

Plant Abiotic Stress: Function of Nitric Oxide and Hydrogen Peroxide



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Abstract The negative effect of various environmental stresses is partially due to the generation of reactive oxygen species (ROS). ROS were originally considered to be detrimental to cells. However, it is now recognized that hydrogen peroxide functions as a trigger for induction of many genes encoding enzymes involved in cellular protection under stress. In a number of abiotic responses, NO generation occurs in parallel with H₂O₂, and both molecules can act synergistically and/or independently. Studies have shown that NO and H₂O₂ function as stress signals in plants, mediating a range of resistance mechanisms. The main place of the signal perception of worsening environmental conditions is the plasma membrane. On the other hand, one of the major proteins of the plant cell membrane is the plasma membrane H⁺-ATPase (PM H⁺-ATPase), a key enzyme in adaptation of plants to abiotic stresses. In plants exposed to different abiotic stresses, e.g., salinity, heavy metals, and low or high temperature, an increase in permeability related to membrane damage is observed. Maintaining ionic balance and replenishing the loss of essential substances are important issues. Support of active transport of ions through the plasma membrane requires increased generation of an electrochemical proton gradient by PM H⁺-ATPase, which results in a proton-motive force used by active transporters for assimilation of various nutrients, as well as for releasing toxic ions from cells. NO and H₂O₂ are important elements for understanding the mechanisms

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of PM H⁺-ATPase modification at both genetic and posttranslational level. Nowadays the role of NO and H₂O₂ as well as the signal cascades by which signaling molecules participate in plant responses to changing environmental conditions is under intensive study.

Keywords Abiotic stress · NO · H₂O₂ · Plasma membrane H⁺-ATPase

1 Introduction

Plants are continuously subjected to numerous environmental stresses, such as heavy metals, salinity, drought, and low or high temperature. These adverse external factors limit plant growth and productivity. Understanding plant responses to abiotic stresses is essential for reasonable production of more resistant crop plants. The process by which plant cells sense the stress signals and transmit them to cellular machinery for activation of adaptive responses is referred as signal transduction. In plants, the signal transduction pathways are still elucidated. Despite many physiological and molecular studies, the knowledge of how plants sense and transduce extreme temperature, drought, salinity, or heavy metals needs to be explained. The negative effect of various environmental stresses is partially due to the generation of reactive oxygen species (ROS). Accumulation of ROS is a general feature of abiotic stresses and together with changes in cell antioxidant capacity can lead not only to oxidative damage but also to ROS signaling (Tanou et al. 2009; Foyer and Noctor 2009; Groß et al. 2013; Arora et al. 2016). In a number of abiotic responses, H₂O₂ generation occurs in parallel with NO (nitric oxide), and these molecules can act both synergistically and independently (Clarke et al. 2000; Neill et al. 2002; Zhang et al. 2007; Qiao et al. 2014). NO is a key signaling molecule that mediates a variety of physiological functions and defense mechanisms against abiotic stresses in plants (Qiao and Faan 2008).

In general, extreme temperature mainly results in mechanical embarrassment, whereas heavy metals, salinity, and drought exert their scathing effects mainly by disrupting the ionic and osmotic equilibrium of the cell. In plants exposed to abiotic stresses, an increase in permeability related to membrane damage is observed (Guy 1990; Breckle and Kahle 1991; Devi and Prasad 1999). Moreover, under heavy metals or salt stress, accumulation of toxic ions in plant cells occurs. Ion homeostasis is the physiological fundament for living cells. Maintaining ionic balance and replenishing the loss of essential substances in repair processes are important issues under such conditions (Janicka-Russak 2011). Support of active transport of ions and organic compounds through the plasma membrane requires increased generation of a proton gradient by plasma membrane H⁺-ATPase (Serrano 1989). The generation of an electrochemical proton gradient across the membrane results in a proton-motive force that is used by active transporters for uptake of various nutrients as well as for releasing toxic ions and substances from cells (Janicka-Russak 2011). It has been proven that the activity of the plasma membrane H⁺-ATPase (PM H⁺-ATPase)

