



REVIEW

Melatonin Biosynthesis, Growth Regulation, and Adaptability to Environmental Stress in Plants

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ABSTRACT: Melatonin is a multifunctional molecule found in all organisms that has been shown to play a crucial role in plant growth, development, and stress response. Plant melatonin is typically synthesized in organelles termed chloroplasts, and the mechanisms of its synthesis and metabolic pathways have been extensively studied. Melatonin serves a significant regulatory function in plant growth and development, influencing the morphological and physiological characteristics of plants by modulating biological processes. While studies on plant melatonin receptors are in their early stages compared to studies in animal receptors, the binding mechanism with melatonin is now recognized as the key initiating step that triggers a series of downstream protective effects. This suggests that melatonin in plants may exert its effects through two main modes of target binding. The CAND2/PMTR1 protein binds to melatonin with a high degree of affinity. This binding activates downstream heterotrimeric G proteins, which trigger rapid intracellular signaling cascades. These cascades include activating the MAPK pathway and modulating ion channel activity. This action swiftly regulates stomatal closure in response to physiological processes such as drought stress. Additionally, melatonin has been demonstrated to regulate the plant stress response through two main mechanisms. First, it directly inhibits the accumulation of reactive oxygen species. Second, it indirectly influences the stress response pathways. This paper examines plant melatonin from three perspectives: the synthesis pathways of melatonin, its effects on plant growth, and its applications in plants under stress. Finally, the prospects for melatonin study and its applications in plants are discussed.

KEYWORDS: Melatonin; biosynthetic pathway; secondary metabolism; growth regulation; environmental stress

1 Introduction

Melatonin is a small molecule belonging to the class of indole heterocyclic compounds. Its chemical designation of the substance is N-acetyl-5-methoxytryptamine. It was first discovered in the pineal gland of cattle in 1958, and it quickly became apparent in a number of other animals [1]. However, plant-related research did not begin to emerge until the 1980s. Although plant melatonin and animal melatonin have the same chemical structure, their functions differ significantly. In animals, melatonin primarily regulates circadian rhythms and sleep, modulates seasonal reproduction, and provides potent antioxidant and immunomodulatory effects. In contrast, the main role of plant melatonin, aside from regulating



circadian rhythms, is to combat oxidative stress. This protects plants from environmental challenges, such as ultraviolet radiation, drought, and pests and diseases [2,3]. Melatonin plays an important role in plant growth and development. It regulates seed germination, root growth and development, leaf senescence, circadian rhythms, and postharvest fruit ripening. Melatonin is also implicated in the process of plant hormone-mediated signal transduction. By interacting with these pathways, melatonin contributes to the regulation of plant development and growth [4]. Recent studies have made significant progress in understanding melatonin's role in plant responses to abiotic stresses. Melatonin has been demonstrated to enhance the plant's capacity to withstand abiotic stresses, such as drought, salt, cold, and heat, by modulating the expression of genes involved in DREB/CBF, HSF, SOS, and ABA pathways [5].

Although the biosynthesis and signaling pathways of melatonin in plants are well understood, the interaction mechanism between melatonin and other hormones under abiotic stress remains unclear. This paper summarizes the synthesis and metabolic pathways of melatonin in plants to establish a foundation for future research on its regulatory mechanisms. The text delineates how melatonin promotes plant growth and development while simultaneously outlining its interactions with plant hormones (e.g., IAA, CTK, and ABA) in regulating plant growth and development. This facilitates a more profound understanding of the combined effects of these hormones on plant growth and development. Despite numerous studies examining the biosynthesis and regulation of melatonin in plants, the connection between melatonin and abiotic stress remains unclear. This review provides an in-depth overview of the crucial role of melatonin in plant growth and its response to abiotic stresses. Additionally, we discuss the effect and related applications of melatonin in response to various abiotic stresses and present the prospects for plant melatonin in agriculture. As a green, highly efficient biostimulant, melatonin offers potential solutions for addressing global food security, reducing chemical pesticide use, and coping with stresses caused by climate change. Melatonin significantly enhances plants' tolerance to various environmental stresses. Spraying melatonin can improve fruit set rates in certain fruits and vegetables, increase individual fruit weight, and thereby boost yields. When pathogenic microorganisms invade, melatonin treatment alerts plants to preemptively accumulate defense compounds and synthesize pathogen-associated proteins. The agricultural value of melatonin lies in its multifunctionality, natural origin, and safety profile. Unlike traditional chemical pesticides or fertilizers, it functions by regulating the plant's own physiological and defense systems, representing an empowering rather than coercive strategy.

2 Biosynthetic and Metabolic Pathways of Plant Melatonin

2.1 Biosynthetic Pathway of Melatonin

Melatonin has a low molecular weight and a stable chemical structure. It occurs naturally in a variety of plants. When ¹⁴C-labeled tryptophan was introduced into plants such as *Hypericum perforatum*, it was found that they contained indoleacetic acid, tryptamine, 5-hydroxytryptophan, and 5-hydroxytryptamine, which are intermediates of the melatonin synthesis pathway. It has also been established that the presence of these substances is an inherent feature of the melatonin synthesis process in animals [6]. The evidence also indirectly shows that the melatonin synthesis pathway of plants is similar to that of animals. However, the synthesis pathways are still different between the two. Animals cannot synthesize the melatonin precursor substance, tryptophan, by themselves, but plants can. We speculate that the synthesis of plant melatonin may be more complex.

