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Physio-biochemical and molecular stress regulators and their crosstalk for lowtemperature stress responses in fruit crops: A review

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Low-temperature stress (LTS) drastically affects vegetative and reproductive growth in fruit crops leading to a gross reduction in the yield and loss in product quality. Among the fruit crops, temperate fruits, during the period of evolution, have developed the mechanism of tolerance, i.e., adaptive capability to chilling and freezing when exposed to LTS. However, tropical and sub-tropical fruit crops are most vulnerable to LTS. As a result, fruit crops respond to LTS by inducing the expression of LTS related genes, which is for climatic acclimatization. The activation of the stress-responsive gene leads to changes in physiological and biochemical mechanisms such as photosynthesis, chlorophyll biosynthesis, respiration, membrane composition changes, alteration in protein synthesis, increased antioxidant activity, altered levels of metabolites, and signaling pathways that enhance their tolerance/ resistance and alleviate the damage caused due to LTS and chilling injury. The gene induction mechanism has been investigated extensively in the model crop Arabidopsis and several winter kinds of cereal. The ICE1 (inducer of Crepeat binding factor expression 1) and the CBF (C-repeat binding factor) transcriptional cascade are involved in transcriptional control. The functions of various CBFs and aquaporin genes were well studied in crop plants and their role in multiple stresses including cold stresses is deciphered. In addition, tissue nutrients and plant growth regulators like ABA, ethylene, jasmonic acid etc., also play a significant role in alleviating the LTS and chilling injury in fruit crops.

However, these physiological, biochemical and molecular understanding of LTS tolerance/resistance are restricted to few of the temperate and tropical fruit crops. Therefore, a better understanding of cold tolerance's underlying physio-biochemical and molecular components in fruit crops is required under open and simulated LTS. The understanding of LTS tolerance/resistance mechanism will lay the foundation for tailoring the novel fruit genotypes for successful crop production under erratic weather conditions.

KEYWORDS

cold tolerance, transcriptome, aquaporins, membrane injury, phytohormone, CBFs

Introduction

Plants are sessile and experience several abiotic stresses episode in their life due to harsh climatic conditions. These circumstances triggered several molecular changes in the plants, which led to a variety of responses of physiology and biochemistry of the plant cells, which determine the fitness of the plant in the environment. Temperature is one of the decisive and essential abiotic factors regulating plant's ontogeny. The process for how Plants sense temperature is elusive. However, several candidate thermos-sensors proteins are known that respond to the membrane changes and other cellular changes triggered by low temperature, leading to downstream events of change in the pattern of gene expressions and plant's responses (Ding and Yang, 2022). Low temperature (LT) influences plant growth and development by causing various morphophysiological and biochemical changes (Nishiyama, 1976; Medina et al., 2011) (Figure 1). LT has an impact on many processes, including, photosynthesis, cell division, membrane stability, absorption and transport of water and nutrients, yield, and ultimately species survival. Plants experience low temperature stress (LTS) in two ways: (i) Chilling stress: when plants are exposed to LT below 10–15°C for a certain period, causing injury without formation of ice crystals within the plant



cells; (ii) Freezing stress: when plants are exposed to sub-zero °C low temperatures, inducing ice crystallization of cellular content, resulting in cell dehydration and freezing injury (Beck et al., 2007; Zhu et al., 2007). Two processes lead to freezing injury: (i) vitrification, which results from the fast freezing of cellular substance, and (ii) supercooling, which results in the formation of ice crystals from intra or extracellular content. An overview of the optimum temperature range for vegetative and reproductive growth for some tropical and subtropical fruit species is given in Table 1. Eco-physiological and biophysical studies on tropical and subtropical fruit species are crucial for understanding how LTS affects fruit crops and how it might be managed. (Barlow et al., 1974). Tropical and subtropical fruit species exhibit distinct damaging symptoms when exposed to LT below 10-15°C for a certain period, resulting in drastic yield reduction or even death over prolonged exposure (Chen and Patterson, 1985; Herner, 1990; Alonso et al., 1997). The ability to withstand mild non-freezing temperature stress has evolved over thousands of years in temperate fruits such as apple, pear, peach, plum, strawberry, etc. The coping mechanism used by temperate fruits to adjust to cold conditions comprises alterations in a number of physio-morphological and biochemical parameters as well as the expression of stress-associated genes (Medina et al., 2011).

Temperate fruit crops undergo dormancy during the winter to avoid the detrimental consequences of LTS. Different physiobiochemical changes are brought on by LTS exposure in tolerant and sensitive genotypes. The extent of changes decides the genotype's tolerance and ability to acclimate to the cold stress (Barlow et al., 1974; Martin and Douglas, 1979). The aim of this review is to understand physio-morphological, biochemical, and molecular processes connected to LTS in fruit crops as well as the role of phytohormones and other signaling molecules in reducing low-temperature damage. A better understanding of the physiological mechanisms underlying LTS tolerance in fruit crops will relate to improved crop damage management and breeding strategies to combat LTS.

Physiological responses

LTS significantly affects the vegetative and reproductive growth of tropical and subtropical fruits (Alonso et al., 1997), thereby supressing their yield. For instance, if the temperature falls <4°C in the winter for over three hours, the papaya plants may die due to the leakage of white (oozing out milky) latex from the stem of frost-damaged plants (Ram, 2005). Frost or unexpected temperature fluctuations during late winter cause severe damage to fruit crop foliage, crown, flowers, and fruits resulting in a limited yield of low-quality papaya fruit, which ripens unevenly (Singh et al., 2010). Moreover, under chilling temperatures and humid conditions, a hermaphrodite flower of papaya may revert to femaleness (carpelloidy of stamens), resulting in deformed fruits (Awada, 1958; Storey, 1969; Ram, 2005; Lin et al., 2016). In coffee plants, temperatures <16°C suppress vegetative growth, limiting net photosynthesis, resulting in irregular maturity and poor yields (Bauer et al., 1985). However, the reduction in vegetative growth was not due to the decrease in leaf water potential but it was attributed to more prolonged exposure to LT (<16°C exposure) (Barros et al., 1997). However, during LTS acclimation, relative water content (RWC) of the leaves in guava was associated with decrease in vegetative growth. The non-acclimated guava leaves had three to four time's higher anthocyanin accumulation than the latter cold-acclimated guava leaves (Hao et al., 2009). Similarly, Pradhan et al. (2017) reported that under the LT regimes,

TABLE 1 Optimum temperature ranges for the growth of some tropical and subtropical fruit species.

