



Exogenous Melatonin Protects Lime Plants from Drought Stress-Induced Damage by Maintaining Cell Membrane Structure, Detoxifying ROS and Regulating Antioxidant Systems

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Abstract: Lime is an important commercial product in tropical and subtropical regions, where drought stress is becoming one of the most severe environmental challenges in the agricultural sector. Melatonin is an antioxidant molecule that helps plants regulate their development and respond to a variety of stresses. In this research, the effects of exogenous melatonin treatments were evaluated at different concentrations (0, 50, 100, and 150 µM) on biochemical aspects and gene expression in two species of lime plants ("Mexican lime" and "Persian lime") under normal (100% field capacity (FC)) and drought stress conditions (75% and 40% FC). The experiments were factorial and based on a completely randomized design (CRD) with four replicates. Drought stress caused electrolyte leakage (EL) as well as accumulations of hydrogen peroxide (H2O2) and malondialdehyde (MDA), indicating the occurrence of damage to cellular membranes. In contrast, the melatonin pretreatment at various concentrations reduced the levels of EL, H₂O₂ and MDA while mitigating the negative effects of drought stress on the two lime species. The application of melatonin (100-µM) significantly increased the level of proline content and activity of antioxidant enzymes in plants under drought stress compared to control plants. According to real-time PCR analysis, drought stress and melatonin treatment enhanced the expression of genes involved in ROS scavenging, proline biosynthesis, and cell redox regulation in both species, as compared to their respective controls. According to these findings, melatonin is able to detoxify ROS and regulate antioxidant systems, thereby protecting lime plants from drought stress-induced damages.

Keywords: antioxidant; Citrus; gene expression; hydrogen peroxide; malondialdehyde; melatonin

1. Introduction

Limes are the smallest citrus fruits in terms of size, and are one of the most important commercial products in the tropics and subtropics. In these regions, drought is a major environmental problem that can cause a substantial decrease in their production [1]. There are two types of acidic lime species: "Persian lime" (*Citrus latifolia* Tanaka) and "Mexican lime" (*Citrus aurantifolia* (Christm) Swingle) [2,3]. While drought acts as a severe environmental restriction and prevents plants from growing or developing properly, it disrupts the morphological, physiological, and anatomical structures of plants in a variety of ways [4–7]. Drought stress generates an overabundance of reactive oxygen species (ROS) in plants and instigates membrane lipid peroxidation as well as electron leakage, thereby disrupting cell membrane structures [8,9]. Plants collect significant amounts of osmolytes that are



Citation: Jafari, M.; Shahsavar, A.R.; Talebi, M.; Hesami, M. Exogenous Melatonin Protects Lime Plants from Drought Stress-Induced Damage by Maintaining Cell Membrane Structure, Detoxifying ROS and Regulating Antioxidant Systems. *Horticulturae* 2022, *8*, 257. https://doi.org/10.3390/ horticulturae8030257

Academic Editors: Pascual García-Pérez and Maria Angeles Pedreño

Received: 13 February 2022 Accepted: 16 March 2022 Published: 17 March 2022

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characterized by low molecular weights and are highly soluble, non-toxic compounds, thereby maintaining plant water content under drought stress [10].

Proline is the most common plant-compatible osmolyte that can reduce membrane permeability and assist plants in maintaining water balance when they are under drought stress. In the production cycle of proline, an increase in proline accumulation in the cell can occur by the stimulation of enzyme production, especially delta 1 proline 5 carboxylate synthetase (P5CS) and proline 5 carboxylate reductase (P5CR) [11]. Protective mechanisms under osmotic stress involve direct ROS scavenging, balancing intracellular redox homeostasis, and cellular signaling enhancement [7]. In this regard, several plants have been studied, including arabidopsis (*Arabidopsis thaliana*) [12], walnut (*Juglans regia*) [13], pistachio (*Pistacia vera*) [14], three grapevine cultivars (*Vitis vinifera* L.) [15], and two olive cultivars (*Olea europaea* L., Arbequina and Empeltre cvs.) [16].

Plant metabolism may be severely harmed ROS, and these can cause permanent damage to vital macromolecules [17-19]. Hydrogen peroxide (H₂O₂), as the most important ROS, is involved in a number of activities related to plant growth. The production of ROS is considered a hazard to cellular metabolism since it generally results in electron leakage, speeds up cell membrane liposuction and generates the harmful chemical malondialdehyde (MDA), thereby making membranes more permeable and compromising their structural integrity [20]. Plants, on the other hand, have evolved a variety of strategies to combat the detrimental consequences of drought and to protect themselves from the harmful effects of high ROS levels. The antioxidant defense system, which is made up of both enzymatic and non-enzymatic components, may effectively scavenge ROS. Superoxide dismutase (SOD), ascorbate peroxidase (APX), peroxidase (POD), and catalase (CAT) are enzyme antioxidants found in plants. A positive correlation between antioxidant enzyme activity, production of osmolytes, and their gene expression has been reported in previous studies [21,22]. The depth-insights to abiotic stress tolerance in plants can be improved by studying the regulation of gene expression. It is plausible that over-expression of genes related to antioxidant enzymes, cellular redox regulation, and proline synthesis may enhance abiotic stress tolerance in plants. For instance, the increased SOD activity helps plants to show resistance to salinity and drought stresses in Brassica juncea plants [23]. Moreover, abscisic acid and melatonin application modulated the expression profile of the Cu/Zn superoxide dismutase (Cu/Zn SOD) gene in cotton (Gossypium hirsutum L.) plants under drought stress [24]. Additionally, the overexpression of Alternative oxidase (AOX) enhanced Arabidopsis thaliana tolerance to drought and salinity stresses [25]. Nonetheless, chronic high-stress exposure can result in severe damage and cell death [26,27]. As a result, several cases of research have aimed to increase crop resistance and reduce the destructive effects of various stressors, including drought stress. While several methods have been used for achieving this goal, plant bio-stimulators are one of the new techniques for improving plant adaptation and protection against unfavorable environmental conditions [28–31]. Findings in recent years have clearly suggested that melatonin may be a very effective substance for reducing stressors, particularly the type of stress that results from water deficits [32,33].

