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Baccharis trimera (Less.) DC responses to water restriction

Respostas de Baccharis trimera (Less.) a restrição de água

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ARTICLE	ABSTRACT
Received: 14 July 2020 Accepted: 22 June 2021	Climate change can cause imbalances in plants. <i>Baccharis trimera</i> is a specie usually found in harsh conditions and has medicinal and agricultural properties. Thus, this study aimed to evaluate the biochemical and physiological responses of this plant under water restriction (WR) levels. <i>B. trimera</i> samples were identified and collected in Erechim/RS and propagated in a greenhouse. When acclimated, they were subjected to 0, 25, 75 and 100% WR, determined according to the water estruction in the soil by confidence of the properties.
Key words: Drought Medicinal plant Oxidative stress Biochemistry Proline	saturation in the soft by capitary action. After 50 days of treatment, the physiological responses: growth, and fresh and dry biomass; and the biochemical responses: activity of superoxide dismutase (SOD), guaiacol peroxidase (GP) and ascorbate peroxidase (APX) enzymes, proline, protein and hydrogen peroxide content, and lipid peroxidation, were determined. Data were submitted to regression analysis and Pearson correlation. The WR of 27.37%, on average, induced an increase in physiological parameters, but the root growth was impaired in conditions above 50% of WR. With the increase in WR there was an increase in the activity of SOD in the shoot and APX in the root. In low WR conditions, proline contents were maintained. Therefore, with low levels of WR, around 27%, <i>B. trimera</i> has increase in root growth and root and shoot biomass. Proline, and SOD and APX activity are a pathway that scavenging the stress generated by WR on <i>B. trimera</i> .
	RESUMO
Palavras-chave: Estiagem Planta medicinal Estresse oxidativo Bioquímica Prolina	Alterações climáticas podem causar desequilíbrios nas plantas. <i>Baccharis trimera</i> é uma espécie frequentemente encontrada em condições adversas e apresenta propriedades medicinais e para uso na agricultura. Assim, objetivou-se avaliar as respostas bioquímicas e fisiológicas desta planta em níveis de restrição de água (RA). Amostras de <i>B. trimera</i> foram identificadas e coletadas em Erechim - RS e propagadas em casa de vegetação. Quando aclimatadas foram submetidas a 0, 25, 75 e 100% de RA, determinada de acordo com a saturação de água no solo por capilaridade. Após 30 dias de tratamento foram determinadas respostas fisiológicas: crescimento e biomassa fresca e seca; e respostas bioquímicas: atividade de enzimas superóxido dismutase (SOD), guaiacol peroxidase (GP) e ascorbato peroxidase (APX), conteúdo de prolina, proteína e peróxido de hidrogênio, e peroxidação de lipídeos. Os dados foram submetidos a análises de regressão e correlação de Pearson. A RA média de 27,37% induziu aumento nos parâmetros fisiológicos avaliados, porém o crescimento das raízes foi prejudicado em condições de RA acima de 50%. Com o aumento na RA houve o aumento na atividade das enzimas SOD na parte aérea e de APX na raiz. Em condição de baixa RA verificou-se a manutenção do conteúdo de prolina. Portanto, com baixos níveis de RA, em torno de 27%, <i>B. trimera</i> tem aumento no crescimento de raiz e na biomassa da parte aérea e raiz. Prolina, SOD e APX são uma via de eliminação do estresse gerado pela RA em <i>B. trimera</i> .



INTRODUCTION

Plants are often exposed to environmental stresses, as drought, light excess, high and low temperatures, freezing, flooding, salinity, heavy metals, among others (BOWNE et al., 2018; LEUNG, 2018). These factors lead to biochemical and physiological imbalances in plants, causing reductions in productivity. Drought is one of the most worrying types of stress and can drastically decrease crop reproduction. Hydric stress in plants affects factors such as stomatal behaviour, reverse mobilization, enzymatic activity, leaf expansion, growth (FAROOQ et al., 2009) and can lead to oxidative stress (MITTLER, 2002) as a result of metabolic interruption due to cytoplasmic free water loss (SMIRNOFF, 1993).