is a key index of plant adaptation to abiotic stresses (Kłobus and Janicka-Russak 2004; Janicka-Russak et al. 2008, 2012a, b, 2013; Sahu and Shaw 2009; Lopez-Pérez et al. 2009). Moreover, the reported studies show that signaling molecules are important elements for understanding the mechanisms of modification of the plasma membrane proton pump activity. Both NO and H₂O₂ play an essential role in regulation of PM H⁺-ATPase activity under stress conditions (Zhang et al. 2007; Siddiqui et al. 2010; Janicka-Russak and Kabała 2012; Janicka-Russak et al. 2012a; Janicka et al. 2018). The pathways by which signaling molecules participate in plant responses to changes in environmental conditions are under intensive study. The purpose of this chapter is to explain the role of NO and H₂O₂ in plant adaptation to abiotic stresses.

2 Abiotic Stress in Plants

Stress is a physical definition. In biological terms, it is problematic to define stress. A physiological condition, stressful for one plant, may be optimal for another plant. The most applied definition of biological stress is an adverse force or a condition which limits the normal functioning of biological systems such as plants (Jones and Jones 1989). It is well known that NO and H₂O₂ play key functions in the regulation of plant tolerance to environmental stress factors such as drought, salinity, low or high temperature, heavy metals, and wounding (Neil et al. 2002, 2008; Vandenabeele et al. 2003; Shapiro 2005; Suzuki and Mittler 2006; Volkov et al. 2006; Qiao and Faan 2008; Janicka et al. 2018).

Exposure to high salinity can affect all the major plant processes, including protein synthesis, photosynthesis, and lipid metabolism (Parida and Das 2005). A high concentration of salt in the soil is one of the stressogenic stimuli. Excessive salinity imposes two stress factors on plants: an osmotic component that results from the reduced water availability caused by an increase in osmotic pressure in the soil and an ionic stress resulting from a solute imbalance, causing changes in the K⁺/Na⁺ ratio and increasing the concentration of Na⁺ and Cl⁻ in the cytosol (Alvarez et al. 2003). Sodium toxicity results from the similarity of the Na⁺ and K⁺ ions that compete for plant transporters and binding sites in enzymes (Silva and Geros 2009). Osmolytes like proline, glycine-betaine, trehalose, mannitol, and sorbitol that are abundantly produced and accumulated in salt-treated plants represent an important component of salt stress responses. These compounds lower the osmotic potential of cells or protect various cellular structures and proteins against stress-induced damage (Sahi et al. 2006). It is well established that the efficient exclusion of Na⁺ excess from cytoplasm is the most important step toward the maintenance of ion homeostasis inside the cell (Zhu 2003; Silva and Geros 2009). Majority of plants adapted to salinity maintain a relatively low concentration of Na⁺ in the cytosol achieved through the active exclusion of sodium ions in the apoplast and vacuole. Removal of sodium ions out of the cell, catalyzed by the specific plasma membrane and tonoplast Na⁺/H⁺ antiporters, depends on the membrane electrochemical proton gradient (Chinnusamy

et al. 2005; Silva and Geros 2009; Kabała and Janicka-Russak 2012). The only pump which generates proton gradient across the plasma membrane is H^+ -ATPase. For this reason, it is believed that plant plasma membrane H^+ -ATPase plays a major role in salt stress tolerance. The importance of the proton pump in plant adaptation to salinity was indicated by the observations carried out on salt-tolerant plants, showing increased activity of this enzyme in both normal and salt conditions (Vera-Estrella et al. 1994; Chen et al. 2007; Sahu and Shaw 2009). Moreover, the salinity induces the activities of the plasma membrane proton pumps both in halophytes and glycophytes (Kłobus and Janicka-Russak 2004; Sahu and Shaw 2009; Lopez-Pérez et al. 2009; Janicka-Russak et al. 2013). NaCl stress induces the expression of PM H^+ -ATPase genes (Niu et al. 1993; Janicka-Russak and Kłobus 2007; Janicka-Russak et al. 2013). The positive correlation between accumulation of mRNAs of H^+ -ATPase and salt tolerance is well documented (Niu et al. 1993; Perez-Prat et al. 1994; Janicka-Russak and Kłobus 2007; Sahu and Shaw 2009). It seems that both NO and H_2O_2 are generated under salt stress to activate resistance mechanisms in plants. These signal molecules maintain adequate K nutrition and a favorable K^+ to Na^+ ratio in cytosol (Zhu 2003). It was shown that NO produced under salt stress can serve as a second messenger to induce H_2O_2 generation mediated by the induction of plasma membrane NADPH oxidase expression (Zhang et al. 2007). Subsequently, the H_2O_2 contributes to induce the expression of PM H^+ -ATPase that creates an electrochemical proton gradient for the establishment of ionic homeostasis to confer salt resistance (Zhang et al. 2007; Janicka et al. 2018).