Melatonin reduces oxidative damage by directly scavenging ROS or by modulating the activity and production of enzymes and non-enzyme antioxidants [34]. Melatonin is reportedly able to reduce oxidative damage caused by water deficit in alfalfa (*Medicago sativa* L.) [35], apple [36], soybean plants [37,38], tobacco seedlings [29], and maize (*Zea mays* L.) [33]. Several cases of research have demonstrated that phytomelatonin can exert effects by modulating various components of the redox network or by interfering with the activity of other phytohormones even as they denote structural similarities between melatonin and indole-3-acetic acid (IAA) [39,40]. In response to its amphiphilic nature, melatonin may readily cross cellular membranes with ease, unlike other antioxidants. Several anti-stress-related genes are up-regulated in plants treated with exogenous melatonin [41]. Melatonin functions as a potent antioxidant agent while depending on its direct ability to scavenge ROS or to stimulate the activity of antioxidant enzymes. It also acts

through its ability to produce a highly effective cascade of free radical scavengers among its metabolites, including N-acetyl-N-formyl 5-methoxyknuramine (AFMK) and N-acetyl-5-methoxykynuramine (AMK). Accordingly, melatonin is usually regarded as one of the most powerful antioxidants in the world when compared to other known chemicals [42].

Despite the fact that previous research has extensively focused on the impacts of this multifunctional molecule on plants, especially when plants are under abiotic stress, more research is needed to update and uncover the functions of melatonin. In fact, a complete image of melatonin-mediated drought tolerance has yet to be pictured. Melatonin acts as a protector against stress by regulating various elements of the redox network so that the expression of many stressor genes can be upheld. The current study aims at the protective functions of melatonin-induced drought-stress tolerance, including biochemical and molecular alterations, and its essential role in regulating the antioxidant system of the two lime species. The results of the current study can help growers to tackle the negative effects of drought stress on lime species through foliar application of melatonin.

2. Materials and Methods

2.1. Plant Material, Experimental Design and Treatments

One-year-old "Persian lime" (Citrus latifolia Tanaka) and "Mexican lime" (Citrus aurantifolia (Christm) Swingle) seedlings were considered as samples. All plants were grown in plastic pots (5 kg each, 33 cm in diameter and 36 cm in height). The pots were filled with soil and leaf litter (3:2 w/w). The field experiment was carried out in a greenhouse at Shiraz University, College of Agriculture in Shiraz, Iran, in September 2019. The plants were placed in a chamber with a mean temperature of 25 ± 2 °C, a relative humidity of 80%, and a day/night cycle of 14/10 h. Before initiating the experiments, all seedlings were thoroughly watered every day. A supplement of 1/2 Hoagland's solution (pH 7.0) was given once a week. Healthy and uniform plants were selected for three watering regimes after three months of growth. The three watering regimes were, namely, (i) well-watered (100% field capacity) (FC), (ii) mild drought-stress (75% FC) and (iii) severe drought-stress (40% FC). Evaporative water loss was assessed by weighing each pot and measuring weight-related variations between each watering episode. Ethanol was used for dissolving melatonin (Sigma-Aldrich Chemie, Steinheim, Germany) and preparing different concentrations of melatonin. Additionally, Tween-20 (0.1%) as a surfactant was applied for the foliar application of melatonin. Exogenous melatonin was given to all plants in the well-watered and drought treatments, at concentrations of 50, 100, and 150 μ M. As a result, four experimental groups were investigated: (i) non-melatonin treatment with well-watered conditions (i.e., well-watered control), (ii) non-melatonin treatment with drought treatments (i.e., moderate and severe stress), (iii) 50, 100, and 150 μ M melatonin treatments with well-watered conditions, and (iv) 50, 100, and 150 µM melatonin treatments with drought treatments. The treatments were stopped after two months in the greenhouse. Fully young expanded leaves, from the middle part of the plant, in each treatment were sampled on day 60 for electrolyte leakage index, lipid peroxidation, proline content, antioxidant enzyme activity assessments, and RNA extraction for carrying out a relative gene expression analysis.

2.2. Estimation of Proline Content

According to the method developed by Bates et al. [43], proline content was determined in freshly collected leaves. Accordingly, 10 mL of sulfosalicylic acid (3%) (w/v) (Merck KGaA, Darmstadt, Germany) was used for homogenizing 0.5 g of fresh leaves. The extract was filtered and combined with an equal ratio of glacial acetic acid and acidninhydrin reagent (2 mL) (Merck KGaA, Darmstadt, Germany). The samples were incubated in boiling water for 40 min, and the reactions were then halted by an ice bath. The solution was thoroughly mixed after adding 4 mL of toluene. On a spectrophotometer (Jenway-7315, Staffordshire, UK), the light absorbance of the toluene phase was measured at 520 nm, and the proline content was evaluated using a standard proline curve. The concentration of proline was estimated in micromoles per gram of fresh weight (μ mol g⁻¹ FW).