Many studies have indicated that there are multiple pathways for plant melatonin biosynthesis. As a general rule, the process of melatonin synthesis can be considered to follow four primary pathways, all of which necessitate the participation of more than six enzymes. These include tryptophan decarboxylase

(TDC), tryptamine 5-hydroxylase (TPH), tryptamine 5-hydroxylase (T5H), 5-hydroxytryptamine-N-acetyltransferase (SNAT), N-acetyl-5-hydroxytryptamine methyltransferase (ASMT), and caffeic acid O-methyltransferase (COMT) [7]. In all four pathways, there is an important intermediate product, 5-hydroxytryptamine (serotonin). The production of 5-hydroxytryptamine requires two-step enzymatic reactions. Firstly, the decarboxylation of tryptophan is initiated by the enzyme tryptophan decarboxylase, resulting in the formation of tryptamine. Subsequently, the subsequent synthesis of 5-hydroxytryptamine is catalysed by the action of tryptophan-5-hydroxylase. An alternative synthetic pathway involves the hydroxylation of tryptophan by tryptamine 5-hydroxylase, followed by decarboxylation by tryptophan decarboxylase (TDC) to produce 5-hydroxytryptamine [4]. In the first and second biosynthetic pathways, 5-hydroxytryptamine is synthesised in the endoplasmic reticulum. In contrast, in the third and fourth biosynthetic pathways, the same process occurs in the cytoplasm (Fig. 1). Studies have also indicated that tryptophan hydroxylase is a copper-dependent enzyme that can catalyze the conversion of tryptophan into 5-hydroxytryptophan. Additionally, 5-hydroxytryptophan in cells is primarily transported to the cytoplasm via plasma membrane transporters and is subsequently catalyzed by decarboxylase to produce 5-hydroxytryptamine. Therefore, in different melatonin biosynthetic pathways, the variance in the synthesis site of 5-hydroxytryptamine—whether in the endoplasmic reticulum or the cytoplasm—may be associated with the localization of tryptophan hydroxylase and tryptophan decarboxylase [8]. The synthesis of plant melatonin is relatively complex, and its crosstalk with other hormones requires further validation.

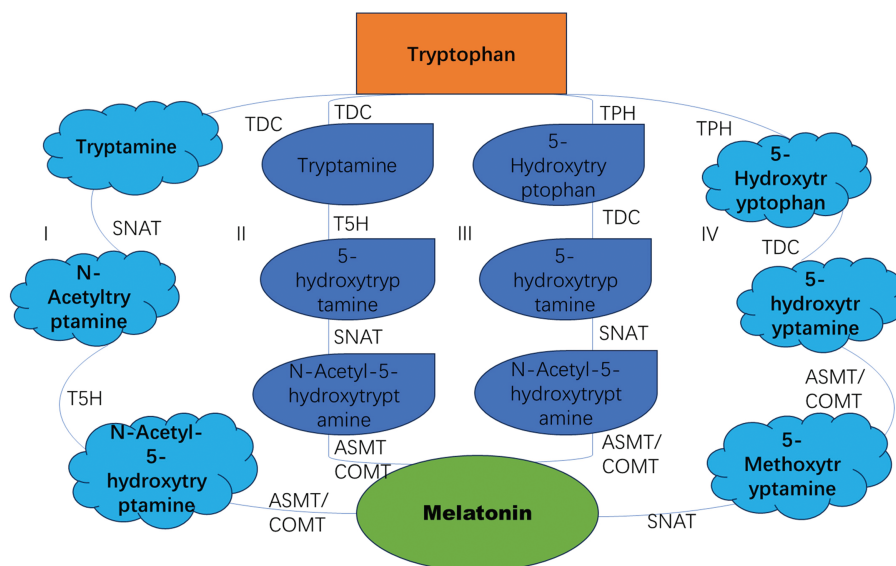


Figure 1: The biosynthesis pathway of melatonin in plants

This is a classic melatonin biosynthesis pathway. TDC (tryptophan decarboxylase), TPH (tryptophan hydroxylase), T5H (tryptamine 5-hydroxylase), SNAT (serotonin N-acetyltransferase), ASMT (N-acetylserotonin methyltransferase), COMT (caffeic acid O-methyltransferase), and ASDAC (N-acetylserotonin deacetylase).