Sl. No.	Fruit crop	Optimum temperature (°C)	Reference (s)			
01	Grapevine (Vitis vinifera)	10-35°C	White et al. (2006)			
02	Banana (Musa spp.)	20-35°C	Viktorova (1983)			
03	Guava (Psidium guajava)	23-28°C	Verheij and Coronel (1992)			
04	Mango (Mangifera indica)	24-27°C	Davanport (2009)			
05	Logan (Dimocarpus longan)	20-25°C	Verheij and Coronel (1992)			
06	Rambutan (Nephelium lappaceum)	25-32°C	Verheij and Coronel (1992)			
07	Jackfruit (Artocarpus heterophyllus)	16-28°C	Haq (2006)			
08	Litchi (Litchi chinensis)	25-35°C	Tindall (1994)			
09	Durian (Durio zibethinus)	24-30°C	Verheij and Coronel (1992)			
10	Langsat (Lansium domesticum L.)	25-35°C	UDPSM (2002)			
11	Pomelo (Citrus maxima L.)	23-30°C	Verheij and Coronel (1992)			
12	Cashew nut (Anacardium occidentale)	20-35°C	Joubert and Thomas (1965)			
13	Coconut (Cocos nucifera)	20-32°C	Louis and Annappan (1980)			
14	Mangosteen (Garcinia mangostana L.)	25-35°C	Osman and Milan (2006)			
15	Citrus (Citrus spp.)	23-27°C	Verheij and Coronel (1992)			

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RWC and fresh leaf weight in papaya decreased, while dry leaf weight reflected the opposite trend. Pradhan et al. (2018) found that the percent change in plant height in control papaya plants was more substantial than its LT-treated counterpart. The decrease in plant growth was due to the reduced photosynthetic rate under LT. In addition, chilling stress in plant roots can alter their metabolic heat rates as detected by micro calorimetry (Criddle et al., 1988), cause cortical destruction and reduce root elongation (Harrington and Kihara, 1960).

Cellular changes

Several studies have found that cell membrane networks are the primary sites of freezing injury in plants (Levitt, 1980; Steponkus, 1984), and that freeze-induced membrane damage is caused primarily by the acute dehydration caused by freezing (Steponkus, 1984; Steponkus, 1993). The extracellular fluids of the apoplastic space contain a lower solute concentration than the intracellular fluid and thus have a higher freezing point; therefore, ice formation is initiated first in the apoplastic space (Jan and Andrabi, 2009). Since ice has a lower water potential than liquid, extracellular ice has a lower water potential than inside the cell, resulting in dehydration. LTS causes some cellular function abnormalities, and membrane damage and the LTS injury is evaluated through electrolyte leakage, changes in membrane lipid composition, and malondialdehyde (MDA) production. Cell dehydration leads to lipid peroxidation and an increase in MDA content, as well as damage to cell membrane fluidity, thus disrupting membrane selectivity, resulting in the permeability of unwanted nutrient elements and ions, leading to ion leakage and disrupting cellular ionic homeostasis (Mahajan and Tuteja, 2005; Yadav, 2010; Shin et al., 2018). The membrane stability index (MSI) of papaya genotypes exposed to LT regimes decreased (Pradhan et al., 2019) while the membrane injury index (MII) increased (Pradhan et al., 2017). Cold stress reduced the MSI gradually, indicating a loss of cell membrane integrity. Campos et al. (2003) reported that electrolyte leakage and lipid degradation influence cold sensitivity in Coffea sp. leaves.

Numerous studies have noted that chilling stress significantly reduces the photosynthetic efficiency of sensitive plants (Yang et al., 2005; Fariduddin et al., 2011). The disruption of the thylakoid ETS pathway and carbon reduction cycle significantly altered photosynthesis under LTS. Furthermore, stomatal control of CO_2 supply may be primarily responsible for decreasing the net photosynthetic rate (Allen and Ort, 2001). Limiting stomatal conductance may be attributed to loss of turgor of guard cells due to LTS induced dehydration. The ultrastructure of chloroplasts has changed after a prolonged cooling period, and thus ability to intercept light energy may be lost (Yang et al., 2005). Potassium (K) and calcium (Ca) were two essential nutrients in improving plant chilling tolerance. Several studies have revealed that electrolyte leakage is primarily related to K^+ efflux from plant cells, mediated by plasma membrane cation conductance. When K+ levels are low, photo-oxidative damage caused by chilling or frost is worsened. This makes plant growth and yield decrease.

High cellular potassium levels can protect against oxidative damage caused by cooling or frost (Waraich et al., 2012). Furthermore, several studies have shown that applying a higher concentration of K+ could reduce LTS-induced damage in crops such as potatoes (Grewal and Singh, 1980), carnations (Kafkafi, 1990), and vegetable seedlings (Hakerlerler et al., 1997). Pradhan et al. (2018) also reported a high level of K⁺ content with a decline in stomatal conductance and transpiration rate, which would help to maintain plant water status. Recently, Maurya et al. (2020) also noted an increase in leaf K⁺ content in papaya seedlings grown under LTS. Similarly, Ca also influences both development and responses to environmental challenges like cold stress by regulating various physiological processes in plants at the tissue, cellular, and molecular levels (Waraich et al., 2011). Plant genotypes that are LTS tolerant can maintain their high leaf water potential by closing their stomata and limiting water loss during transpiration (Wilkinson et al., 2001). However, Yadav (2010) reported that dehydration occurs under exposure to LTS due to low water uptake due to stomata closure.

Furthermore, Ca also modulates the stress responses during cold injury, healing, and adaptation to cold stress (Palta, 1990). LTS activates Ca²⁺channels, resulting in the accumulation of Ca² ⁺ in the cytosol and the induction of phospholipid signaling. The vacuole is the intracellular source of Ca²⁺, and which triggers stomatal closure with increasing Ca²⁺ levels in the cell, a distinguishing feature of plants grown at or exposed to LTS. According to some studies, Ca2+ (released from internal guard cell stores or the apoplast) regulates ABA-induced stomatal closure (Wilkinson et al., 2001). Ca2+ is required for LTS recovery because it activates the plasma membrane enzyme ATPase, which is needed to pump nutrients lost during the cell injury back into the cell (Palta, 1990). Calcium also acts as calmodulin under LTS conditions, regulating plant metabolic activity and promoting plant development (Waraich et al., 2012). Compared with different Carica papaya genotypes, Pradhan et al. (2018) found the highest Ca²⁺accumulation in the leaf and roots of a cold-tolerant wild relative of papaya, V. cundinamarcensis.