Unpredictable falls in temperature are a common phenomenon in a temperate climate. Membranes are most susceptible to damage resulting from low temperature. An increase in membrane permeability and a change in its viscosity and fluidity result in a decrease of cell turgor due to low temperature (Guy 1990). It is well known that lipids containing saturated fatty acids solidify at temperatures higher than those containing unsaturated fatty acids. Therefore, the relative proportion of unsaturated fatty acids in the membrane strongly influences its fluidity (Mahajan and Tuteja 2005). Membrane changes at cold concern positional redistribution of saturated and unsaturated fatty acids. Membrane reconstruction has been observed in response of plants to low temperature. Chilling-sensitive plants usually have a higher proportion of saturated fatty acids. On the other hand, chilling-resistant species are marked by higher proportion of unsaturated fatty acids (Mahajan and Tuteja 2005). Probably, the change in membrane fluidity in sensitive species initiates a signal chain reaction that leads to acclimation to cold stress. Furthermore, the increase in H_2O_2 accumulation at low temperatures occurs (Kocsy et al. 2001). Glutathione is an important component of the ascorbate-glutathione cycle, which is indirectly involved in the regulation of H_2O_2 concentration in plants (Foyer et al. 1997). At low temperature, high GSH content and NADPH-dependent glutathione reductase (GR) activity were detected in plants indicating their possible contribution to chilling tolerance and cold acclimation. Changes in H_2O_2 concentration and GSH/GSSG ratio alter the redox state of the cells and activate specific defense mechanisms through a redox signaling chain (Kocsy et al. 2001). ROS produced in response to low temperature contributes to membrane damage. The integrity of intracellular

organelles is also disrupted, leading to a loss of compartmentalization and impairing of photosynthesis, protein synthesis, and general metabolic processes. Low temperatures induce alteration in cellular components, including unsaturated fatty acids, glycerolipids, proteins, and carbohydrate composition as well as the activation of ion channels (Knight et al. 1996). Cold acclimation results in protection and stabilization of the integrity of cellular membranes, enhancement of the antioxidative mechanisms, increased intracellular sugar level, as well as accumulation of cryoprotectants like polyamines (Krishna et al. 1995; Polisensky and Braam 1996; Xiong and Zhu 2001; Mahajan and Tuteja 2005; Chinnusamy et al. 2006). Cold acclimation involves precise signaling and regulation of the transcriptome. Cold-induced ROS (such as H_2O_2) may activate a mitogen-activated protein kinase cascade that regulates tolerance to freezing (Chinnusamy et al. 2006). Altered transcript levels of PM H^+ -ATPase genes were also observed (Ahn et al. 1999; Janicka-Russak et al. 2012a) due to exposure of plants to low temperature. Published data also indicate that activity of PM H^+ -ATPase is modulated under low temperature by changes in plasma membrane lipids (Martz et al. 2006). The activity of plasma membrane H^+ -ATPase may increase more than twofold following cold acclimation (Martz et al. 2006). Moreover, fatty acid-regulated PM proton pump activity is involved in the cellular response underlying cold acclimation and de-acclimation (Martz et al. 2006).