2.2 Metabolic Pathway of Melatonin

Melatonin, as an important bioactive molecule, can be degraded through enzymatic conversion pathways. In plants, melatonin degradation is primarily catalyzed by melatonin N-acetyltransferase (NAT), which converts it into N-acetylmelatonin, and by melatonin-O-methyltransferase (HIOMT), which further

catalyzes the production of 6-hydroxymelatonin. This compound is subsequently metabolized and degraded by chloroplast esterase. A similar degradation pathway is present in vertebrates and other eukaryotic organisms; NAT and HIOMT also play crucial roles in melatonin metabolism in mammals [9]. The metabolic pathway of melatonin is subject to influence by many factors, including oxidative stress, the endocrine system, and environmental changes. Studies have confirmed that the metabolite of melatonin in water hyacinth, N1-acetyl-N2-formyl-5-methoxykynuramine (AFMK), has a high antioxidant effect and found that the content of AFMK and melatonin is high under strong light, which is believed to be related to plant photosynthesis or light protection [10]. AFMK has shown distinct rhythms in water hyacinth, with peaks occurring during the photophase. As light intensity increases, the levels of melatonin in the water hyacinth appear to rise as well [10]. Plant melatonin undergoes hydroxylation by M2H and M3H to form two hydroxylated metabolites, namely 2-hydroxymelatonin (2-OHM) and cyclic 3-hydroxymelatonin (C3-OHM) [11]. 2-hydroxymelatonin (2-OHM) is derived from the hydroxylation of M2H, among which M2H is one of the most important metabolic pathways. The gene of melatonin hydroxylation metabolite C3-OHM was cloned in rice and expressed in the cytoplasm, catalyzed by the M3H enzyme [12]. To better understand the biological function of melatonin, it is critical to further reveal its metabolism.

3 Effect of Melatonin on Plant Growth and Development

3.1 Melatonin Stimulates Plant Growth

Exogenous melatonin can increase the fresh weight and root number of plant growth. For instance, the spraying of exogenous melatonin has been demonstrated to be a highly efficacious method to promote the growth parameters of wheat seedlings, including root length, root number, fresh weight, and dry weight [13]. Additionally, it was demonstrated that the addition of various concentrations of melatonin to pineapples and tomatoes significantly enhances plant rooting. This effect is somewhat reminiscent of the effects of the growth hormone IAA [14]. The effects of varying concentrations of melatonin on plant root growth and development are entirely divergent. Melatonin does exhibit significant dose-dependent effects in its application: low concentrations may promote growth, while high concentrations may have the opposite effect. High concentrations of melatonin inhibit the growth and development of roots while slowing their accumulation of beneficial compounds. Conversely, low concentrations of melatonin significantly boost nutrient accumulation in plants, thereby enhancing their stress resistance and growth potential [15]. This paradoxical effect stems from the fact that melatonin triggers multiple distinct signaling pathways, each with different activation thresholds and ultimate effects at different concentrations [16,17]. Low concentrations of melatonin directly scavenge ROS and upregulate endogenous antioxidant enzyme systems, whereas extremely high concentrations of melatonin molecules may react with certain metal ions, unexpectedly generating small amounts of ROS with signaling functions. ROS signaling may inhibit the expression of genes associated with cell division and elongation, leading to growth arrest [18]. Furthermore, melatonin-induced reactions require significant energy expenditure. High concentrations of melatonin strongly activate systemic defense responses like systemic acquired resistance, consuming significant energy and carbon skeleton resources [19]. After 24 h of treatment, melatonin promotes the growth of Arabidopsis roots, stems, and leaves. This effect is efficient, further demonstrating the significance of melatonin in root growth and development [20]. Moreover, varying concentrations of melatonin influence the elongation of rice tiller buds. The application of appropriate concentrations of melatonin can not only encourage the elongation of rice tiller buds but also mitigate the inhibition of bud elongation caused by high concentrations of basic amino acids [21]. Melatonin not only influences root growth but also impacts the photoperiod and flowering of plants. It has been demonstrated that the exogenous addition of melatonin can ameliorate photoperiod disorders and promote the flowering of Arabidopsis plants when subjected to a 12-h light cycle.

The transcription levels of flowering genes *FT*, *SOC1*, and *LFY* increase significantly, thereby enhancing the flowering process [22]. Therefore, melatonin can serve as a small-molecule active substance to regulate plant physiological processes, such as cell expansion and root growth, with varying concentrations producing different effects (Fig. 2).