2.3. Measurement of Electrolyte Leakage (EL)

The electrolyte leakage was measured using the Gulen and Eris [44] method. Leaf samples were sliced into one-centimeter pieces. The samples were put in test tubes and incubated in the dark for 24 h with 45 mL of distilled water (23 °C). The test tubes were stored at 25 °C. A conductivity meter was used to measure the electrical conductivity (EC₁) of the electrolytes after shaking the test tubes vigorously (Hanna, HI8633, North Smithfield, RI). After the EC₁ measurements, all samples were autoclaved for 15 min at 121 °C. Then, the samples were cooled (25 °C) and their electrical conductivity (EC₂) was measured once more. The electrolyte leakage was calculated using the following formula:

Electrolyte leakage (%) = $(EC_1/EC_2) \times 100$

2.4. Measurement of Lipid Peroxidation and Hydrogen Peroxide (H_2O_2) Concentration in Leaves

When malondialdehyde (MDA) was generated by the thiobarbituric acid reaction, its quantity was measured by evaluating lipid peroxidation. Accordingly, 100 mg of leaves were homogenized in 2 mL of trichloroacetic acid (0.1%) (TCA) solution (Merck KGaA, Darmstadt, Germany) and then centrifuged at 12,000 rpm for approximately 25 min at 4 °C (SIGMA 1-14 k, Osterode, Germany). The supernatant was then treated with 4 mL of thiobarbituric acid (0.5%) (TBA) in TCA (10%) (Merck KGaA, Darmstadt, Germany). The solution was centrifuged at 10,000 rpm for approximately 10 min (SIGMA 14 k, Osterode, Germany). The absorbance of the mixture was measured at 532 and 600 nm [45]. The concentration of MDA was determined using a correction factor of 0.155 (mol⁻¹ cm⁻¹) and was represented in micromoles per gram of fresh weight (μ mol g⁻¹ FW).

The amount of hydrogen peroxide was calculated using the method developed by Alexieva et al. [46], which involved reacting H₂O₂ with potassium iodide. In this technique, 0.5 g of fresh leaf tissue was crushed with 0.1% trichloroacetic acid (TCA). The resultant extract was centrifuged at 12,000 rpm for 15 min. Then, 2 mL of 1 M potassium iodide were added to 500 μ L of supernatant and 500 μ L of 100 mM potassium phosphate buffer (pH 7.0). The reaction mixture was kept at room temperature (25 °C) for one hour before the samples' absorbance was measured at 390 nm.

2.5. Antioxidant Enzymes Activity

Extraction and measurement of antioxidant enzyme activity were performed using Ozden et al. [47] method. To prepare the extracts, 0.5 g of fresh leaf tissue was homogenized in 5 mL potassium phosphate buffer (50 mM) (pH 7.0) (Merck KGaA, Darmstadt, Germany) along with 2 mM ethylene diamine tetraacetic acid (EDTA) (Merck KGaA, Darmstadt, Germany) and 1% polyvinylpyrrolidone (PVP) (as extraction buffer) (Merck KGaA, Darmstadt, Germany). The resultant homogeneous mixture was centrifuged (15,000 rpm) for 15 min at 4 °C using a SIGMA 3-16PK refrigerated centrifuge (SIGMA 1-14 k, Osterode, Germany). The resultant supernatant was then stored at -84 °C until relevant experiments were performed. Using a spectrophotometer (Jenway 7315, Staffordshire, UK) to detect the increase in light absorption induced by the oxidation of guaiacol in the presence of H₂O₂ at a wavelength of 470 nm, the activity of the POD enzyme was calculated based on Hemeda and Klein [48]. The POD enzyme activity was determined in terms of oxidized μ M of guaiacol at a rate of one min per gram of fresh sample weight using a quenching coefficient of 26.6 mM cm⁻¹.

SOD activity was evaluated by measuring the decrease in light absorption of the nitroblutetrazolium chloride (NBT) complex [49]. In this method, the reaction mixture (3 mL) contained 50 μ L extracted enzyme extract, 50 mM potassium phosphate buffer (pH 7.0), 13 mM L-methionine, 75 μ M NBT, 0.1 mM EDTA, and 4 μ M riboflavin. To perform the reaction, the mixtures were situated in a light chamber, which was sourced with four 20-watt fluorescent lamps. The samples were placed under the lamps for 15 min. The reaction was then stopped by turning off the lamps and placing the samples in the dark.

The absorbance of each sample was read at a wavelength of 560 nm. The enzyme activity was reported as units per mg of fresh sample weight.

Dhindsa et al. [50]'s method was used for measuring the activity of CAT. According to this method, 50 μ L of the extract was mixed with 1 mL of catalase-measuring solution, which contained potassium phosphate buffer (50 mM) (pH 7.0) and H₂O₂ (10 mM). Then, the adsorption of the solution was measured at 240 nm for one minute with a spectrophotometer (JENWAY-7315, Staffordshire, UK). The extinction coefficient for the activity of this enzyme is usually 39.4 mM⁻¹ cm⁻¹, and an enzyme unit (CAT) equal to the decomposition of one mM of H₂O₂ per minute was considered.

To measure the activity of APX, the reaction mixture consisted of 150 μ L enzyme extract, 50 mM phosphate buffer, 1.2 mM oxygenated water, 0.5 mM ascorbic acid, and 0.1 mM EDTA. Due to ascorbic acid peroxidation, the decrease in light absorption was read at 290 nm with a spectrophotometer. The extinction coefficient for enzyme activity is usually 2.8 mM⁻¹ cm⁻¹. Absorption changes per minute, per gram of fresh sample weight, were used for calculating enzyme activity [51].

The activity of glutathione reductase (GR) was measured using the Foyer and Halliwell [52] technique in a reaction mixture containing 25 mM sodium phosphate buffer (pH 7.0), 0.1 mM EDTA, 0.5 mM oxidized glutathione (GSSG), 0.12 mM NADPH, and 100 μ L of enzyme extract. In accordance with NADPH oxidation, an extinction coefficient of 6.2 mM⁻¹ cm⁻¹ was used for evaluating GR activity by monitoring the reduction in absorbance at 340 nm. As a result, GR activity was expressed as specific activity units per mg⁻¹ protein.