Temperature is a major factor in determining the natural distribution of plants. High temperatures can damage several plant metabolic and physiological processes (Larkindale and Huang 2004; Johnston et al. 2007). Plants are able to sense, respond to, and acclimate to mild high temperatures. This property allows them to survive extreme temperature shocks. High temperature causes protein dysfunction. Maintaining proteins in their functional conformations and preventing their aggregation are important for cell survival under high temperature stress (Wang et al. 2004; Timperio et al. 2008). The treatment of plants with temperatures near the upper limit for survival induces a specific set of proteins, the heat shock proteins (HSPs). This response is common to all living organisms. Königshofer et al. (2008) observed that as the final indicator of the heat signaling cascade, the synthesis of HSPs occurs. Their results indicated that elevated production of hydrogen peroxide, in the early stages of the heat signaling pathways, which responds rapidly to changes in membrane fluidity, is required for the activation of HSP synthesis. Similar observations were made by Miller and Mittler (2006) and Volkov et al. (2006). It was demonstrated that increased H_2O_2 production in apoplast under abiotic stresses occurs as a result of elevated levels of ABA (Hu et al. 2005, 2006). There is some evidence that ABA may be involved in heat stress responses in plants (Robertson et al. 1994; Gong et al. 2002). It was shown that ABA added to plants protected them from heat-induced oxidative damage (Larkindale and Knight 2002; Larkindale and Huang 2004).

Bioaccumulation of heavy metals in the environment has become a problem for all living organisms including plants. Heavy metals have the competence to interact with several cellular biomolecules such as proteins and DNA. Elevated levels of heavy metals lead to the increased generation of reactive oxygen species and induction of oxidative stress (Howlett and Avery 1997; Pál et al. 2005; Hossain et al. 2012). In response to heavy metals, plants are equipped with a range of mechanisms to counteract

their toxicity. These mechanisms include chelation of metals by forming complexes with phytochelatins or metallothioneins, which is followed by the removal of heavy metals ions from cytoplasm. The plant plasma membrane may be regarded as the first structure that is a target for heavy metal toxicity. An increase in permeability related to membrane damage is observed in plants that have been subjected to heavy metal stress (Demidchik et al. 1997; Murphy and Taiz 1997; Murphy et al. 1999). It is well known that metal ions are easily bound to both the sulfhydryl groups of proteins and hydroxyl part of phospholipids (Devi and Prasad 1999). They can also replace the calcium ions at essential sites of cell membranes (Breckle and Kahle 1991). All these events result in disruption of membrane integrity and ionic homeostasis of cells. A factor that may be important in the maintenance of plasma membrane integrity in the presence of heavy metals could be enhanced membrane repair after damage (Salt et al. 1998). This could involve heat shock proteins. Moreover maintaining ionic balance and replenishing the loss of essential substances are important under heavy metal stress. A few observations have indicated that H^+ -ATPase activity was changed under heavy metal stresses (Demidchik et al. 1997; Astolfi et al. 2003, 2005; Janicka-Russak et al. 2008, 2012b; Kabała et al. 2008). Moreover, the relative expression of PM- H^+ -ATPase genes in cucumber roots was affected due to Cd treatment (Janicka-Russak et al. 2012b).

Many studies indicate that NO and H_2O_2 could be a common and mutual elements of different signaling pathways activated by plants in response to adverse environmental factors. Hydrogen peroxide and nitric oxide are involved in both salt and heat stress tolerance. Pretreatment of seedlings with H_2O_2 or SNP (NO donor) before salt or heat treatment caused decreases in deleterious effects of stress in plants (Uchida et al. 2002). Cross-resistance factors are able to reduce the effects of other stresses. For instance drought-hardened maize plants become more chilling-tolerant as well (Pérez de Juan et al. 1997). These phenomena are common in wide range of stresses (Uchida et al. 2002). This suggests that environmental factors activate similar signal transduction pathways. Stress conditions often lead to similar disturbances in plants, including dehydration and membrane damage. Control of ion movement across plasma membrane is an essential element of cellular adaptation to abiotic stresses. Because active transport of ions across the plasma membrane requires increased generation of a proton gradient by PM H^+ -ATPase, it is suspected that this enzyme could play a key role in abiotic stress tolerance.