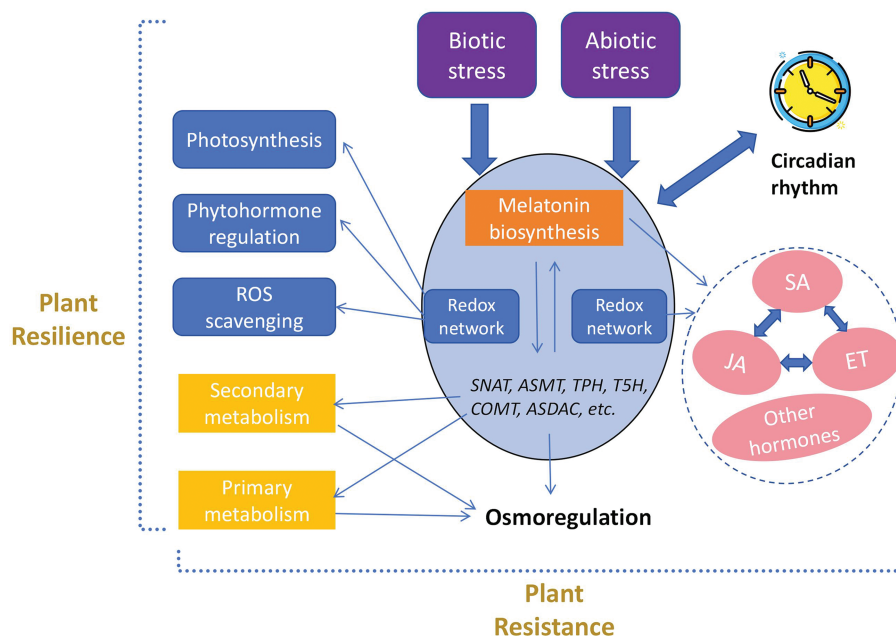


Figure 2: The role of melatonin in regulating stress responses through the mediation of redox networks

Melatonin enhances plant resilience by improving photosynthetic efficiency, regulating the synthesis of other plant hormones, promoting the accumulation of primary and secondary metabolites, and scavenging reactive oxygen species (ROS). It combats oxidative stress caused by various biotic and abiotic stresses by regulating the synthesis of key enzymes within the redox network.

3.2 Melatonin Inhibits Leaf Senescence

Leaf senescence, an important organ of plant photosynthesis, is directly related to crop yield [23]. A proteomic analysis of apple trees subjected to prolonged melatonin root irrigation indicated that it can inhibit the activity of hydrolases associated with photosynthesis and redox responses in apple leaf plastids, thereby effectively delaying the senescence of apple leaves [24]. Furthermore, melatonin administration to the roots of grapevines led to a reduction in the extent of photosynthetic activity inhibition caused by NaCl, while simultaneously enhancing the maximum fluorescence and maximum photochemical efficiency of leaves under light conditions. Therefore, the root application of melatonin mitigates the injury to the photosynthetic process of grape leaves under NaCl stress [25]. Moreover, melatonin delays plant leaf senescence primarily by boosting the activity of antioxidant enzymes, maintaining the activity of leaf photosynthetic enzymes, improving the efficiency of photosystem II, and decelerating the degradation rate of chlorophyll. It also prevents protein degradation by regulating sugar and nitrogen metabolism [26]. First, melatonin directly scavenges ROS by upregulating certain antioxidant enzyme genes and enhancing the endogenous antioxidant system, maintaining cellular redox homeostasis and thereby delaying the onset of leaf senescence [27]. Melatonin strongly inhibits the expression of the *NYE1/SGR1* genes, thereby blocking the chlorophyll degradation

pathway at its source and maintaining photosynthetic capacity [28]. Melatonin also regulates the senescence-associated transcription factor network. ORE1 is a NAC family transcription factor that positively regulates aging. By inhibiting the expression of *ORE1*, melatonin abrogates its activation of numerous downstream senescence-associated genes (SAGs) [29]. ANAC016, another pro-aging NAC transcription factor, is also inhibited by melatonin [30]. Certain WRKY family members, such as WRKY6, also promote senescence, and melatonin inhibits their expression [31]. Melatonin has been shown to upregulate the expression of core autophagy-related genes (*ATGs*). Some studies suggest that moderate and orderly autophagy is beneficial for delaying aging, and melatonin-induced autophagy tends to be a type of “maintenance autophagy,” promptly removing damaged mitochondria and misfolded proteins [32]. Therefore, melatonin does not act through a single gene but rather as a system regulator that works together at multiple levels to delay leaf senescence.

3.3 Interaction between Melatonin and Other Phytohormones

Melatonin and ABA have antagonistic effects in regulating leaf senescence, yet the specific mechanism of action remains unclear. Exogenous melatonin can delay the senescence of mature melon leaves under ABA stress and protect the functionality of photosystem II. Under high concentrations of ABA stress, melatonin induced the expression of the respiratory burst oxidase homolog D (*CmRBOHD*) gene in melon and promoted the accumulation of H_2O_2 [33]. Moreover, melatonin has been demonstrated to regulate the synthesis and metabolism of the plant growth hormone (IAA), thereby exerting a significant influence on the growth, development, and physiological metabolism of plants. In an experiment where melatonin and IAA interacted to regulate the growth of *Arabidopsis* taproots, melatonin inhibited the expression of key genes in the cytokinin signaling pathway (*AHK4*, *AHP*, *ARR15*), consequently affecting the regulatory role of cytokinin on root growth. Therefore, the similar effects of melatonin and IAA may stem from their regulatory influences on plant growth hormones rather than merely acting as substitutes for them [20].

GA is a plant hormone that plays a pivotal role in various processes, including plant growth and development. It has been demonstrated that under conditions of salt stress, melatonin is capable of upregulating the expression of genes involved in the synthesis of GA, specifically, *CsGA20ox* and *CsGA3ox*, in cucumber seeds. In addition to this, melatonin appears to enhance the synthesis of GA, particularly the hydroxy-GA (HA) form, whilst concomitantly mitigating the inhibitory effects of NaCl-induced stress on seed germination. Melatonin interacts with GA to mediate plant growth by regulating its biosynthesis, transport, and signal transduction [34]. Ethylene is involved in the control of various physiological processes, including plant growth, development, and responses to adversity. The application of exogenous melatonin was found to inhibit ethylene release in pear fruit to a significant extent. It was also demonstrated to reduce peel colour change, as well as to delay the decline in titratable acid in the fruit. Furthermore, chlorophyll content was observed to be increased by the treatment. Melatonin can impact plant nutrient absorption, fruit ripening, and stress resistance by regulating ethylene synthesis and release [35].