2.6. RNA Extraction, cDNA Synthesis and Quantitative Real-Time PCR

RNA was extracted according to the manufacturer's protocol using Iraizol kit (RNA Biotechnology Co., Isfahan, Iran). The total RNA concentration and its quality were measured using a Nano-Drop (ND) 1000 spectrophotometer (Implen Nano Photometer, NPOS 3.0 version 12984, Westlake Village, CA, USA). Agarose gel electrophoresis (1.2% w/v) was also used for evaluating total RNA integrity. DNaseI (Promega, Madison, WI, USA) was employed to eliminate genomic DNA before cDNA synthesis. Following the manufacturer's instructions, cDNA was synthetized from 1 μ g of total RNA using the cDNA synthesis kit (RNA Biotechnology Co., Isfahan, Iran). Primers for genes were involved in ROS scavenging (cytosolic ascorbate peroxidase, cAPX; Cu/Zn superoxide dismutase, *Cu/Zn SOD*; Fe superoxide dismutase, *Fe SOD*), cellular redox regulation (alternative oxidase, AOX; NADH dehydrogenase, NaDde), and proline synthesis (Pyrroline-5-carboxylate synthase, *P5CS*; pyrroline-5-carboxylate reductase, *P5CR*). The sequences of the primers (Table 1) were designed based on the conserved sequences in Rutaceae found in the NCBI GenBank database. The ABI Step One (Applied Biosystems, Waltham, MA, USA) was used for running the real-time PCR test in a 10 μ L reaction mixture. Each reaction included 5 μ L SYBR Green (2X RealQ Plus Master Mix Green, AMPLIQON, Odense, Denmark), 0.1 µL of each primer (10 μ M), and 1 μ L of five times diluted cDNA template, and deionized double distilled water up to 10 μ L.

The amplification conditions had several steps, including initial denaturation at 95°C for 10 min, followed by 40 cycles of 95 °C for 15 s, appropriate annealing temperature (Table 1) for 30 s, and finally, 72 °C for 45 s. To confirm the purity of the amplified products, a melting curve was generated for each sample when each cycle ended. The $2^{-\Delta\Delta Ct}$ technique was used for calculating relative gene expression levels [53] using the Actin gene (ACT) as an internal reference gene. The experiment on gene expression was conducted with three replicates in both technical and biological samples.

Row	Primer Name	Primer Sequence	Length (bp)	Annealing Temperature (°C)	
1	AOX_F	5'- GCGTAAGTTCCAGCATAGTG -3'	20	60	
	AOX_R	5'- CCTCCAAGTAGCCAACAAC -3'	19		
2	Sod-cu/zn-F	5'- TGATGACGGGAACTAACGGT -3'	19	60	
	Sod-cu/zn-R	5'- AGTGTGAATAATGAGTGCGTGA -3'	22		
3	Sod-fe_F	5'- CGTAAGGAGCGGCGAGTA -3'	18	55	
	Sod-fe_R	5'- GTGGCTAATGCGGTGAAT -3'	18		
4	APX_F	5'- AGCAGTTCCCTACCATCTCC -3'	20	58	
	APX_R	5'- TTCAGCCTTGTCATCTCTTCC -3'	21		
5	P5CS_F	5'- TGGACTAGGTGCTGAGGTTG -3'	20	55	
	P5CS_R	5'- ACCCGCTTCCTTTGAGAATC -3'	20		
6	P5CR-F	5'- TCTGCTGTAGGTGAGGCTGC -	20	58	
	P5CR-R	5'- ATATGCTGGACCGCTGCCAC -3'	20		
7	NADH-F	5'- CTTCATGCCCAAGGTGTCTGAT -3'	22	60	
	NADH-R	5'- ATCAAGCAGCCCTCCAACAA -3'	20		
8	ACT-F	5'- CCAGGCTGTTCAGTCTCTGTAT -3'	22	55	
	ACT-R	5'- CGCTCGGTAAGGATCTTCATCA -3'	22		

Table 1. Primer sequences used for Real Time PCR analysis in "Mexican lime" (*Citrus aurantifolia* (*Christm*) Swingle) and "Persian lime" (*Citrus latifolia* Tanaka).

2.7. Statistical Analysis

The experiment was performed as a CRD in a factorial arrangement with three factors, including four concentrations of melatonin, three levels of drought stress, and two *Citrus* species. The experiment had 24 treatments with four replications (96 experimental units). The SAS software (SAS Institute, Cary, NC, USA, V.9.1) was used for carrying out the analysis of variance (ANOVA). The LSD test was used to compare mean values at a 5% probability level ($p \le 0.05$).

3. Results

3.1. Proline Content

Based on the results of comparing the mean effects of treatments, drought stress increased proline concentration in leaves, and a significant difference was observed between species. Comparison of means at different levels of stress showed that Mexican lime control plants with (55.5 μ mol g⁻¹ FW) had the lowest and Mexican lime and Persian lime species under severe drought stress and 100 μ M melatonin foliar application with 155.3 and 204.3 μ mol g⁻¹ FW had the highest proline concentration, respectively. Proline levels in both species of lime under drought treatment showed a significant increase compared to the control. Overall, treatments of 100 and 50 μ M melatonin significantly increased proline levels, and at the highest concentration (150 μ M) this increase was less (Table 2).

3.2. Electrolyte Leakage

Drought stress caused an increase in EL into the intercellular space. This increase was 59.4% in "Mexican lime" and 33.4% in "Persian lime" in response to 40% FC, compared to the control (100% FC), implying that drought stress impaired cellular membrane integrity and fluidity. Exogenous melatonin, on the other hand, significantly controlled the rise of EL under drought conditions and ultimately led to a decrease of approximately 20% in the EL in plants sprayed with a concentration of 100 μ M melatonin, compared to the control group under severe drought stress (Table 2). The foliar application of 100 μ M melatonin was the most efficient treatment in preserving cellular membrane integrity.