3 Plasma Membrane H^+ -ATPase

The main place of the signal perception of worsening environmental conditions is the outer cell membrane, i.e., the plasma membrane. One of the most important proteins of the plant cell membrane is PM H^+ -ATPase. This protein belongs to a large superfamily of pumps termed P-type ATPases. Plasma membrane H^+ -ATPase is a tightly bound and integral transmembrane protein. The enzyme is a single polypeptide of ~100 kDa. By the use of the chemical energy of ATP hydrolysis, PM ATPases extrude protons from cells to apoplast to generate an electrochemical

proton gradient (Serrano 1989). PM H⁺-ATPase as a proton pump plays a central role in many physiological functions such as nutrient uptake, intracellular pH regulation, stomatal opening, and cell growth. Besides regulation of physiological processes, this proton pump also participates in the adaptation of plants to changing environmental conditions, especially stress conditions (Janicka-Russak 2011). Generation of an electrochemical proton gradient across the membrane is necessary for active transporters involved in accumulation of nutrients as well as removal of ions and other toxic substances from cells. Thus, the plasma membrane H⁺-ATPase can be a key element for resistance mechanisms activated under various stress conditions. Therefore, the regulation of enzyme activity may represent an important cellular mechanism for abiotic stress resistance. Modification of the plasma membrane H⁺-ATPase activity could take place at the genetic and posttranslational level.

The molecular study of plant H⁺-ATPase has shown that this enzyme is encoded by a multigene family. In addition to tissue-specific expression, the plasma membrane H⁺-ATPase genes are differentially expressed by environmental factors. Several studies have indicated that the H⁺-ATPase genes might be activated by various abiotic stresses. With such a phenomenon, the amount of H⁺-ATPase might be increased under conditions requiring greater transport activity. The external signals resulting in changes in plant plasma membrane H⁺-ATPase gene expression include salinity (Niu et al. 1993; Janicka-Russak and Kłobus 2007), low temperature (Ahn et al. 1999; Janicka-Russak et al. 2012a), heavy metals (Janicka-Russak et al. 2008, 2012b), and mechanical stress (Oufattole et al. 2000). It was shown that signaling molecules NO and H₂O₂ contribute to increased activity of PM H⁺-ATPase by stimulation of expression of gene encoding this enzyme (Zhang et al. 2007; Janicka et al. 2018).

As to posttranslational regulation, the best known mechanism described to date involves the auto inhibitory action of the C-terminal domain (approximately 100 amino acids) of the enzyme protein. The activity of the enzyme is well known to be regulated by 14-3-3 proteins, the association of which requires phosphorylation of the penultimate H⁺-ATPase residues of Thr 947 (Svennelid et al. 1999; Fuglsang et al. 2006). The binding of 14-3-3 regulatory protein displaces the inhibitory domain, activating the enzyme. One 14-3-3 protein dimer binds two C-terminal polypeptides simultaneously, so a high activity state of H⁺-ATPase could involve formation of dimers or multimeric complexes (Kanczewska et al. 2005). Recent studies have shown that NO participates in posttranslational modifications of plasma membrane proton pump because it leads to an increased level of enzyme phosphorylation and to an increased H⁺/ATP coupling ratio (Janicka et al. 2018).

It is well known that the plant plasma membrane H⁺-ATPase requires lipids for activity. This lipid dependency suggests a possible mode of regulation via modification of lipid environment (Kasamo 2003). Abiotic stresses lead to changes in the plasma membrane lipid composition altering the fluidity of the membrane. The modulation of the phospholipid environment of the plasma membrane regulates the activity of H⁺-ATPase (Kasamo 2003). The activation of H⁺-ATPase is dependent on the degree of saturation or unsaturation of the fatty acyl chain and its length. The activity decreased with an increase in the length of the chain and in the degree of unsaturation (Kasamo 2003; Martz et al. 2006).