Chlorophyll

The most critical component of the photosystem is chlorophyll. LTS represses chlorophyll bio-synthesis and accumulation in actively growing leaves (Glaszmann et al., 1990). For example, the chlorophyll content of papaya seedlings at a 20°/10°C (day/night) temperature regime was

10.3389/fpls.2022.1022167

8.16% lower than the control plants at 28°/18°C (day/night). Cold-tolerant genotypes may accumulate higher chlorophyll content under LTS than cold-sensitive lines (Pradhan et al., 2019). An alternative measure for determining the freezing tolerance and accessing freezing damage and cold acclimation responses of leaves is chlorophyll fluorescence (Ehlert and Hincha, 2008). The chlorophyll fluorescence technique has been employed in various other crops to assess the extent of photodamage at LT, including Arabidopsis (Ehlert and Hincha, 2008), soybean (Tambussi et al., 2004), maize (Aroca et al., 2001), etc. Regardless of the plant species investigated, Maxwell and Johnson (2000) found that the value of chlorophyll fluorescence (Fv/Fm) in healthy leaves was close to 0.80. The lower value of chlorophyll fluorescence (Fv/Fm) indicates photoinhibition of PS II reaction canters, disrupted due to LTS. Smillie (1979) observed that the quantum efficiency (QE) of the PS II center (Fv/Fm) in papaya was 0.42 in the winter season, having a temperature range of 6°/17°C (min./max.) and 0.72 in the summer, in a temperature range of 18°/26°C (min./ max.), indicating that LT likely lowered the PSII activity. Perturbance of photosynthetic machinery in Strawberry, also reflected by chlorophyll fluorescence ((Fv/Fm) and LTS significantly reduced the chlorophyll fluorescence (Fv/Fm) value (Zareei et al., 2021). Pradhan et al. (2019) also found reduction in Fv/Fm during LTS. The treated plants expressed a significantly lower Fv/Fm value than the control. The lowest observed values were 0.438 and 0.584 in the temperature regime of 20°/10°C (day/night), followed by 22°/12°C (day/night), respectively. In contrast, the control plants (28°/18°C day/ night) exhibited the highest (0.736) value of Fv/Fm. Classified mango cultivars based on LTS and Fv/Fm responses. An Fv/Fm ratio of less than 0.5 is denoted as chilling sensitive, while those with an Fv/Fm value greater than 0.6 are considered as chilling tolerant cultivars.

Photosynthesis

With the lowering of temperature, the rate of metabolic processes slows steadily and eventually stops under extreme stress (Taiz and Zeiger, 2002). LT affects different dimensions of photosynthesis in fruit crops. Cellular photosynthesis under LTS is substantially inhibited or reduced due to the disruption of all major components, including the thylakoid electron transport system and carbon reduction cycle. After a long cooling period, the chloroplast ultrastructure has altered, and the capacity to efficiently trap light energy may be lost by the thylakoids membranes (Yang et al., 2005). LTS thermodynamically reduces enzymatic activities, such as Calvin cycle enzymes and ROS (Reactive Oxygen Species) scavenging enzyme activities. LT causes a photostatic imbalance in thylakoid membranes, leading to an over-reduction of the electron transport chain and ROS generation at PSII and PSI (Soitamo et al., 2008; Ruelland et al., 2009; Yun et al., 2010). Papaya plants exposed to an LT regime of 20°/10°C (day/night) exhibited a 57.96% reduction in the photosynthesis rate compared to the control plants. This reduction was genotype-dependent and decided their degree of tolerance. The chilling susceptible Red Lady papaya had a drastic reduction (82.10%) compared to the other genotypes (Pradhan et al., 2018). Similarly, Grau and Halloy (1997) also reported a 15% reduction in the photosynthesis rate of papava plants exposed to the LT regime (15°/5°C; day/night; 4 days) as compared to the control (25°/15°C; day/night). LTS also contributes to ROS generation by reducing the leaf gas exchange in fruit crops due to decreased stomatal conductance. The genotypes that can withstand LTS can keep their leaf water potential high by closing their stomata and avoiding water loss through transpiration. (Wilkinson et al., 2001). In mango, reported that LTS was attributed to a rise in stomatal resistance of net CO₂ assimilation rate and a decrease in Rubisco activity, chlorophyll concentrations, etc., with increased activity of the chlorophyll-degrading enzyme chlorophyllase. Similarly, among various papaya genotypes, the cold-tolerant wild relative V. cundinamarcensis exhibited a lower transpiration rate than other susceptible genotypes (Pradhan et al., 2018). Grau and Halloy (1997) also reported that the reduction in stomatal conductance under cold stress was up to 25-30% compared to the control in papaya.

Biochemical responses

The biochemistry of various cellular components and reactions changed during LTS to help the plant acclimatize to the cold. LTS induced changes range from changes in membrane lipids composition to cellular metabolites, disruption in the equilibrium of gel to liquid-crystalline phase, synthesis of diverse cryoprotective molecules *viz.* soluble sugars (raffinose, saccharose, stachyose, trehalose), sugar alcohols (inositol, ribitol, sorbitol) and low-molecular-weight nitrogenous compounds (glycine betaine, proline) (Janska et al., 2009).

Cell membrane

Phospholipids are the fundamental components of the plasma membrane, and their composition and properties give membrane fluidity and selectivity. The cell membrane is the first place affected by LTS damage, which results in alterations in the plasma membrane's lipid composition. Compared to nonacclimated plants, the amount of unsaturated fatty acids in the cellular membranes of acclimated plants increases, minimizing injury caused by threshold temperature (Theocharis et al., 2012). These modifications protect the plasma membrane and chloroplast envelope from LTS injury (Matteucci et al., 2011). However, in susceptible genotypes, the proportion of saturated fatty acids in the phosphatidyl glycerol (PG) backbone of the chloroplast membrane was higher under the LTS. (Yokoi et al., 1998).

The ROS produced under LTS due to photostatic imbalance induced lipid peroxidation in the cell, damaging the cell membrane's structural and dynamic features, and hence responsible for the non-functionality of the membrane by modifying the fluidity, ion leakage, and producing malondialdehyde (MDA). For example, lipid order in a phospholipid bilayer of a cell membrane diminishes when lipid peroxidation products are added. Pore formation might occur if all phospholipids are oxidized. Reactive species such as reactive oxygen and nitrogen species (RONS) will be able to cause oxidative damage to intracellular macromolecules like DNA and proteins, too (Alonso et al., 1997; Van der Paal et al., 2016). Coffee seedlings, while exposed for six days at the different treatments of temperatures (10°, 15°, 20°, and 25°C), a higher amount of MDA is generated under low temperatures (10°C), leading to higher electrolyte leakage from the root tip. Under completely controlled conditions. Pradhan et al. (2019) evaluate the effects of five LT regimes on five Carica papaya genotypes and one distantly related cold-tolerant plant, V. cundinamarcensis, They observed lower MDA content in the cold-tolerant papaya genotype and V. cundinamarcensis than in the other C. papaya genotypes. The higher rate of oxidation of membrane lipids in susceptible genotypes was linked to a higher amount of MDA, which caused damage to the membrane.