Lime Species	Melatonin (µM)	% Drought Stress	Proline (μmol g ⁻¹ FW)	%EL	H_2O_2 (µmol g ⁻¹ FW)	MDA (µmol g ⁻¹ FW)
	0	100	65.5 ⁿ	14 ^{kl}	0.5 ^j	0.16 ^k
		75	116.3 ^{gh}	29.2 ^e	3.5 ^e	0.36 ^f
		40	117.8 ^{fg}	33.4 ^d	4.5 ^d	0.39 ^e
	50	100	70.2 ^m	18 ^{hij}	0.9 ^{hij}	0.25 ⁱ
		75	179.1 ^b	23.2 ^{fg}	2.7 ^f	0.33 ^f
Persian		40	180.1 ^b	28.5 ^e	3.3 ^e	0.35 ^f
reioluli	100	100	100.5 ^j	12 ^m	0.3 ¹	0.11 ^m
		75	202.5 ^a	14.9 ^{jkl}	0.7 ^{ij}	0.18 ^k
		40	204.3 ^a	20.7 ^{gh}	1.7 ^g	0.30 ^g
		100	83.6 ^k	19.3 ^{hi}	1.1 ^{hi}	0.28 ^h
	150	75	156.2 ^c	21 ^{gh}	1.5 ^g	0.30 ^g
		40	75.6 ¹	25 ^f	2.7 ^f	0.34 ^f
	0	100	55.5 ^p	16.8 ^{jkl}	0.7 ^{ij}	0.21 ^j
		75	110.7 ⁱ	48.4 ^b	7.8 ^a	0.60 ^b
		40	112.7 ⁱ	59.4 ^a	7.8 ^a	0.64 ^a
	50	100	60.5 °	17.7 ^{hij}	0.8 ^{hij}	0.25 ⁱ
		75	125.2 ^e	34.1 ^d	5.3 ^c	0.45 ^d
Mexican		40	139.9 ^d	38.7 ^c	6.7 ^b	0.52 ^c
1110/ticult	100	100	97.8 ^j	13.1 ¹	0.5 ^k	0.14^{1}
		75	141.1 ^d	17.6 ^{hij}	0.7 ^{ij}	0.21 ^j
		40	155.3 ^c	33.6 ^d	4.7 ^d	0.40 ^e
	150	100	83.1 ^k	19.9 ^{ghi}	1.2 ^h	0.28 ^h
		75	122.3 ^{ef}	37.8 ^c	5.7 ^c	0.46 ^d
		40	72.5 ¹	46.1 ^b	6.9 ^b	0.53 ^c

Table 2. Effect of melatonin concentrations, lime species, and percentage of drought stress on biochemical responses including proline content, percentage electrolyte leakage (%EL), hydrogen peroxidase (H₂O₂) and malondialdehyde (MDA).

Means in each column followed by same letters at superscript are not significantly different according to LSD at p < 0.05.

3.3. Hydrogen Peroxide and Malondialdehyde Contents in Leaf Extracts

As important indicators of stress-induced ROS levels and oxidative damage, H_2O_2 and MDA levels were measured in the control and melatonin-pretreated plants during drought stress treatments. Melatonin had no influence on the levels of H_2O_2 or MDA in the control group. When drought stress was applied, melatonin-pretreated lime plants had significantly lower levels of H_2O_2 and MDA, compared to non-treated lime plants, indicating less oxidative damage. These findings suggest that exogenous melatonin treatment can reduce abiotic stress-induced ROS production and mitigate oxidative damage in both lime species. H_2O_2 and MDA accumulated extensively in plant leaves after the drought treatment, but this accumulation was partially mitigated by melatonin application. In plants treated with drought and melatonin, the leaf H_2O_2 concentration was lower than in control plants, although drought stress treatment alone raised the H_2O_2 content in "Mexican lime" and "Persian lime" by 4.5 and 7.8 µmol g⁻¹ FW, respectively. Similarly, drought stress alone increased the leaf MDA content, whereas applying 100 µM melatonin caused a maximum decline in MDA content, regardless of the level of drought stress (Table 2).

3.4. Changes in the Activities of Antioxidant Enzymes

Drought stress affected SOD, POD, GR, and APX levels in the leaves of both lime species, either with or without melatonin treatment. The results demonstrated that severe drought-stressed plants (particularly at 40% FC) had higher SOD, GR, and APX activity, compared to the control plants. CAT activity showed a different trend than other enzymes. The highest activity of this enzyme was observed in Persian lime species under moderate drought stress (75% FC) and foliar application with 100 μ M melatonin, which was not significantly different from plants under severe drought stress at the same concentration. The application of 100 μ M melatonin caused plants to have higher antioxidant enzyme activity than plants that had been exposed to severe drought stress (Figure 1).

Both species showed significantly enhanced activities of SOD under stress, and those activities were further increased by exposure to melatonin. Generally, the results of this experiment showed that 100 μ M melatonin under severe drought stress (40% FC) significantly enhanced the SOD activities in both Persian lime (78.8 units g^{-1} FW) and Mexican lime (73.5 units g^{-1} FW). Additionally, in moderate drought stress (75% FC), a similar trend was observed and melatonin foliar application showed a positive effect on increasing the activity of this enzyme. The lowest activities of SOD were observed in both species under unstressed condition without melatonin foliar application. According to these findings, there was a significant difference in POD activity caused by the different melatonin concentrations under normal and drought-stress conditions. The 100 μ M melatonin treatment led to the increase in POD activity in "Mexican lime" (10.6 μ mol guaicol min⁻¹ g⁻¹ FW) and "Persian lime" (11.9 μ mol guaicol min⁻¹ g⁻¹ FW). Without using melatonin, the lowest POD activity (1.2 μ mol guaicol min⁻¹ g⁻¹ FW) was obtained in the unstressed "Mexican lime" (Figure 1b). In general, the findings of this experiment demonstrated that 100 μM melatonin substantially increased CAT activity in both "Persian lime" (9.7 µmol H₂O₂ min⁻¹ g^{-1} FW) and "Mexican lime" (4.3 µmol H₂O₂ min⁻¹ g^{-1} FW) under severe drought stress, compared to the unstressed condition. At moderate drought stress (75% FC) similar results were obtained compared to severe drought stress (40% FC) and the application of 50 and 100 µM melatonin increased CAT activity. The lowest level of CAT activity in both species was observed in control plants (Figure 1c).