Oxidative stress is characterized by accumulation of reduced and very reactive forms of molecular oxygen known as Reactive Oxygen Species (ROS), like singlet oxygen (10²), superoxide radical (O²⁻), hydrogen peroxide (H₂O₂) and hydroxyl radical (OH•) (MITTLER, 2002; GILL; TUTEJA, 2010). The ROS accumulation can lead to lipid peroxidation causing intense damage to the plant attributed to lipid-derived radicals that further aggravate oxidative stress (GILL; TUTEJA, 2010). A consequence of water stress is also related to changes in secondary metabolites, which can be verified in the concentration of some essential oil compounds (RIOBA et al., 2015) or in monoterpenes concentration in medicinal plants under moderate drought conditions (NOWAK et al., 2010). The ROS scavenging pathways may be enzymatic, including the enzymes superoxide dismutase, ascorbate peroxidase, guaiacol peroxidase, glutathione S-transferase and catalase, or non-enzymatic, via ascorbate, a-tocopherol, carotenoids, flavonoids and proline (GILL; TUTEJA, 2010).

Worldwide, including Brazil, the reduction in water availability has been worsened by climate change. Research has shown a tendency to increase temperatures and arid areas with water scarcity due to a decrease in the frequency of precipitation and prolonged droughts (AMBRIZZI et al., 2012). After 1990s there is an increasing in the minimal temperature allied with rate of warming more evident (AHMED et al., 2020). There is a relationship between the reduction of rainfall and the relative humidity with increasing temperatures for Brazilian conditions (SANCHEZ et al., 2017). The negative effects of the climatic situation, generated by anthropogenic actions that interact with natural processes in the environment, will cause changes in precipitation patterns that can provoke floods and droughts. Therefore, it is important to study the mechanisms by which plants develop to tolerate water restriction. A plant that is usually found in harsh conditions is the Baccharis trimera, popularly known in Brazil as "carqueja". This plant has high rustic growth habits, found in poor and acidic soils, preferring full sun conditions to grow (BONA, 2002).

B. trimera has been studied for its numerous medicinal properties, being commonly used for hepatic and gastric disorders treatment due to its antioxidant, anti-inflammatory and antimicrobial activity (RABELO; COSTA, 2018). In addition, it has applications in plant pathogenic bacteria control and phytoalexins induction in sorghum (MOURA et al., 2014) as a tool to reduce pesticides use. Thus, its uses in medicine and

sustainable agriculture are in the process of being elucidated. However, the physiological and biochemical mechanisms by which B. trimera tolerates harsh environments still need to be unraveled. These studies are important in the sense of providing tools for breeding programs for cultivated plants tolerant to different types of stresses, and for seek strategies to species preservation (SOUZA, 2015). Besides that, it is known that the global changes tend to prolonged periods of drought, which makes it important to verify if this fact affects the quality of B. trimera and its benefits for use in medicine and agriculture, so that, if applicable, they must be appropriate adaptation strategies (BASU; SHAW, 2013). Elucidation of B. trimera biochemical and physiological responses can guarantee the adequacy of a better system, pursuing a higher active ingredients final yield by area and can providing tools for breeding programs for cultivated plants tolerant to hydric stress. Thus, physiological and biochemical changes in B. trimera plants submitted to different levels of water restriction were evaluated.

MATERIALS AND METHODS

This study was performed to analyze the responses of B. trimera to water restriction, which can be important for understanding, analyzing, and improving the defense strategies through various parameters. B. trimera sample were collected along BR 153 (Erechim, Brazil), at 27°37'41.07"S, 52°14'10.35"W geographical coordinates, according to the World Geodetic System (WGS-84). Phylogenetic classification of plant samples at species level was determined according to The Angiosperm Phylogeny Group system. Plant propagation occurred by 20 cm cuttings in pots containing Plantmax® substrate arranged in a greenhouse. The indole-3-butyric acid (1g L⁻¹) was used as a rooting hormone by immersing the cuttings in solution for thirty seconds before being inserted into pots containing substrate. Irrigation occurred daily. During the period of execution of the tests, the total precipitation was 480 mm, the average temperature was 19.26 °C and average relative humidity was 74.49%. After thirty days the cuttings with root development were then transplanted into new pots, two plants per pot, with 10 cm spacing between.

After acclimation period, plants were exposed to different levels of water deficit for thirty days in a greenhouse, to assess responses during the plant's growing season. Substrate waterhold capacity was determined in laboratory with using a precision scale and considering the water saturated soil mass that ascended by capillarity as Pot Capacity (PC) excluding the pot mass. From the total water mass retained by the substrate in PC, 25, 50 and 75% were removed to determine the specific mass that the pots of each treatment should present. In order to know evapotranspiration water amount, pots without plants were weighed at two-day intervals. Water lost by evapotranspiration in the period was replaced by manual irrigation, respecting the water omission in each treatment (25, 50 and 75% water deficit) and PC at 0% water deficit treatment (MAROSTICA et al., 2019).