LTS related proteins

LTS affects a variety of metabolites and metabolic processes, including lipoproteins. Cryoprotection, which is the change of lipid biosynthetic enzymes during LTS (Miura and Furumoto, 2013), keeps the plasma membrane, chloroplastic envelope, and other cellular membranes safe and stable. Plant cells synthesize the specific proteins under the LTS, primarily hydrophilic in nature. Cold-regulated proteins (CORs), Low-Temperature Induced (LTI), Late Embryogenesis Abundant (LEA), Responsive to Abscisic acid (RAB), Early Responsive to Dehydration (ERD), Cold Induced (KIN), heat-shock proteins (HSPs), etc. are examples of such protein families. Dehydrins are the most common LEA proteins during cold acclimation and stabilize cell membranes against FI (Bies-Etheve et al., 2008; Sun et al., 2013). Hao et al. (2009) performed the leaf protein analyses after cold acclimation in guava and observed the accumulation of dehydrin protein in FI tolerant guava variety (Lucknow-49). Several freeze-induced membrane injuries have been associated with ice nucleation in the apoplast initiated by freezing. It has also been reported that denaturation of proteins occurs in plants under LT, potentially resulting in cellular damage (Guy et al., 1985). For instance, Pradhan et al. (2017) noted a 35.51% higher total soluble protein content in the cold-treated papaya leaves

than in the non-treated plants. The cold-tolerant wild relative, *V. cundinamarcensis*, reported the highest increase (79.06%), while the lowest was in the cold susceptible variety, Pusa Nanha (8.78%).

Antioxidant

Low temperature induced membrane injuries also accelerate generation of reactive oxygen species (ROS), and enhanced antioxidants activities plays key role in cold acclimation. (Arora, 2018). Antioxidant enzymes influence plant abiotic stress tolerance as they play a crucial role in balancing the ROS concentration within the cell. The antioxidant defense system of plants is comprised of non-enzymatic antioxidants such as glutathione, vitamin C (ascorbic acid), phenolic compounds, flavonoids, carotenoids, α -tocopherol, the osmolyte proline, and antioxidant enzymes such as CAT (catalase), PPO (polyphenoloxidase), SOD (superoxide dismutase), POD (peroxidase), GR (glutathione reductase), DHAR (dehydro ascorbate reductase), MDHAR (monodehydro ascorbate reductase), and APOX (ascorbate peroxidase) (Cao et al., 2009; Kumar and Yadav, 2009; Zhang et al., 2009 and Ding and Wang, 2018). Geng et al. (2019) reported that overexpression of antioxidant enzymes such as CAT, POX, and SOD under cold stress dramatically lowers the ROS levels in Citrus grandis. Lee and Lee (2000) also reported increasing antioxidant enzyme activity in cucumbers under chilling stress. The LT induces CAT and SOD, which act as antioxidant system and are integral parts of the plant's defensive responses under LTS (Figure 2). It is evident that V. cundinamarcensis, a cold-tolerant wild relative of papaya, also expressed higher SOD, GPX, APX, and GR (Pradhan et al., 2017). However, SOD and APX activities increased in cultivated papaya regardless of genotype under LT conditions (Maurya et al., 2020). In strawberries, the increased levels of H2O2 and MDA are associated with decreased activity of SOD, POD, and CAT under LTS (Xu et al., 2020).

Non enzymatic antioxidants also play a crucial role in ROS balancing during abiotic stresses, including LTS (Ahmad et al., 2010). For instance, higher lycopene containing peels of grapes show better ROS scavenging, and higher chilling tolerance, without significant effect of enzymatic antioxidants activities (Lado et al., 2016). Though, Rooy et al. (2017) also observed cold tolerance of grapevines was associated with both, carotenoids as well as enzymatic antioxidants activities. In cold sensitive, mandarin, chilling injury was associated with low carotenoid content (Rey et al., 2020). However, effective antioxidant defense against ROS, is dependent upon the coordinate action of enzymatic and non-enzymatic antioxidants (Kerchev and Van Breusegem, 2022). For instance, activities of GR, DHAR, MDHAR and APX regulate glutathione-ascorbic acid (GSH/ AA) cycle (Fover and Noctor, 2011) and thus exogenous application of compounds which boosts the plants



antioxidants defense, ameliorate the chilling injury in fruits (Zhang et al., 2021).

Cryoprotectant

Accumulation of compatible osmolytes or osmoprotectants in the cytoplasm is a typical response under various abiotic stresses, where dehydration is involved, including LTS. For instance, plants exposed to chilling stress accumulated more fructans and had better freezing tolerance (Livingston et al., 2009). They safeguard the structure of proteins and enzymes; help to maintain the osmotic gradient between the cell's surroundings and the cytoplasm (Kempf and Bremer, 1998; Fedotova, 2019). Under cold stress, the organic osmolytes like sugars, proline, glycine betaine, amino acids, etc., change the cell's osmotic potential. This change narrows the water potential gradient between the apoplastic region of ice generation and the cytoplasm. As a result, the pace at which water moves away from the cell slows down, and the cell membrane becomes more stable, making it more resistant to LTS stress (Elbein et al., 2003). The role of osmoprotectants in ameliorating LTS is well known in fruit crops. Red Delicious apple's cortical tissues and buds' cold hardiness were associated with the concentration of total sugars, sorbitol, and RFO (raffinose family oligosaccharides) (Stushnoff et al., 1993). Another compatible solute, trehalose (α-D-glucopyranosyl-1, 1-α-D- glucopyranoside), a nonreducing disaccharide, is found in many organisms, including bacteria, fungi, insects, and plants. One of the crucial functions of trehalose is to maintain biological membranes and proteins through H-bonding during dehydration due to FI (Donnamaria et al., 1994; Goddijn and Van Dun, 1999; Elbein et al., 2003). Plants exposed to chilling stress accumulated more fructans and had better freezing tolerance (Livingston et al., 2009).

Proline, a type of amino acid, is found abundantly in higher plants and is known to accumulate in considerable amounts in response to environmental challenges (Hayat et al., 2012; Jogawat, 2019). Proline helps maintain osmotic balance and helps in scavenging free radicals (Figure 2) and their crosstalk. It also serves to preserve the integrity of subcellular structures (e.g., membranes and proteins), limits electrolyte leakage, and buffers cellular redox potential during stress by bringing ROS concentrations into normal ranges (Ozturk and Demir, 2002; Hsu et al., 2003; Kishor et al., 2005). There is a significant correlation between cold stress tolerance and increased proline content in various plants (Swaaij et al., 1985; Dorffling et al., 1990; Kaplan et al., 2007). Kushad and Yelenosky (1987) observed the effect of proline and polyamines in LTS tolerance in three citrus genotypes: sour orange (SO), Citrus aurantium L., 'Valencia' (VAL), Citrus sinensis (L.) Osbeck, and rough lemon (RL), Citrus jambhiri Lush. After three weeks of exposure to cold hardening (temperature range: 15.6°/4.4°C day/night) and nonhardening (temperature range: 32.2°/21.1°C day/night), Sour orange had the highest proline content (7.9mg/g dry weight), whereas the lowest content (4.2 and 3.9mg/g dry weight) was found in VAL and RL, respectively. According to the authors,

there is a direct association between higher spermidine (Spd) content and citrus cold hardiness. Pradhan et al. (2019) also found a significant proline accumulation in papaya genotypes as temperatures decrease.