The findings showed that drought treatment at 40% FC increased APX activity in Mexican lime (9.2 units g^{-1} FW) and Persian lime (8.2 units g^{-1} FW), as compared to the control. Under moderate and severe drought conditions, treating lime plants with 100 μ M melatonin increased the activity of this enzyme (Figure 1d). In drought-stressed plants, GR activity increased in a manner similar to other enzyme activities. During severe drought stress, using 100 μ M melatonin increased the activity of GR in "Persian lime" and "Mexican lime" (4.5 and 2.8 units g^{-1} FW, respectively). Compared to the non-drought controls treated with different melatonin dosages, drought-stressed plants with the same melatonin treatments showed substantial GR activity in leaf extracts (Figure 1e).



Figure 1. The effects of different concentrations of exogenous melatonin, *Citrus* species, and various levels of drought stress on the activities of antioxidant enzymes including (**a**) superoxide dismutase, (**b**) peroxidase, (**c**) catalase, (**d**) ascorbate peroxidase, and (**e**) glutathione reductase. Based on the LSD test, columns with similar letters are not significant at the 5% probability level.

3.5. Relative Gene Expression

The formation of 28srRNA and 18srRNA bands on the gel electrophoresis indicated no fracture or damage to the RNA structure. Additionally, the PCR products confirmed the length of the amplified fragments on a 1.2% agarose gel. It showed that all primers were able to amplify the expected fragments well (Figure 2).



Figure 2. Agarose gel electrophoresis shows total RNA extracted from Persian lime leaves in some treatments (1: 50 μ M melatonin under mild drought-stress; 2: 100 μ M melatonin under mild drought-stress; 3: 150 μ M melatonin under mild drought-stress; 4: 50 μ M melatonin under severe drought-stress; 5: 100 μ M melatonin under severe drought-stress; 6: 150 μ M melatonin under severe drought-stress; M: Marker III).

The results of the real-time PCR reaction in the "Mexican lime" revealed that the relative expression of *SOD Cu/Zn* and *Fe SOD* genes increased by about 19.04 and 27.412-fold in response to severe drought stress and foliar application (100 μ M melatonin), respectively. Additionally, the highest expression of these genes was observed in Persian lime species in response to mild drought stress and foliar application (100 μ M melatonin), which caused these genes to increase their expression by about 20.066 and 28.147-fold, respectively, compared to the control sample (Figure 3a,b).

The results of *APX* gene expression showed an increase in the expression of this gene in both Mexican and Persian lime species. Under severe drought stress and foliar application of 100 μ M melatonin, the gene expression of *APX* increased in each species, as compared to the corresponding control. The largest increase in the transcript level occurred in the Mexican lime (24.86-fold higher) and in the Persian lime (27.04-fold higher) (Figure 3c). In drought-stressed plants, the *P5CS* and *P5CR* genes displayed two similar patterns of expression. Plants that were exposed to drought quickly responded by increasing the *P5CR* (Figure 4a) and *P5CS* expression (Figure 4b) in both lime species. When plants under drought stress were sprayed with 100 μ M melatonin, the expression of the *P5CR* gene increased in both species. In response to severe drought stress and foliar application (100 μ M melatonin), the expression of the *P5CR* gene in "Persian lime" and "Mexican lime" became 31.07 and 28.25 times higher than in the control plants, respectively. Furthermore, the transcript level in Persian lime was higher than in Mexican lime in each treatment.



Figure 3. The effect of different concentrations of exogenous melatonin, *Citrus* species, and various levels of drought stress on the expression of ROS scavenging genes including (**a**) Cu/Zn superoxide dismutase, *Cu/Zn SOD*, (**b**) Fe superoxide dismutase, *Fe SOD*, and (**c**) cytosolic ascorbate peroxidase, *cAPX*. Data describe the means of three biological replicates with three technical replicates each. Error bars represent standard deviation (SD).



Figure 4. The effect of different concentrations of exogenous melatonin, *Citrus* species, and various levels of drought stress on the expression of proline biosynthesis genes including (**a**) pyrroline-5-carboxylate reductase, *P5CR* and (**b**) pyrroline-5-carboxylate synthase, *P5CS*. Data describe the means of three biological replicates with three technical replicates each. Error bars represent standard deviation (SD).

As shown in Figure 5a, the *AOX* gene was upregulated after drought treatments. When plants under drought stress were treated with melatonin foliar spray, *AOX* gene expression was upregulated, compared to the expression in control plants. However, the foliar application of melatonin alone had no effect on increasing the *AOX* gene expression. The highest levels of gene expression in both Persian lime and Mexican lime were observed in response to the 100 μ M melatonin treatment and severe drought stress. These expression levels were about 35.235 and 41.798-fold higher than in the control plants, respectively. While drought stress increased *NaDde* gene expression levels, pretreating the plants with melatonin also resulted in a substantial upregulation of *NaDde* gene expression. Its highest level of expression was observed in "Persian lime" in response to the 100 μ M melatonin treatment and severe drought stress. The observed in "Persian lime" in response to the 100 μ M melatonin treatment and severe drought stress to the 100 μ M melatonin treatment and severe drought stress increased *NaDde* gene expression (9.600-fold higher than in control plants. In "Mexican lime", the highest level of gene expression (9.600-fold higher than the control) was observed in response to 100 μ M melatonin and mild drought stress (Figure 5b).