B. trimera growth and biomass development were evaluated thirty days after treatment beginning. Growth (GW) was determined by measuring root and shoot systems with a

ruler. After harvesting, plants were immediately weighed on a precision scale in order to obtain the fresh biomass (FB). To determine the dry biomass (DB), plants were left at 70 °C until no mass variation.

Prior to biochemical analysis, shoot and root systems were macerated separately with liquid nitrogen until they become a homogeneous powder and then stored at -80 °C. After that, sample preparation occurred according to Zhu et al. (2004) followed by centrifugation at 14,000 g and 4 °C for 20 min. Superoxide dismutase (SOD) activity was measured according to Giannopolitis and Ries (1997) adopting a lighting time of 15 min. Guaiacol Peroxidase (GP) and Ascorbate Peroxidase (APX) activity were used to determine the antioxidant activity of hydrogen peroxide removal. Techniques were performed according to Zeraik et al., (2008) and Zhu et al., (2004), respectively. The Hydrogen peroxide (H_2O_2) content in B. trimera was determined according to Loreto and Velikova (2001). The levels of lipid peroxidation products were estimated by the method described by El-Moshaty et al. (1993) measuring Malondialdehyde (MDA) concentration as the final product of lipid peroxidation by reaction with Thiobarbituric Acid (TBA). Protein concentration in B. trimera was measured by the Comassie Blue method according to Bradford (1976) using serum albumin as standard. In shoot samples, proline was measured by rapid colorimetric procedure described by Bates et

al. (1973), based in the reaction between ninhydrin and amino acids.

The experiment was done as completely randomized design. Consisting of 3 replicates and 10 cuttings per replicate, in total 120 observations per water restriction (WR) assessed. Normality of residues were verified by Shapiro-Wilk test and homogeneity of variances by Barlett's test ($p \le 0.05$). Nonnormal data, GP and APX, were transformed by the square root. Once assumptions confirmed, data was submitted to analysis of variance with F test. Water restriction and systems (root and shoot) and the interaction between them were tested. Significant results equal or lower than 5% probability level, were submitted to regression analysis with linear and quadratic models, the maximum and minimum point of curve were obtained by formula: $y = -b/2^*c$. Pearson's correlation coefficient was calculated ($p \le 0.05$). The analyses were performed at the statistical software RStudio (R CORE TEAM, 2020).

RESULTS AND DISCUSSION

We found a significant interaction between water restriction (WR) and the system (S) evaluated for absolute growth, SOD, APX and TBARS. Dry biomass show significant effect for water restriction. GP and fresh biomass showed significant difference to both, WR and the S. Protein and H_2O_2 just show difference in system evaluated. To proline contents, we verified significant effect of water restriction (Table 1).

Table 1. Summary of the Analysis of Variance for the dry biomass (DB), fresh biomass (FB), growth (GW), superoxide dismutase (SOD), guaiacol peroxidase (GP), ascorbate peroxidase (APX), thiobarbituric acid-reactive substances (TBARS), hydrogen peroxide (H_2O_2) , protein (PTN) and proline (PRL) in root and shoot systems of *Baccharis trimera* submitted of water restriction

SV	<i>p</i> -value										
	DB	FB	GW	SOD	GP	APX	TBARS	H_2O_2	PTN	PRL	
Water											
restriction	0.0000^*	0.0002^{*}	0.0001^{*}	5.9E-07*	0.0011^{*}	0.1162 ^{ns}	0.1942 ^{ns}	0.1192 ^{ns}	0.2828^{ns}	3E-05*	
(WR)											
System	0 0546ns	0.0078^{*}	0.0000^{*}	7E-07*	0.0000^*	0.0000^*	0.0000^{*}	0.0000^{*}	0.0000^*		
(S)	0.0340									-	
Interaction	0 2806ns	0 748ns	0.0074*	3 3E 06*	0 2780 ^{ns}	0.0006*	0.0020*	0.0067 ^{ns}	0.0886ns		
(WR*S)	0.2800	0.748	0.0074	5.5E-00	0.2780	0.0000	0.0029	0.0907	0.0880	-	
Linear	0.0019^{*}	0.0063^{*}	r0.023*	r0.000*	0.9322 ^{ns}	0.0001*	0.0006*			0.0024*	
regression			s0.059 ^{ns}	s0.708 ^{ns}		0.0001	0.0000	-	-	0.0024	
Quadratic	0.0032^{*}	0.0068^*	r0.001*	r0.001*	0.0009^{*}	0 5257ns	0 0800ns			0 672 1 ns	
regression			s0.692 ^{ns}	$s0.004^*$		0.3237	0.0899	-	-	0.0731	
CV (%)	30.18	32.64	8.59	14.96	11.87	9.74	18.79	13.01	23.36	9.72	