Strawberry (Fragaria x ananassa Duch.) crown sugar content was also positively correlated with cold tolerance (Paquin et al., 1989). The amount of sucrose decreased, and total sugar content in the crowns of Bounty and Redcoat cultivars increased during cold hardening and peaking in January, coinciding with maximum hardiness. The high glucose levels, fructose, raffinose, and stachyose content are attributed to cold hardiness in Chardonnay and Riesling grape (Vitis vinifera) buds and cortical tissues (Hamman et al., 1996). An endophytic gram-negative bacteria (Burkholderia phytofirmans) altered carbohydrate metabolism in the grapevine (V. vinifera) and helped to acclimatize to LTS (Fernandez et al., 2012). Most sugars, such as mannose, glucose, fructose, sucrose, galactinol, raffinose, and maltose, were higher when grapevine was treated with B. phytofirmans under LTS. Yooyongwech et al. (2009) found that the high-chill peach cv. Kansuke Hakuto had a higher total soluble sugar content in the bud cushion sample during December and January (temperature about 2°C), but the low-chill peach cv. Coral had a lower total soluble sugar content. Pradhan et al. (2019) found that cold (20°/10°C; day/night) exposed papaya plants had 117.10 percent higher total soluble sugars than control plants. Polyamines, such as putrescine (Put), spermidine spermidine: (Spd), and spermine (Spm), play a significant role in the green mango (Mangifera indica L. 'Kensington Pride') LT tolerance during storage (Nair and Singh, 2004). Similarly, membrane integrity and fluidity were protected from LTS, in pomegranates due to higher polyamine levels during storage (Mirdehghan et al., 2007). In addition, experimental evidence suggests that polyamines also play a role in protecting photosynthetic function (Figure 2). Polyamines interact with proteins present in the thylakoid membranes, particularly the light-harvesting complex II (LHCII) and the PSII, through hydrogen bonding and stabilize the tertiary structure of proteins under stress (Kotzabasis et al., 1993; Hamdani et al., 2011). Glycine betaine is an ammonium molecule synthesized in chloroplasts that accumulates under stress conditions, including high salt levels and cold temperatures (Sakamoto and Murata, 2002). In four weeks of cold-acclimation treatment of strawberry (Fragaria x ananassa Duch.), LTS tolerance was raised from -5.8° to -17°C with a twofold increase in endogenous glycine betaine levels in the leaves (Rajashekar et al., 1999). Exogenous glycine betaine (2 mM) to unhardened plants boosted cold leaf tolerance nearly two-fold within 72 h. Furthermore, it promoted whole-plant freezing survival and regeneration. Sirooeinejad et al. (2020) investigated the level of freezing tolerance of four pomegranate (Punica granatum L.) varieties, including 'Alak Torsh' (AT), 'Tabestaneh Torsh' (TT), 'Poostsefid Torsh' (PT), and 'Poostsyah Shirin' (PS), and its association with several physiological events and antioxidant response during cold acclimation and de-acclimation. The lethal temperature (LT_{50}) of pomegranate shoots was positively associated with MDA, starch, and antioxidant enzyme activity but negatively related to total phenolic content, proline, and soluble carbohydrates. Similarly, Pradhan et al. (2019) also noted a significant positive correlation of LT exposed papaya plantlet survival (%) with relative water content (RWC), membrane stability index (MSI), and chlorophyll fluorescence, while noting a significant negative association with MDA, proline, and sugar contents.

Aquaporins

Aquaporins are membrane-bound water channels that facilitate water flow with small solutes across the membrane (Moshelion et al., 2015). Plant aquaporins are present in the plasma membrane and other cellular membranes like the tonoplast, endoplasmic reticulum, plastids, and membrane compartments that interact with symbiotic organisms in some species (Baiges et al., 2002; Maurel et al., 2015). Aquaporins are classified into four clades: AQP1, AQP2, AQP3, and AQP4, of a broad superfamily of major intrinsic proteins (MIPs). Various reports show that aquaporins play a significant role in imparting abiotic stress tolerance, including LTS tolerance (Yooyongwech et al., 2009; Lee et al., 2012; Li et al., 2015; Wei et al., 2019) in various crops. Several reports have suggested that aquaporins play a significant role in the cold tolerance of temperate fruits during winter dormancy. Yooyongwech et al. (2009) reported that gamma tonoplast intrinsic protein ($Pp-\gamma TIP1$) and plasma membrane intrinsic protein (Pp-PIP1) play essential roles in intra- and intercellular membrane transport, enhancing cold resistance in the bud cushions of high-chill peach cultivars.

Among the tropical fruits, the role of aquaporins was studied in cold tolerance in bananas. According to Sreedharan et al. (2013), transgenic banana plants that overexpressed a natural plasma membrane aquaporin (MusaPIP1;2) showed remarkable tolerance to cold stress. In addition, transgenic lines showed lower MDA accumulation, higher proline content, RWC, and increased photosynthetic efficiency (Fv/Fm) than controls under exposure to cold stress. Moreover, these transgenic plants exhibited lower damage to the cell membrane and chloroplastic membrane under LTS. Cold-tolerance of banana Dajiao, also associated with aquaporins viz. MaPIP1;1, MaPIP1;2, MaPIP2;4, MaPIP2;6, MaTIP1;3, and membrane bounds Peroxidase 52 and Peroxidase P7. These proteins are the vital cellular adaptations that led to a reduction in lipid peroxidation and maintained leaf cell water potential (WP) (Figure 2) (He et al., 2018). Xu et al. (2020) reported that the overexpression of MaPIP2-7, an aquaporin gene in bananas, improved tolerance to various stresses, including cold, salt, and drought. When compared to wild-type (WT) banana plants under stress, *MaPIP2-7* transgenic plants had reduced levels of MDA and ion leakage (IL) but higher levels of chlorophyll content, proline, soluble sugar, and abscisic acid (ABA). Furthermore, Xu et al. (2021) determined the function of the aquaporin gene MaPIP1;1 and confirmed that overexpression led to multiple stress tolerance in bananas, including drought and cold. Recently, Lopez-Zaplana et al. (2022) elucidated that aquaporins (*PIP2;1*, *PIP2;5* and *PIP2;6*) decreases with the magnitude of different abiotic stresses in melon.