The crosstalk between ABA and melatonin constitutes a key hub in the finely regulated growth-defense trade-off in plants. In many circumstances, ABA and melatonin exhibit antagonistic physiological effects, including on processes such as stomatal closure, seed germination, and leaf senescence. However, this antagonism is not simply a cancellation but rather involves complex interactions at multiple levels of the signaling pathway. In the presence of melatonin, it scavenges or attenuates ROS generated in the ABA signaling pathway, thereby interrupting or weakening ABA downstream signaling outputs, such as stomatal closure [27]. ABA and melatonin both activate the MAPK pathway, but they may activate different members or induce distinct phosphorylation patterns, ultimately activating different transcription factors and leading to opposing gene expression [36]. At the transcriptional level, ABA activates the expression of ABRE-binding genes and upregulates senescence-related transcription factors, while melatonin inhibits the expression

of these ABA-induced senescence and stress-related genes [37]. Furthermore, exogenous ABA treatment downregulated the expression of genes (*T5H* and *ASMT*) involved in melatonin biosynthesis, leading to reduced endogenous melatonin levels [38]. Therefore, ABA and melatonin have a relationship similar to the “brake and accelerator” in plant growth and stress. In summary, melatonin coordinates with various plant hormones during growth and development, regulating plant growth, development, and responses to adversity by modulating hormone synthesis, transport, and signal transduction.

4 Roles of Melatonin in Plant Abiotic Stress

In light of the recognised function of melatonin in mitigating abiotic stress in plants, researchers have initiated studies on the administration of exogenous melatonin prior to stress events to enhance plant tolerance. Some studies suggested that exogenous administration of melatonin to plant seeds or the most vulnerable stages of plant growth and development, either through soaking, irrigation, or foliar spraying, can attenuate adverse impacts associated with abiotic stresses such as cold, heat, heavy metals, drought, waterlogging, and salt stress (Table 1). The current frontier and critical focus of plant melatonin studies is exploring the variations in melatonin efficacy across different stress types and plant species. It indicates that the efficacy of melatonin exhibits significant ‘stress-species’ specificity [39]. This variation primarily stems from two mechanisms. Firstly, different abiotic stresses (such as drought, salinity, heavy metals, and extreme temperatures) cause primary damage and secondary oxidative stress through distinct pathways [40,41]. Meanwhile, melatonin’s core function lies in its role as a potent antioxidant and regulator of oxidative stress signalling. Thus, for stress types where oxidative bursts constitute primary secondary damage (e.g., drought, salt, and heavy metal stress), the application of exogenous melatonin typically exhibits higher protective efficacy by directly scavenging ROS and upregulating endogenous antioxidant enzyme systems [42]. Conversely, for stresses causing direct physical damage, their effects may be weaker and more indirect. Additionally, inherent variations in melatonin sensitivity exist among different plant species and even cultivars. These variations are closely linked to baseline endogenous melatonin levels, metabolic rates, receptor protein affinities, and the completeness of downstream signalling pathways [43]. For example, species with lower endogenous melatonin levels may exhibit heightened sensitivity and more pronounced responses to exogenous supplementation [44]. Conversely, certain species may possess more efficient melatonin synthesis pathways or signalling networks, enabling the rapid amplification of exogenous melatonin signals to coordinate a series of physiological stress responses, such as stomatal closure, the induction of stress-resistant gene expression, and the regulation of ion homeostasis [45]. The species that are inherently vulnerable to specific stresses, or that have limited endogenous antioxidant capacity, often benefit more from melatonin supplementation.

Table 1: Exogenous application of melatonin in regulating plant abiotic stress tolerance

Abiotic stress	Plant species	Melatonin treatment		Changes and mechanisms	Reference
		Concentration	Application		
Drought	Rice	200 µM	Root application	Melatonin activated the expression of ALMT1, OsPOX1, OsCATC, and OsAPX2 and reduced MDA and ROS content.	[46]

(Continued)

Table 1 (continued)

Abiotic stress	Plant species	Melatonin treatment		Changes and mechanisms	Reference
		Concentration	Application		
	Maize	50 μ M	Foliar spray	Melatonin enhanced the drought tolerance of maize by modulating the metabolic processes of carbon (C) and nitrogen (N).	[21]
	Tomato	100 μ M	Root application	Melatonin application enhances transcript levels of plant antioxidant defense enzymes and antioxidant stress-responsive genes (SICAT1, SIAPX, SIGR, etc.).	[47]
	Pepper	100 μ M	Foliar spray	Administration of melatonin has the potential to reduce oxidative stress and improve nitrogen metabolism by activating some enzymes, including NR, NiR, GS, GOGAT, and GDH.	[48]
Salt	Banana	100 μ M	Root application	Melatonin has been demonstrated to induce the accumulation of MDA, and members of the transcription factor family, including MYB, NAC, bHLH, and WRKY, have the potential to contribute to the reduction of salt stress tolerance in banana.	[49]
	Rice	150 μ M	Root application	Maintaining a low ROS state by increasing total antioxidant capacity, promoting lutein cycling and increasing lutein pool size to dissipate excess light energy, and increasing the activity of key photosynthetic enzymes.	[50]