SV: source of variation; CV: coefficient of variance; ^{ns}: not significant; *: significant at 5% probability by F-test; r: root, s: shoot.

We found that water restriction of 27.37%, on average, induces an increase in physiological parameters. The dry biomass of *B. trimera* plants showed a significant statistical difference response to water deficit, but not to the system evaluated. The maximum point of DB was in 26.37% of water restriction. Shoot dry biomass development had been 1.9 times higher when compared to no water restriction (Figure 1A). The maximum point of FB (Figure 1B) was in 25.78% WR and, of root growth (Figure 1C) was in 30% WR. Therefore, these parameters show a tendency to increase with a little WR, followed by a decrease due to more restrictive conditions. Others results confirm higher

shoot development when coccoa varieties ICS-9, and MA-15 were exposed to water deficit (SANTOS et al., 2014).

Data of the present study showed that in 30% WR the root length development was 1.3 times higher than 0% WR (Figure 1C). Physiological data obtained in this study are in accordance with results reported by Price et al. (2002), observing root growth increase when *Hedyosmum brasiliense* were submitted to hydric deficiency. The observed root growth may be related to the increased exploitation of soil areas in search for water considering that the availability is higher in deeper horizons (PINHEIRO et al., 2005). However, the root growth of *B. trimera* is impaired in drought conditions above 50% of WR. **Figure 1.** Physiological parameters measured by dry biomass $(y=3.971+0.063x-0.0012x^2; R^2=0.341)$ (A) fresh biomass. $(y=8.614+0.139x-0.0027x^2; R^2=0.538)$ (B) and growth (ROOT $y=12.286+0.144x-0.0024x^2; R^2=0.882;$ SHOOT y= not significant) (C) in root and shoot of *Baccharis trimera* plants exposed to water restriction.



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There are interaction between WR and system evaluated for SOD activity. Without restriction the enzymatic activity are higher in the roots (intercept for root = 4508.04 and shoot = 1925.18). The increase of WR caused decrease SOD activity in the roots and increased activity in the shoot. Under the conditions of highest WR tested, an inversion occurred, and the SOD activity in shoot become greater than that of the root. The minimum point of root SOD activity was in 70.26% WR and of shoot SOD activity was in 36.24% WR (Figure 2A). SOD constitutes the first line in plants defense being found in all cell compartments and activated under stress conditions (MITTLER, 2002). Generally, SOD activity increases when plants are exposed to water deficit (JALEEL et al., 2008; HAMEED et al., 2011; HOJATI et al., 2011). As it was verified in shoot of *B. trimera*.

Results that corroborate the analyses in *B. trimera* root was reported by Campos et al. (2011), according to the authors SOD and POD activity in Swingle Citrumelo was reduced during water deficit. This reduction of SOD activity may be a consequence of increase in proline contents, since pyrroline existent in proline molecules presents low capacity to provide electrons thus forming a charge transfer complex capable of sequestering free O_2 and reduce superoxide ions production (REDDY et al., 2004). This may represent that *B. trimera* can tolerate a little water restriction, associated with proline activity and SOD activity balance. Even of drought stress effects, the application of proline can reduce and mitigate these effects, keeping productivity closer to normal water condition (GHAFFARI et al., 2019).

GP activity analyses in root and shoot systems from B. trimera shows a reduction in activity when exposed to milder WR, followed by an increase in activity for shoot in the most restrictive conditions (Figure 2B). The minimum point of GP was in 41.75% WR. The GP enzyme is related to lignin biosynthesis, and in the defense against biotic stress with H₂O₂ consumption (GILL; TUTEJA, 2010). The POD and APX enzyme are a ROS scavenging pathway, ROS fine tuning for signaling using H₂O₂ micro concentrations is the APX main biochemical role (MITTLER, 2002). APX showed no significant difference in shoot system, but for APX activity in root system it was found an increasing tendency when water availability was reduced (Figure 2C). A negative correlations (-0.776 and -0.877) in the root system was obtained with hydrogen peroxide and SOD activity, respectively (Table 2). The increase in H_2O_2 has a positive correlation with the increase in SOD activity on roots (0.642), as it is a product of the activity of this enzyme (MITTLER, 2002). This is suggesting that SOD activity and hydrogen peroxide concentration in root was decreasing as APX activity increased.