Figure 5. The effect of different concentrations of exogenous melatonin, *Citrus* species, and various levels of drought stress on the expression of cellular redox regulation genes including (**a**) alternative oxidase, *AOX* and (**b**) NADH dehydrogenase, *NADHde*. Data describe the means of three biological replicates with three technical replicates each. Error bars represent standard deviation (SD).

4. Discussion

Excessive production of ROS under stress conditions damages the plant structure and disrupts the plant's natural metabolism [54,55]. Plants prevent these disorders by regulating the expression of stress-related genes, physiological and biochemical responses, and the antioxidant defense system [56,57]. Plants can be safeguarded against stress by osmoprotectants, such as soluble proteins, proline, and total free amino acids, because these assist with cellular osmotic adjustment, enzyme stability, and protein detoxification of ROS, all of which contribute to the maintenance of membrane integrity [58]. As an osmotic protectant, proline aids in the preservation of turgor pressure in stressed cells and allows key proteins to be produced for a more efficient response to stress [59]. Melatonin treatments significantly increased proline content in lime seedlings under drought stress, indicating how melatonin can potentially assist plants in coping with drought stress (Table 2). These findings are in accordance with earlier studies that proline content reportedly increased in melatonin-treated plants [60-64]. Melatonin may have increased proline levels by inducing the expression of pyrroline-5-carboxylate synthetase 1 ($P5CS_1$), an enzyme involved in proline biosynthesis [65]. The proline content may also increase in response to a decrease of proline oxidase activity during drought stress. When it is present at a proper concentration, melatonin can modulate osmotic metabolism and improve plant tolerance to stress by increasing proline accumulation [60].

Reactive oxygen species interact with phospholipids and fatty acids to accelerate cell membrane liposuction, producing malondialdehyde toxins, thereby increasing membrane permeability in plants and damaging membrane structural integrity [66,67]. Melatonin is thought to enhance the redox state of cells, reducing ROS and reactive nitrogen species levels and stabilizing biological membranes in plant cells [28]. The findings of this study showed that melatonin treatment significantly reduced MDA levels and membrane leakage in plants under drought stress. In melatonin-treated plants, this is accompanied by a decrease in H_2O_2 concentration. It is assumed that melatonin was responsible for the decrease in oxidative stress in both lime species. Melatonin is reportedly able to situate itself between the polar heads of polyunsaturated fatty acids in cell membranes, thereby lowering the level of lipid peroxidation and maintaining natural membrane fluidity [68]. According to several cases of research, melatonin plays a crucial role in preventing lipid peroxidation through its ability to react with lipid peroxyl (LOO•) and lipid alcoxyl (LO•) radicals so that the peroxidation cycle is interrupted and stopped [66,69]. Melatonin is reportedly capable of supporting membrane integrity while limiting lipid peroxidation products and electrolyte leakage in drought-stressed cucumber seedlings [70]. These findings demonstrate that the melatonin treatment can reduce MDA, EL, and H_2O_2 levels, and this observation appears to be consistent with prior drought-related research [71,72]. According to Meng et al. [73], under drought stress, melatonin-treated cuttings of grapevine accumulated smaller amounts of MDA in their leaves and had lower levels of relative electrolytic leakage, compared to grapevine cuttings with no melatonin treatment.

In addition to its direct interaction with ROS, melatonin increases antioxidant enzymes in plants. Plants have evolved defense systems, enzymatic and non-enzymatic, to limit ROS production [74,75]. In the current study, drought stress caused an increase in the activities of CAT, POD, SOD, APX, and GR enzymes in plants. Additionally, the exogenous application of melatonin further increased the activity of these antioxidant enzymes (Figure 1). Increased plant tolerance to stress using melatonin foliar application is due to the antioxidant properties of melatonin, which directly counteracts the harmful effects of reactive oxygen species by stimulating increased production of antioxidant enzymes, and thus improving plant antioxidant capacity [63,76,77]. Overall, melatonin is a broadspectrum antioxidant and a receptor-independent free-radical scavenger that can increase the activity of antioxidant enzymes and other antioxidants to protect plant tissues from oxidative damage [78,79]. After treating the plants with 100 μ M melatonin, the activities of SOD, POD, and APX increased significantly. In this regard, Gantait and Mukherjee [80] demonstrated that melatonin usually serves as a potent long-distance signal and has the capacity to be translocated via vascular bundles from treated leaves or roots to distant tissues, ultimately causing a systemic induction of various abiotic tolerances. According to Li et al. [81], exogenous melatonin is able to reduce stress-induced oxidative damage in Malus hupehensis by directly scavenging H₂O₂ and increasing antioxidant enzyme activity. Previous research by Wei et al. [82], Xia et al. [83], and Arnao and Hernández-Ruiz [41] showed that melatonin treatment increased antioxidant enzyme activity, but reduced ROS content in apple and kiwifruit under abiotic stress. These findings suggest that melatonin therapy reduces the negative effects of drought stress on lime seedlings and, thus, increases their tolerance.