(Continued)

Table 1 (continued)

Abiotic stress	Plant species	Melatonin treatment		Changes and mechanisms	Reference
		Concentration	Application		
	Sugar beets	100 μ M	Root application Foliar spray	Melatonin maintains K/Na homeostasis and reduces Na toxicity. In addition, melatonin enhances antioxidant defence systems by regulating the ASA-GSH cycle and mediating the phenylalanine pathway, thereby maintaining plant cell membrane integrity.	[51]
	Pigeon pea	50 μ M	Root application	Melatonin application of wood beans promotes lignan biosynthesis via the CcPCL1 and CcF3'H-5 pathways, resulting in salt tolerance.	[52]
Cold	Arabidopsis	10 and 30 μ M	Root application	Melatonin enhanced the expression of C-repeat-binding factors (CBFs)/drought response element-binding factors (DREBs) in response to cold stress.	[53]
	Bermudagrass	100 μ M	Root application	Differential regulation of melatonin responses to cold stress is mainly through influencing the antioxidant system, photosystem II, and metabolic homeostasis.	[54]
Heat	Naked oat seedlings	100 μ M	Root application	Melatonin induces the expression of MAPKs and TFs and regulates the expression of downstream stress-responsive genes, thereby improving plant tolerance.	[55]

(Continued)

Table 1 (continued)

Abiotic stress	Plant species	Melatonin treatment		Changes and mechanisms	Reference
		Concentration	Application		
	Chrysanthemum seedlings	200 μ M	Root application Foliar spray	Melatonin regulates heat shock transcription factors (HSFs) and heat shock proteins (HSPs), and gene 2+ signaling involved in Ca (CNGC and CAM/CML).	[56]
	Tomato seedlings	100 μ M	Root application	Melatonin pretreatment improved the heat tolerance of tomato seedlings by improving their antioxidant defence mechanisms, inducing the ascorbate-glutathione cycle, and reprogramming the PAs metabolism and NO biosynthesis pathways.	[57]

4.1 Application of Melatonin under Drought Stress

Drought is a factor that seriously affects crop growth. Plants can replenish water in dry root systems during drought conditions through hydraulic redistribution, which increases the water potential of the root systems and helps maintain normal physiological functions [58]. Under drought stress, an imbalance between the generation and removal of ROS in plant cells has been shown to result in damage to the membrane system. The application of antioxidants can maintain the balance of reactive oxygen species in cells [59]. As a natural antioxidant, melatonin can increase the content and activity of antioxidant substances in plants, thereby alleviating the effects of drought stress on plants. The optimal growth temperature for pepper is 25–30°C; however, temperatures below 15°C negatively impact seed development. The application of 5 μ M melatonin to the roots of plants such as *Capsicum* peppers has various positive effects, including an increase in seed biomass, improvements in photosynthesis and antioxidant activity, and a reduction in membrane permeability, MDA, hydrogen peroxide, and other indicators. Melatonin pretreatment can also double the early yield of pepper under cold stress [31]. Studies have shown that foliar spraying with melatonin reduces oxidative damage and enhances net photosynthesis in plants under cold stress [60]. The administration of melatonin has been shown to protect photosynthetic pigments, counteract the impact of drought stress on *Amorpha fruticosa* seedlings, and increase antioxidant enzyme activity [61]. The application of melatonin to *Psoralea corylifolia* and sweet sorghum can enhance their drought tolerance, decrease lipid peroxidation and oxidative stress, facilitate the recovery of leaf structure and function, boost photosynthetic pigment levels, and increase antioxidant enzyme activity [62,63]. Thus, exogenous melatonin can reduce drought-stressed crops' damage. When the ambient temperature exceeds the optimal temperature of the plant, heat stress occurs. *Festuca arundinacea* is sensitive to high temperatures during the seedling stage. Eight-day-old tall fescue seedlings exhibited improved tolerance to elevated temperatures when 20 μ M melatonin was added to their growth medium. Treatment with melatonin increased the fresh weight, plant

height, chlorophyll content, and protein levels of the seedlings. It also enhanced the activities of antioxidant enzymes such as SOD, POD, and CAT, and reduced electrolyte leakage, MDA, and hydrogen peroxide levels. Additionally, melatonin treatment upregulated several genes associated with heat response, while downregulating other genes such as FaF-box, FaHSFA6B, and FaCYP710A. These results indicate that the application of melatonin can enhance the adaptability of tall fescue to heat stress [64]. Drought negatively impacts crop growth and yield, but variations in melatonin-treated crops and their mechanisms of adaptation determine their sensitivity to drought and their ability to withstand adverse conditions.