Thiobarbituric Acid-Reactive Substances (TBARS) had interaction between WR and system evaluated. There was an increase in lipid peroxidation in the root and a decrease in the shoot, resulting from increased levels of WR (Figure 3A). The drought resistance can be measured by lipid peroxidation, through the content of the MDA that represents the degree of damage to the cell membrane (WANG et al., 2011). It's well known that lipid peroxidation is considered one of the worst plants damages and occurs when there is ROS accumulation (GILL; TUTEJA 2010). Petridis et al. (2012) reported in a study with olives that cultivars with greater drought resistance are those whose lipid peroxidation was lower. Biochemical parameters analyses between cultivated and wild plants allowed Ghorecha et al. (2014) to conclude that wild plants tolerance is related to lower MDA accumulation, and induction of SOD activation. The best yield in wheat plants according to Hameed et al. (2011) is related to ROS detoxification systems and decreased lipid peroxidation. There are a positive relation between TBARS and APX on root (0.776). But has a negative correlation between TBARS and SOD activity on root (-0.619). That can indicate APX activity on root in higher WT as being a more relevant pathway (Table 2).





Table 2. Significant Pearson's correlation coefficients for the dry biomass (DB), fresh biomass (FB), growth (GW), superoxide dismutase (SOD), guaiacol peroxidase (GP), ascorbate peroxidase (APX), thiobarbituric acid-reactive substances (TBARS), hydrogen peroxide (H_2O_2), protein (PTN) and proline (PRL) in root (r) and shoot (s) systems of *Baccharis trimera* submitted of water restriction

Respons	TBARSr	HaOas	H2O2r	PRLs	PTNr	SODs	SODr	GPr	GWs	GWr	FBs	FBr	DBs	DBr
e	IDARSI	112023	112021	I IXLS	1 1111	50D3	SODI	UII	0.03	GWI	1 D3	I DI	003	DDI
APXs	-	0.782	-	0.784	-	-	0.653	-	0.700	-	0.713	0.641	0.588	-
APXr	0.776	-	-0.776	-	-0.862	-	-0.877	-	-	-	-	-	-	-
TBARSs	-	-	-	0.58	-	-	-	-	-	-	-	-	-	-
TBARSr	-	-	-	-	-0.681	-	-0.619	-	-	-0.668	-0.577	-0.624	-0.672	-0.599
H_2O_2r	-	-	-	-	0.828	-	0.642	-	-	-	-	-	-	-
PRLs	-	-	-	-	-	-	-	-	0.819	-	0.754	-	0.782	0.662
PTNs	-	-	-	-	-	-0.618	-	-	-	-	-	-	-	-
PTNr	-	-	-	-	-	-	0.831	0.593	-	-	-	-	-	-
SODs	-	-	-	-	-	-	-	-	-	-0.611	-	-	-	-
GWs	-	-	-	-	-	-	-	-	-	-	0.833	0.652	0.831	-
GWr	-	-	-	-	-	-	-	-	-	-	0.676	0.853	0.628	0.780
FBs	-	-	-	-	-	-	-	-	-	-	-	0.943	0.728	0.643
FBr	-	-	-	-	-	-	-	-	-	-	-	-	0.685	0.733
DBs	-	-	-	-	-	-	-	-	-	-	-	-	-	0.883

The content of hydrogen peroxide (H_2O_2) and protein concentration were different between system evaluated, bigger in the shoot (Figure 3B and 3C). There is a positive correlation between both when evaluated at the root (0.828). Peroxide is an endogenous molecule involved in plants stress signaling. Increases in H_2O_2 concentration leads to a higher potential of hydroxyl radical (OH•) production resulting in lipid peroxidation (NEILL et al., 2002). When evaluating the water deficit effect on *Catharanthus roseus* metabolism, Jaleel et al. (2008) reported a reduction in H_2O_2 content, relating it to the effective activity of both the enzymatic and non-enzymatic antioxidant system. As well as *B. trimera* analyses, drought-resistant sugarcane cultivars studied by Boaretto et al. (2014) showed that with 70% water deficit there were no differences in H_2O_2 content, only a

difference occurring for susceptible cultivar and under severe stress. In *Jatropha curcas* (L.) plants, Moura et al. (2016) reported that less significant losses caused by water deficit are associated with a lower reduction in protein content. The maintenance of these variables may indicate the ability of *B*. *trimera* to tolerate the stress caused by WR.