Drought significantly increased the expression of genes involved in ROS scavenging, cellular redox regulation, and proline production. Melatonin increased plant tolerance by further inducing the expression of genes associated with the antioxidant system in response to stress [84–86]. Melatonin-mediated gene regulation during drought plays an important role in controlling cellular signaling pathways [87]. The activities of antioxidant enzymes are connected to the control of important genes that are involved in antioxidant expression. In fact, this supports the redox balance in cells that are under stressful conditions [88]. Plants activate ROS-scavenging gene families when they are under stress. SODs are the first

line of defense against ROS in plants, and they are categorized by the metal ions bound to their active sites, such as copper/zinc (Cu/Zn SOD) and iron (Fe SOD). In drought-stressed plants, SODs defend the photosynthetic apparatus against ROS [89]. In a previous case of research, it was shown that increasing the expression of the SOD Cu/Zn gene in Poncirus trifoliata caused enhancements in antioxidant capacity and assisted in plant tolerance to stress [90]. Exogenous melatonin may promote the production of endogenous nitric oxide as an essential signaling molecule in plants, thereby activating ROS scavenging enzymes under drought conditions [32]. Meanwhile, melatonin can reduce the expression of miR398s and, thus, increase the production of ROS-scavenging enzyme genes such as Cu/Zn SOD and Fe SOD [91].

APX plays a role in the first cyclic phase of AsA-GSH. In fact, AsA-GSH scavenges ROS and protects plants from stress. H_2O_2 is scavenged by this heme enzyme through the AsA-GSH cycle, which converts H_2O_2 to water and dehydroascorbate DHA [92,93]. Meanwhile, *APX* activity usually increases in response to various abiotic stressors. The *APX* gene is capable of many isoforms that have been discovered in the genome. Under abiotic stress, the overexpression of *APX*₂ enhances *APX* activity and leads to a decrease in H_2O_2 and MDA content [94]. According to ElSayed et al. [95], both *APX*₃ and *APX*₄ genes are involved in drought tolerance. At times of drought stress, melatonin usually upregulates genes, such as *APX*₄, that are involved in ascorbate metabolism. Ascorbate biosynthesis is regulated by the *APX*₃ gene, whereas H_2O_2 reduction is regulated by the *APX*₄ gene [94]. Hydrogen peroxide helped apple seedlings resist oxidative stress more when the expression of *APX* was increased [42].

The regulation of proline metabolic genes by drought is a matter of frequent scientific report. It is generally known that dehydration causes an increase in gene expression, resulting in proline production [96]. As a major response to drought stress, the activities of P5CS and P5CR genes reportedly increased [97]. The P5CS gene, which codes for proline, is known to play a key role in the stress response. It brings an accumulation of proline at times of abiotic stress. The accumulation of proline is one of the primary reasons for an increase in osmotic pressure, which improves plant capacity for water retention [98]. P5CS expression was reportedly upregulated in Arabidopsis, oil palm, and three wheat cultivars under osmotic stress [99,100]. Another study considered the expression of P5CS and P5CR genes in two rose cultivars during drought stress. There was no significant increase after the stress period, and *P5CR* gene expression was very low when compared to *P5CS* gene expression. These findings suggested that the *P5CS* gene plays a more important role in proline accumulation in roses under drought stress, as compared to the role of the P5CR gene [101]. We observed appreciably enhanced accumulation of proline, accompanied by a 20-fold increase in P5CS₁ transcript levels and P5CR in lime plants under melatonin-treated compared to control plants. This implies that melatonin positively influences proline accumulation during drought stress in higher plants. These findings are consistent with previous studies showing that melatonin use increases proline and P5CR activity in stressed plants compared with plants not treated with melatonin [102,103].

In plant cells, the mitochondrial transport chain is the prominent site of ROS production. It contains a number of enzymes such as *AOX* and *NADHde* that are involved in the detoxification of ROS [104]. The *AOX* intermittent pathway supposedly plays a role in modulating the production of ROS that are produced during the mitochondrial electron transfer chain [105]. In one study, it was stated that mild-to-moderate drought stress led to a gradual increase in *AOX*₁ and in various components that contribute to ROS inhibition in tobacco [106]. In another study, the use of melatonin (10 μ M) in alfalfa increased the expression of *NADHde* and *AOX* genes under stress conditions. Overall, the recent results showed how melatonin can be a promising agent for enhancements of plant tolerance to drought in ways that involve regulating nitro oxidative homeostasis and protecting plant structures [107].

5. Conclusions

The foliar application of melatonin to lime plants, especially at a concentration of 100 μ M, substantially improved drought stress damage in both types of lime. This occurred by inhibiting membrane damage and reducing the concentration of malondialdehyde and hydrogen peroxide, despite moderate and severe drought stress. These effects were probably achieved by regulating the activity of antioxidant enzymes. Melatonin induced the accumulation of compatible osmolytes such as proline, facilitating its synthesis and thus instilling drought tolerance in plants. In addition, there were increases in gene expressions involved in ROS scavenging, cellular redox regulation, and proline biosynthesis genes, as a result of melatonin treatment on lime leaves. This suggests that exogenous melatonin is an effective protectant that improves drought tolerance in lime seedlings by enhancing antioxidant enzymes and reducing oxidative damage. The use of melatonin can be considered a promising method to reduce the negative impacts of drought stress, although the mechanism by which melatonin aids in drought tolerance would require further research.

Author Contributions: Conceptualization, M.J., A.R.S. and M.T.; methodology, M.J., A.R.S. and M.T.; software, M.J.; validation, M.J., A.R.S. and M.T.; formal analysis, M.J.; investigation, M.J.; resources, M.J. and A.R.S.; data curation, M.J., A.R.S. and M.T.; writing—original draft preparation, M.J.; writing—review and editing, M.J., A.R.S., M.T. and M.H.; visualization, M.J. and M.H.; supervision, A.R.S.; project administration, A.R.S. All authors have read and agreed to the published version of the manuscript.

Funding: There were no available funding resources for this study.

Institutional Review Board Statement: Not applicable.

Informed Consent Statement: Not applicable.

Data Availability Statement: Not applicable.

Acknowledgments: The authors would like to express their gratitude to the College of Agriculture at Shiraz University and the Department of Biotechnology at Isfahan University of Technology for their cooperation during this study.

Conflicts of Interest: The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript, or in the decision to publish the results.

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