4.2 Application of Melatonin in Salt Stress

Salinisation is a significant threat to the ecological environment and agricultural production, ranking among the most prevalent abiotic stresses affecting global agriculture. Salt stress has a negative impact on plant growth and development. The damage caused by excessive salt mainly includes osmotic stress, ion toxicity, and oxidative stress [65]. Exogenous melatonin can enhance the antioxidant and osmotic regulation capabilities of rice, remove a large amount of accumulated ROS, alleviate the oxidative damage, and improve the salt tolerance of seedlings [46]. The application of melatonin can reduce the harm of salt stress to plants and promote plant growth and development. Cotton seedlings showed growth inhibition under salt stress. The levels of hydrogen peroxide and superoxide anions in the plant increased, while the levels of osmotic regulators in the leaves decreased. This data indicates that the seedlings produced excessive ROS, and the cell membrane was damaged. The application of exogenous melatonin to cotton seedlings inhibits the production of reactive oxygen species (ROS), enhances the activity of the antioxidant enzyme system, and increases the levels of substances that regulate osmosis. It also reduces membrane lipid peroxidation and protects the integrity of the lipid membrane under salt stress. These actions mitigate the damage caused by salt stress to seedlings and effectively reduce its negative impact on their growth [66]. Under salt stress conditions, wheat seedlings treated with melatonin showed a certain degree of growth recovery. Dry weight, IAA content, leaf photosynthesis rate, maximum photochemical efficiency of photosystem II, and chlorophyll content were all higher than in the salt stress control group. However, they remained lower than in the blank control group under normal conditions. Melatonin alleviated the inhibitory effect of salt stress on the growth of the whole wheat plant to a certain extent. It significantly alleviated salt-induced oxidative damage to upland cotton, which was achieved by increasing the accumulation of osmotic-regulating substances and activating the activity of antioxidant enzymes [67]. In addition, melatonin also enhances the tolerance of upland cotton to salt stress by regulating the expression of stress response genes and ion channel genes [68]. Salt stress negatively affects the expression patterns of melatonin biosynthetic enzymes and increases endogenous melatonin levels. Overexpressing SNAT significantly increases plant salt tolerance. For instance, Arabidopsis plants overexpressing VvSNAT1 exhibited greener leaves, more vigorous growth, and higher germination rates in response to salt stress than wild-type lines [69].

Melatonin fundamentally helps maintain ion balance to combat salt stress. For example, the application of melatonin increased the accumulation of potassium ion, decreased the absorption of sodium ion, and maintained a high K^+/Na^+ ratio, thereby enhancing the salt tolerance of corn seedlings. Melatonin positively influences the response of plants to salt stress. It can regulate the growth and development of plants, boost their antioxidant capacity, and mitigate the oxidative damage caused by salt stress. It can also increase their tolerance to these conditions [70]. Melatonin upregulates and enhances the activity of SOS1, SOS2, and SOS3 proteins, which regulate the functionality of the entire SOS pathway by driving the SOS1 protein to actively pump excess sodium out of the cytoplasm. This process utilizes the proton gradient established by the plasma membrane H^+ -ATPase, which directly contributes to the reduction of intracellular sodium concentrations [71,72]. Melatonin significantly upregulates the expression and protein

activity of sodium/hydrogen transporters such as NHX1. It also enhances the activity of the tonoplast H^+ pump, providing a stronger driving force for sodium compartmentalization [73]. Under salt stress, high sodium concentrations inhibit potassium absorption channels. Melatonin, through its potent antioxidant properties, reduces oxidative damage to potassium channels, such as AKT1, caused by ROS generated by salt stress [74,75]. Melatonin also reduces potassium leakage by maintaining membrane integrity. However, the effects of melatonin can vary between different plant species and varieties. Further studies are needed to uncover how it works and how it can be used.

4.3 Application of Melatonin in Cold Response

Low-temperature environments will affect the fluidity and enzyme activity of plant cell membranes, inhibiting photosynthesis and nutrient transport, causing damage to the plant body, and leading to a reduction in crop yield [76]. Melatonin improves rice's tolerance of low-temperature stress by regulating the activity of enzymes in the antioxidant system, the levels of osmotic substances and chlorophyll, the levels of plant hormones, and the expression of genes that confer cold resistance in rice [77]. Spraying melatonin can promote the growth of cucumber seedlings in low temperatures, enhance the osmotic regulation capability of the leaves, maintain the balance of plant hormones, and thus improve the cold resistance of cucumber seedlings [78]. By adding appropriate concentrations of exogenous melatonin, the growth of seedlings can be enhanced while reducing the levels of MDA and H_2O_2 in the leaves and increasing the content of AsA and GSH, which helps mitigate the damage caused by low-temperature stress to potato seedlings, thereby improving their resistance to low-temperature stress to varying degrees [79]. Exogenous melatonin has also been used to enhance the growth of highland barley seedlings under cold stress. The results showed that exogenous melatonin could restore circadian oscillations in the clock genes *HvCCA1* and *HvTOC1*. These genes had lost their circadian phenotype due to environmental cold stress [80]. These studies have demonstrated the potential of exogenous melatonin in enhancing plant tolerance to low-temperature stress. Spraying exogenous melatonin can promote the growth of plant seedlings, increase the activity of antioxidant enzymes and the capability of osmotic regulation, and maintain the balance of plant hormones. This increases the resistance of plants to low-temperature stress. Furthermore, exogenous melatonin can regulate plant clock genes and restore circadian oscillations, thereby mitigating the effects of low temperatures on plants. Using exogenous melatonin to improve crop resistance to low-temperature stress provides a theoretical basis and practical guidance.