Figure 3. Levels of lipid peroxidation products measured by thiobarbituric acid-reactive substances (ROOT y=7.183+0.051; R²=0.821, SHOOT y= not significant) (A), hydrogen peroxide contents (y= not significant) (B) and concentration of protein (y= not significant) (C) in root and shoot of *Baccharis trimera* plants exposed to water restriction



Water restriction levels significantly influence proline content in shoot. Condition of low water restriction maintains the content of this amino acid. We found a slight increase in response to few WR. In the conditions of higher WR, proline content tends to drop. The minimum point was observed in 66.5% of WR (Figure 4). Proline is one of the amino acids most commonly associated with responses to water stress (CHEN; KAO, 1993). Proline synthesis has been related to a mechanism for cytoplasmic acidosis relief and maintenance of NADPH at compatible values with metabolism (GILL; TUTEJA, 2010). In addition, free proline acts as an osmoprotectant, protein stabilizer, metal chelator, lipid peroxidation inhibitor, and OH• and 10² eliminator (VALLIYODAN; NGUYEN, 2006). Corroborating our study, citrumelo swingle transgenic plants containing a gene encoding the key enzyme for proline biosynthesis showed 2.5 times higher proline content under moderate water stress (CAMPOS et al., 2011). Primary nitrogen metabolism modulation by water deficit through dependent and independent abscisic acid (ABA), in Medicago truncatula demonstrated that water deficit induced ABA accumulation, consequently inducing asparagine and proline accumulation, such osmolite increase contributed to the osmotic adjustment (PLANCHET et al., 2011).

In *B. trimera* the higher proline content (13.21 μ mol of proline g⁻¹ of plant tissue) obtained under 25% water restriction may have led to osmotic adjustment, which was possibly responsible for the 1.3 times increase in root and shoot length, and 1.9 times in fresh biomass. Under conditions of abiotic stress, the transcription of genes related to proline content is increased in tolerant genotypes (BENITEZ et al., 2016). This demonstrates that the overcoming abiotic stress mediated by proline is an interesting route for breeding programs. Proline has a positive correlation with APX and TBARS on shoot (0.784 and 0.580 respectively), and a positive correlation with GW (0.819) and FB (0.754) shoot and DB root (0.782) and shoot (0.662). It can

indicated proline as being relevant for maintaining high physiological parameters in WR conditions up to 30% (Table 2).

Figure 4. Proline content in *Baccharis trimera* plants exposed to water restriction ($y=10.491-0.027x-0.0002x^2$; R²=0.157).



In *B. trimera* low water restriction, 27.37% on average, increase growth, dry biomass and fresh biomass, limited until drought conditions above 50% of WR. *B. trimera* can tolerate a little water restriction, related with proline activity associated, in milder conditions, with SOD activity in the roots. In conditions of higher restriction can be related with APX in the root and SOD in the shoot. Therefore, *B. trimera* maintains root growth and root and shoot biomass under low WR levels. Proline, and APX and SOD activity can be a pathway that scavenging the stress generated by WR and maintains biomass and growth (Figure 5).

Figure 5. Summary of physiological responses measured by dry biomass (DB), fresh biomass (FB) and growth (GW) and biochemical responses measured by superoxide dismutase (SOD), guaiacol peroxidase (GP), ascorbate peroxidase (APX), thiobarbituric acid-reactive substances (TBARS), hydrogen peroxide (H₂O₂), protein (PTN) and proline (PRL) in root and shoot systems of *Baccharis trimera* submitted of water restriction.



CONCLUSION

B. trimera maintains root growth and root and shoot biomass under low WR levels. The proline and the APX and SOD activity can be a pathway that scavenging the stress generated by WR on *B. trimera*. This pathways can be an important tool to be considered in breeding programs to enhance drought tolerance in plants and for seek strategies to medicinal species cultivation in harsh conditions. In addition, this study provides a way to optimize the sustainable production of these species, in view of a greater biomass production.

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