Signaling and molecular responses

Phytohormones

Plant growth-regulating hormones such as abscisic acid (ABA), ethylene, jasmonic acid, and salicylic acid (SA) play a significant role during abiotic stresses, including cold, salinity, drought, light, and heavy metal stresses and they acts as links between stress regulator and responses at cellular, tissue and organ level to external stimuli (Rachappanavar et al., 2022). When Vitis vinifera cv. 'Pinot Gris' was sprayed with ABA @ 400 mg L⁻¹ between veraison and three weeks post veraison, it efficiently accelerated and induced deeper dormancy and increased bud frost tolerance without influencing vine size, yield, or fruit quality (Li et al., 2015). Postharvest foliar treatments of ABA @ 5000 ppm to Merlot grapevines promoted leaf abscission and boosted autumn bud cold tolerance. However, ABA at a lower concentration of 1000 ppm encourages a delay in bud break in the spring season. It has been noted that after ABA treatments, fruit yield and elemental composition were less affected (Bowen et al., 2016). Foliar application of ABA (500 mg· L⁻¹) increased freeze tolerance in grapefruit trees (Wang et al., 2020). In grapevine, ethylene is released under low-temperature stress, which induces the expression of ethylene-responsive factor (VaERF057) and enhances cold tolerance (Sun et al., 2016). Ethylene plays a crucial role in preventing chilling injury in nectarines (Zhou et al., 2001). In Arabidopsis, cold stress reduces the amount of active gibberellin (GA) content and the accumulation of DELLAs, which are negative regulators of GA (Achard et al., 2006; Achard et al., 2016). SA treatment (2 mM) is highly effective in reducing the chilling injury effects, electrolyte leakage, and ascorbic acid loss in the husk of pomegranates (Sayyari et al., 2009). Furthermore, SA treatment alleviates chilling injury in 'Qingnai' plum fruit during postharvest storage (Luo et al., 2011). Exogenous application of methyl jasmonate (MeJA) (10⁻⁴ M) reduced the chilling injury index and ion leakage in Tommy Atkins mangos during storage at 7°C (Gonzalez-Aguilar et al., 2000). Application of MeJA vapor for 20 h at 20°C to 'Kent' mango fruits has enhanced the shelf-life under cold storage with a lower chilling injury score (Gonzalez-Aguilar et al., 2001). Similar reports are also available for guava

(Gonzalez-Aguilar et al., 2004), loquat (Cao et al., 2009), and pomegranate (Sayyari et al., 2011).

Molecular responses

The response of a plant to LTS is to limit or eliminate the negative consequences of LT. It leads to biochemical and physiological changes in the plant cells, like membrane rigidification, lowering of enzyme kinetics, ROS generation, metabolic disequilibrium, etc. Plant species' responses to cold stress vary, and the expression of several stress-associated genes are redirected to alter their metabolism accordingly (Chinnusamy et al., 2010; Guo et al., 2018). Plants can activate a cascade of processes that produce changes in gene expression (Zuther et al., 2019) and, as a result, promote biochemical and physiological adaptations that improve their resistance to low non-freezing temperatures. This process is known as cold acclimatization. Plants follow the adaptive mechanism against cold stress by changing the proteins at the translational level, the composition of the cell membrane, and triggering the ROS scavenging system. In addition, gene expression plays a crucial role in these adaptive mechanisms. The molecular mechanisms of cold stress tolerance and freezing tolerance have been studied in detail in Arabidopsis and winter cereals. It has been shown that temperate crops mainly acquire adaptive tolerance, whereas tropical and subtropical crops are vulnerable to cold stress (Chinnusamy et al., 2010). Many cold-regulated (COR) genes require specific signal transduction pathway activation for acclimatization under LTS (Thomashow, 1998) such as membrane linked receptor proteins, and the most extensively studied pathway: ICE-CBF-COR pathway (Zinta et al., 2022). (Figure 3).

The list of various gene families/gene(s) responsible for LTS tolerance in different fruit crops is listed in Table 2. The COR genes in Arabidopsis plants are regulated by CRT/DRE binding factors (CBF) at the transcription level to enhance the freezing tolerance (Jaglo-Ottosen et al., 1998; Gilmour et al., 2000). CBF related genes, namely, CBF1, CBF2, and CBF3, are rapidly and transiently expressed in response to LT (Gilmour et al., 1998). In the transgenic Arabidopsis plants, susceptible to cold stress, it has been found that ectopic synthesis of CBF proteins leads to constitutive expression of COR genes, resulting in enhanced freezing tolerance (Jaglo-Ottosen et al., 1998; Gilmour et al., 2000). The inducer of C-repeat binding factor (CBF) expression 1 (ICE1), part of the CBF transcriptional cascade, regulates transcription (Zhu et al., 2007). In addition, the activation of transcriptional cascades involving CBF is mediated by Ca²⁺ signaling (Figure 3). Under cold stress, the function of ICE1 is to induce the constitutive expression of CBFs (Chinnusamy et al., 2007). PIF4 (Phytochrome Interacting Factor 4) and PIF7 (Phytochrome Interacting Factor 7) negatively regulate CBFs during the long day. On the other hand, when days are



short, PIF4 and PIF7 are turned off, which makes plants more resistant to freezing. However, the actual mechanisms of photoperiod-induced repression of *CBFs*, *PIF4* and *PIF7*, are still not very evident; therefore, it may not be the sole mechanism engaged in acclimation processes (Fennell, 2014).

Despite this, the CBF/DREB1 dependent low-temperature signaling network is the most crucial regulatory system (Chinnusamy et al., 2007). Maurya et al. (2020) investigated the similarity of the putative amino acid sequence of CBF1 of Carica papaya through a BLAST search. They reported higher similarities with Quercus suber (53%), Malus domestica (50%), Durio zibethinus (49%), Theobroma cacao (49%), Ziziphus jujube (48%) and Rosa chinensis (48%) (Table 3). Furthermore, the phylogenetic tree revealed the evolutionary link between the CBFs of different plants (Figure 3). The phylogenetic tree of CBFs was classified into three major groups, and C. papaya CBF1/DREB belonged to group-II, as presented in Figure 4. The comparison of amino acid sequence and multiple sequence alignment of Carica papaya's CBF1/ DREB protein and other plant species was shown in Figure 5. Dhekney et al. (2007) found cold-inducible regions and CBF sequences in C. papaya and Vasconcellea and reported sequence homology with the Arabidopsis genome. Recently, Maurya et al. (2020) studied the comparative expression profile of CBF1 and CBF2 gene transcripts under LTS and observed that the CBF1 expression profiles was up-regulated while CBF2 was downregulated in six C. papaya genotypes (Red Lady, Pusa Nanha, P-