4.4 Application of Melatonin in Heat Stress

High temperatures cause plant stomata to close, affect photosynthesis, and interfere with water balance and mineral metabolism. It can also cause antioxidant system disorders and increase oxidative damage. These changes have a significant impact on plant growth and reproduction, resulting in lower crop yields [81]. In *Arabidopsis*, melatonin-mediated heat tolerance has been found to be associated with the increased expression of *HSEF2*, *HSA32*, *HSP90*, and *HSP101*. Furthermore, Hsp40 interacts with SNAT to regulate melatonin biosynthesis, thereby enhancing heat tolerance by stabilising the Rubisco enzyme during periods of heat stress [82]. The administration of exogenous melatonin can inhibit the upregulation of genes involved in pyruvate synthesis and the downregulation of genes related to pyruvate consumption in celery. This prevents pyruvate accumulation from rapidly increasing the expression of peroxidase genes and enhancing peroxidase activity. Physiologically, the application of exogenous melatonin in celery significantly enhanced the plant's ability to remove ROS under heat stress and improved the plant's heat tolerance [83]. Exogenous melatonin alleviated the heat damage to plants by improving antioxidant capacity, reducing ROS accumulation, and upregulating the transcription of *HSF7*, *HSP70.1*, and *HSP70.11*. Specifically, the heat

damage symptoms were mild, and the heat damage index was reduced. Applying melatonin improves a plant's resistance to high temperatures. This enhancement occurs through several mechanisms, including increasing the activity of antioxidant enzymes APx and GR, boosting the antioxidant compound levels of AsA and GSH, inhibiting ROS production, safeguarding cell membrane stability, and enhancing the plant's ability to endure high temperatures [45].

Endogenous melatonin increases significantly under high-temperature stress, which may reflect a response mechanism for plants to resist such stress. This enhancement occurs as it can induce the expression of *HSFA1a*, *HSFA1b*, *HSFA1d*, and *HSFA1e* genes, thereby activating downstream heat shock protein genes and improving the tolerance to high temperatures. Thus, both high temperature and melatonin can jointly induce the expression of the *HSFA1* gene; however, melatonin does not enhance the resistance of *HSFA1* mutant plants to high temperatures. Additionally, exogenous melatonin inhibits the expression of downstream heat shock protein genes that are activated by *HSFA1* [84]. Both high temperature and exogenous melatonin induce the expression of the *HSFA1* gene, enhance the expression of *HSP*, and improve plant tolerance to high-temperature stress. The results suggest that melatonin plays a significant regulatory role in how plants respond to high-temperature stress. Using exogenous melatonin can increase a plant's antioxidant capacity, maintain the stability of its cell membranes, and regulate the expression of heat shock transcription factors and heat shock protein genes. This ultimately enhances a plant's tolerance to elevated temperatures.

5 Conclusion Remarks

The first melatonin study demonstrated its consistent action across plants, animals, and humans. For instance, melatonin may play a role in life phenomena such as light morphogenesis by acting as a time regulator. Since plant melatonin shares structural similarities with IAA, a previous study sought to determine whether melatonin exhibits similar physiological functions as IAA, such as promoting growth, root development, and seed germination. Both the methoxy group in the fifth position of the indole ring of melatonin and the N-acetyl group in the side chain of melatonin are crucial for its ability to scavenge ROS. Consequently, the antioxidant effects of melatonin in plants have also become a focal point of research, encompassing resistance to adversity and responses to both biological and abiotic stresses, as well as aging and apoptosis related to ROS. Recently, the primary focus has shifted toward verifying the hypotheses from earlier studies and expanding to include all aspects of melatonin. These include different plant organs, cultivated and wild species, plant-derived products, the molecular mechanisms of melatonin, and metabolic pathways. Despite advances in plant melatonin research, some scientific issues remain. First, while numerous studies have examined the synthesis pathway of plant melatonin, fewer have investigated plant melatonin receptors. An attractive and challenging direction for future research is exploring whether melatonin has additional receptors by leveraging existing research, along with advanced facilities and technologies. What role does exogenous plant melatonin play in the core clock-phytomelatonin-redox network? The signaling pathway of plant melatonin is not fully understood. The synergistic effects of plant melatonin and other plant hormones in regulating physiological processes remain unclear. In addition, the role of melatonin signaling at the interface between soil microorganisms and roots in relation to plant growth, ion transport, and nutrient distribution requires further exploration.

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