, which is a physiological defense response to osmotic stress. Pretreatment of *A. annua* cells with Cor increased antioxidant enzyme activity (Fig. 4) which regulates the ROS rate and increased the cell tolerance to sorbitol stress; resulting in increased cell biomass (Fig. 2). Moreover, Cor enhanced the expression of *ADS*, *CYP71AV1*, and *ALDH1* genes but did not affect the expression of the *DBR2* gene (Fig. 6). Uppalapati et al. (2005) showed that Cor enhances proteinase inhibitors and as a result enzymes involved in the SM biosynthesis pathway would possess a more continuous activity in cell cultures treated with this elicitor, resulting in higher SM contents. In Cor + sorbitol, the highest relative turnover potential of *ADS*, *ALDH1*, and *DBR2* was observed (Table 1). Also, relative turnover of *ALDH1* and *DBR2* are nearly equal in Cor + sorbitol treatment (Table 1). The simultaneous application of Cor and osmotic stress was more efficient to produce SM owing to a synergistic or potentiating result.

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## Compliance with ethical standards

**Conflict of interest** The authors declare that they have no competing interests.

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