7-2, P-7-9, P-7-14, and V. cundinamarcensis). In addition, they noted that V. cundinamarcensis had the highest expression profile of CBF1, followed by P-7-9, whereas the lowest level was found in Red Lady. On the other hand, the CBF2 expression profile followed the opposite pattern. According to Zhu et al. (2007), high-temperature stress (HTS) had minimal influence on the CpCBF2 expression profile, whereas LT inhibited it. The induction of CBF1/DREB1B and CBF3/DREB1A is simultaneous and before CBF2/DREB1C under cold stress (Medina et al., 2011). Fursova et al. (2009) found that overexpression of ICE2 increased the expression profile of CBF1/DREB1B and improved freezing tolerance in Arabidopsis. Takuhara et al. (2011) found that the grapevine cold-induced transcription factors (VvCBF2, VvCBF4, VvCBFL, VvZFPL) also enhance LT tolerance in transgenic Arabidopsis plants. Transgenic strawberry leaves over expressing wheat acidic dehydrin gene (Wcor410a) improved their cold tolerance when exposed to 5°C (Houde et al., 2004). In addition, Hara et al. (2004) reported that a dehydrin gene, i.e., CuCOR19, from Citrus unshiu, can operate as a hydroxyl radical scavenger under LT and found more tolerance to freezing when over expressed in Nicotiana tabacum (Hara et al., 2003). Apart from this, they also reported a negative association between the accumulation of CuCOR19 and electrolyte ion leakage. The approximate size of peach's dehydrin gene (ppdhn1) is 50 kDa, which encodes PCA60 protein, consisting of 472 amino acids. In the deciduous genotype, the protein accumulates earlier and is retained at

Sl. No.	Fruit crop	Gene (s)	Reference (s) Xiao et al. (2008)		
01	Grapevine (Vitis vinifera)	VvCBF1, VvCBF2, VvCBF3, and VvCBF4			
02	Apple (Malus x domestica)	MdCBF1, MdCBF2, MdCBF3, MdCBF4 and MdCBF5	Feng et al. (2012)		
03	Almond (Prunus dulcis)	PdCBF1 and PdCBF2	Barros et al. (2012)		
04	Strawberry (Fragaria × ananassa)	FaCBF1	Owens et al. (2002)		
05	Crab apple (Malus baccata)	DREB1/CBF(Mb-DREB1)	Yang et al. (2011)		
06	Sour cherry (Prunus cerasus)	PcCBF1	Owens et al. (2002)		
07	Sweet cherry (Prunus avium)	PaCBF	Kitashiba et al. (2004)		
08	Blueberry (Vaccinium corymbosum)	BB-CBF	Polashock et al. (2010)		
09	Vitis riparia	VrCBF4 and VrCBF1a	Xiao et al. (2008)		
10	Poncirus trifoliata	COR11, COR19	Thomashow (2010)		
11	Peach (Prunus persica)	ppdhn1	Artlip et al. (1997)		
12	Citrus unshiu	Cu-COR19	Hara et al. (2004)		
13	Banana (Musa spp.)	MusaDHN-1	Upendra et al. (2011)		
14	Apple (M. domestica)	Dehydrin (MdDHN)	Liang et al. (2012)		
15	Grape (V. vinifera)	Dehydrin (DHN)	Yang et al. (2012).		
16	Banana (Musa spp.)	MaCslD4, MaCslA4/12, MaCslE2	Yuan et al. (2021)		

TABLE 2 Various gene (s) responsible for LTS tolerance in different fruit crops.

TABLE 3 Percent identity matrix of Carica papaya CBF1/DREB with other plants CBF/DREB factors (Table adapted from Maurya et al., 2020).

CBF1/DREB Carica papaya	100.00															
DREB 1B Juglans regia	47.55	100.00														
DREB 1F Quercus suber	53.16	59.67	100.00													
DREB 1F Durio zibethinus	49.48	59.69	55.00	100.00												
DREB 1F Gossypium arboretum	46.45	60.10	61.83	67.51	100.00											
DREB 1B Ziziphus jujube	48.80	59.02	61.58	66.50	82.27	100.00										
DREB 1B Theobroma cacao	49.50	47.06	53.63	50.00	45.59	43.78	100.00									
DREB 1B Cinnamomum micranthum	45.93	47.37	51.89	49.48	46.95	46.67	66.82	100.00								
DREB 1B Ricinus communis	45.58	51.44	54.79	52.82	48.61	48.84	52.68	51.42	100.00							
CBF6 Populus trichocarpa	47.09	51.00	52.91	51.05	53.02	53.70	44.72	43.27	50.00	100.00						
CBF like Vitis riaparia	45.60	51.91	58.60	51.18	56.52	54.89	43.24	45.79	50.54	53.59	100.00					
CBF/DREB Rosa chinensis	48.24	47.03	55.37	52.94	54.95	56.28	48.06	45.02	47.29	57.00	55.32	100.00				
A)2/ERF Artemisia annua	46.94	53.54	54.70	58.51	57.58	56.63	46.67	46.73	52.76	56.10	62.36	56.37	100.00			
CRT binding factor 3 Solanum tuberosum	46.63	55.88	54.59	52.36	56.87	56.87	44.66	45.50	49.53	53.49	62.23	53.85	62.44	100.00		
AP2/ERF Trema orientale	46.12	55.88	62.78	56.08	58.17	59.71	45.15	44.08	52.86	57.77	58.42	55.98	61.93	61.86	100.00	
DREB 1D Malus domestica	50.54	55.88	61.18	57.14	59.36	63.04	47.85	45.79	52.38	58.20	58.38	62.77	59.79	61.78	71.05	100.00



Ziziphus jujube DREB 1B Like (XP_015901948.1), Theobroma cacao DREB 1B Like (EOY08142.1), Cinnamomum micranthus DREB 1B Like (RWR82367.1), Ricinus communis DREB 1B Like (EEF33807.1), Populus trichocarpa C-repeat binding factor 6 (ABO48367.1), Vitis riparia CBF-Like transcription factor (ARD27023.1), Rosa chinensis putative CBF/DREB transcription factor (ABQ53132.1), Artemisia annua AP2/ERF domain-containing protein (PWA50609), Solanum tuberosum CRT binding factor 3 (ACB45095.1), Trema arientale AP2/ERF domain-containing protein (PON57830.1), Malus domestica DREB 1D (XP_028948628.1) (FIGURE adapted from Maurya et al., 2020).

higher levels for a longer time than in the evergreen genotype, indicating that it plays a role in cold tolerance (Artlip et al., 1997). Furthermore, the PCA60 protein acts as an anti-freeze protein that modifies ice crystal growth in the cell and is present in the cytosol and nucleus but absent in the vacuole when investigated under the light electron microscope (Wisniewski et al., 1999). LT tolerance is increased when MpRCI, a cold-tolerance-related plasma membrane protein gene of the 'Dajiao banana cultivar, is overexpressed in transgenic tobacco. (Feng et al., 2009).

Overexpression of the *MbDREB1* transcription factor improved cold tolerance and other abiotic stresses in apples (Yang et al., 2011). Additionally, *Prunus persica CBF1* gene overexpression significantly enhanced freezing tolerance in transgenic apples (Wisniewski et al., 2011). Feng et al. (2012) studied the adaptive responses to cold tolerance in apples and discovered that the transcription factor MdCIbHLH1 plays an important role in the LTS.

Bananas are highly susceptible to chilling stress (Chen et al., 2020). Recently, Meng et al. (2020) reported that the accumulation of fasciclin-like arabinogalactan protein (*FLA*) in banana under low temperatures could improve chilling tolerance by facilitating cold signal pathways, which increases the biosynthesis of plant cell wall components. The results provided background information on *MaFLAs* and suggested their involvement in plant chilling tolerance and their potential as candidate genes to be targeted when breeding for cold tolerance. Yuan et al. (2021) reported that the *Csl* (*Cellulose*

synthase-like) gene family imparts tolerance against LT in bananas by synthesizing a special kind of hemicellulose, an essential component of the cell wall, and strengthening the stiffness of the cell wall. The expression of genes *MaCslD4* (*Musa acuminate Cellulose synthase-like gene*), *MaCslA4/12*, and *MaCslE2* (member of the Csl gene family) were found to be significantly higher in the tolerant variety than in the susceptible banana variety under LTS (Yuan et al., 2021).

The commercially important diversity of citrus is available for cultivation. However, it has been observed that most coldacclimated citrus cultivars can only sustain temperature variations of -4° to -10°C. However, Poncirus trifoliata, a citrus relative, is a genetic resource for cold-hardiness that can survive at temperatures as low as -26°C (Yelenosky, 1985). Zhang et al. (2005) studied the gene expression profile of P. trifoliata by mRNA DDRT-PCR (differential display-reverse transcription) and quantitative relative-RT-PCR in a cold acclimation temperature regime. The up-regulated genes showed high similarities at the amino acid level with genes with known functions like water channel protein, betaine or proline transporter, early light-induced protein, nitrate transporter, aldo-ketoreductase, tetratricopeptide-repeat protein, F-box protein, ribosomal (r) protein L15. These cold-acclimation up-regulated genes of P. trifoliata are also regulated by osmotic and photo-oxidative signals in other plants.

Anthocyanins, a water-soluble pigment available across the plant kingdom, have a significant role in mitigating cold stress by increasing their content in the cell and maintaining the osmotic CBF1/DREB Carica papaya DREB 1B Juglens regia DREB 1F Quercus suber DREB 1B Durio zibethinus DREB 1F Gossypium arboretum DREB 1B Ziziphus jujube DREB 1B Theobroma cacao DREB 1B Theobroma cacao DREB 1B Theobroma cacao DREB 1B Thinus communis CBF6 Populus trichocarpa CBF like Vitis riparia CBF/DREB Rosa chinensis AP2/ERF Artemisia annua CRT binding factor 3 Solanum tuberosum AP2/ERF Trema orientale DREB 1D Malus domestica

CBF1/DREB Carica papaya DREB 1B Juglans regla DREB 1F Quercus suber DREB 1B Durio zibethinus DREB 1F Gossypium arboretum DREB 1B Theobroma cacao DREB 1B Theobroma cacao DREB 1B Theobroma cacao DREB 1B Cinnamomum micranthum DREB 1B Ricinus communis CBF6 Populus trichocarpa CBF like Vitis riparia CBF/DREB Rosa chinensis AP2/ERF Artemisia annua CRT binding factor 3 Solanum tuberosum AP2/ERF Trema orientale DREB 1D Malus domestica

CBF1/DREB Carica papaya DREB 1B Juglans regla DREB 1F Quercus suber DREB 1B Durio zibethinus DREB 1F Gossypium arboretum DREB 1B Theobroma cacao DREB 1B Theobroma cacao DREB 1B Theobroma cacao DREB 1B Theobroma cacao CBF6 Populus trichocarpa CBF6 Populus trichocarpa CBF like Vitis riparia CBF/DREB Rosa chinensis AP2/ERF Artemisia annua CRT binding factor 3 Solanum tuberosum AP2/ERF Trema orientale DREB 1D Malus domestica

CBF1/DREB Carica papaya DREB 1B Juglans regia DREB 1F Quercus suber DREB 1B Durio zibethinus DREB 1F Gossypium arboretum DREB 1B Ziziphus jujube DREB 1B Theobroma cacao DREB 1B Theobroma cacao DREB 1B Ricinus communis CBF6 Populus trichocarpa CBF like Vitis riparia CBF/DREB Rosa chinensis AP2/ERF Artemisia annua CRT binding factor 3 Solanum tuberosum AP2/ERF Trema orientale DREB 1D Malus domestica



DDDGND

-VDD

WG--NNTO

MIL

STR

FIGURE 5

Alignment of the predicted amino acid sequences of the *Carica papaya CBF1/DREB* from with another plant *CBFs/DERB*. All the designated sequences showed conserved domain of AP2, the characteristics of c-repeat binding factors. (FIGURE adapted from Maurya et al., 2020).

balance to improve the chilling tolerance (Chalker-Scott, 1999; Hao et al., 2009). Lo Piero et al. (2005) reported an accumulation of anthocyanin in red-orange juice vesicles under LTS. The biosynthesis pathway of anthocyanin is complex and involves several structural genes: *PAL* (phenylalanine ammonia-lyase), *CHS* (chalcone synthase), *DFR* (dihydro flavonol 4-reductase), and *UFGT* (UDP-glucose flavonoid glucosyltransferase), which are strongly triggered during LTS.

TGLESSVWARGWLAKCTYGTGLAPS

According to Crifo et al. (2011), cold stress causes transcriptome changes in blood oranges that trigger enhancement of flavonoid biosynthesis, including the metabolic pathways of anthocyanin biosynthesis. Similarly, in red-skinned grapes, cold stress promoted anthocyanin biosynthesis but did not increase the accumulation of endogenous levels of abscisic acid (Gao-Takai et al., 2019). Using an RNA-seq study of leaves from the 'Gala' apple variety after exposure to LT (16°C), Song et al. (2019) identified additional regulatory genes ROS1 in apple (*M. domestica*) that may be involved in controlling LT-induced anthocyanin production.

Conclusion

LTS unfavorably affects the fruit crop's overall growth and development because it affects almost all aspects of cellular activity and, ultimately the quality production of the crop. However, according to the current review, the method of comprehending low-temperature damage have been adequately tried by researchers on many field crops, while very limited studies have been conducted on fruit crops. These include primarily temperate fruit crops and some tropical crops like bananas. Therefore, new avenues like the role and use of aquaporin genes to alleviate the LTS need to be explored in tropical and sub-tropical fruit crops. Furthermore, under open and simulated low-temperature conditions, more detailed research is required to elucidate the underlying physiobiochemical and molecular mechanisms for LTS tolerance and resistance in fruit crops. These strategies will immensely help in genetic improvement and standardizing cultural practices for successful cultivation of the fruit crops, particularly in the erratic weather conditions of sub-tropical and tropical regions.

Author contributions

AKG, NKM, SG, and KB conceived the idea of the review and prepared the initial outline, and wrote the first draft. AKG

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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