4 Function of H₂O₂ in Abiotic Stress in Plants

It is known that the negative effect of the various environmental stresses, e.g., salinity, heavy metals, and extreme temperature, is partially due to the generation of reactive oxygen species and the inhibition of the systems which defend against them (Alscher et al. 1997; Quan et al. 2008). The reactive oxygen species H₂O₂ is a harmful cellular metabolite. To some extent, excess H₂O₂ accumulation can lead to oxidative stress in plants, which then triggers cell death (Quan et al. 2008). Although ROS were originally considered to be detrimental to cells, they are now widely recognized as key factors modulating cellular activities (Suzuki and Mittler 2006; Quan et al. 2008). There are several possible sources of H₂O₂ in plants which can be activated during abiotic stress, e.g., electron transport chains in chloroplast and mitochondria, photorespiration in peroxisomes, or enzymatic sources including plasma membrane NADPH oxidase and cell wall-bound peroxidases/amine oxidase. However, under stress conditions, much attention is focused on plasma membrane and apoplast compartments, in which increases in H₂O₂ could be associated with the activities of NADPH oxidases and peroxidases (Bolwell et al. 2002). Plants tolerate much higher concentration of endogenous H₂O₂ than other organisms (Queval et al. 2008). This increased tolerance could be attributed to the fact that accumulation preferentially occurs in the apoplast and that intracellular concentrations are much lower. ROS originating from the plasma membrane NADPH oxidase have been focus in ROS signaling (Jiang and Zhang 2002). It has been shown that NADPH oxidase is involved in plant growth and development and in plant responses to pathogens, elicitors, and wounding (Sagi and Fluhr 2006), chilling (Piotrovskii et al. 2011), drought (Jiang and Zhang 2002), and salt stress (Yang et al. 2007). This enzyme oxidizes cytoplasmic NADPH and transfers an electron to molecular O₂ to form O₂⁻; the latter is then converted to H₂O₂ by superoxide dismutase (SOD) (van Gestelen et al. 1997). NADPH is mainly generated by the pentose phosphate pathway (Yu et al. 2014). Glucose-6-phosphate dehydrogenase (G6PDH) is the rate-limiting enzyme of this pathway. Recent studies demonstrated that G6PDH plays important role under salt stress and heavy metals (Wang et al. 2008; Jakubowska et al. 2015). However, studies that have been carried to elucidate the physiological role of this protein in plant tolerance to abiotic stresses are still limited.

In addition, apoplastic ROS generation can be mediated by two amine oxidases, the diamine oxidase (DAO) and polyamine oxidase (PAO). DAO and PAO are localized in peroxisomes and the apoplast (Moschou et al. 2008). DAO oxidizes putrescine, whereas PAO oxidizes spermidine and spermine and yields Δ¹-pyrroline and 1,5-diazabicyclononane, respectively, along with 1,3-diaminopropane and H₂O₂. Increased DAO activity resulted in increased tolerance to salinity (Waie and Rajam 2003). It was also shown that upon salt stress spermidine is secreted into apoplast and catabolized by PAO to produce H₂O₂ (Moschou et al. 2008). Polyamines (PAs) are low molecular weight polycationic compounds present ubiquitously in all living cells. Data suggest that PAs are able to affect the activity of H₂O₂-scavenging enzymes, moderating this signal at molecular level (Kubiś 2003). Some evidence

has shown that PA oxidation is directly involved in plant adaptation to abiotic stresses (Moschou et al. 2008). H_2O_2 derived from PA catabolism could also exert signaling effects. Toumi et al. (2010) have shown that the intrinsic ABA signal upregulates PA metabolism, which in turn increases endogenous H_2O_2 load through the apoplastic PAs exodus/catabolism pathway. Also Paschalidis et al. (2010) have indicated that ABA is partly responsible for the induction of the polyamine exodus pathway in plants. They demonstrated that ABA is an upstream signal for the induction of the polyamine catabolic pathway.

Increasing evidence indicates that hydrogen peroxide functions as a signaling molecule responsible for induction of many genes encoding enzymes involved in cellular protection under stress conditions (Vandenabeele et al. 2003; Volkov et al. 2006; Quan et al. 2008). It was observed that treatment of plants with H_2O_2 contributes to increased expression of PM H^+ -ATPase genes in cucumber roots (Janicka-Russak et al. 2012a, 2018). Furthermore, it was shown that abscisic acid causes enhanced generation of ROS (Pei et al. 2000; Laloi et al. 2004). ABA is known as a stress hormone, which mediates responses to a variety of stress factors. Some studies demonstrated that increased H_2O_2 production in apoplast under abiotic stresses occurs as a result of elevated levels of ABA (Hu et al. 2005, 2006). Furthermore, ABA stimulates H_2O_2 production by NADPH oxidases (Hu et al. 2005). So, it seems that under abiotic stress, the elevated level of H_2O_2 could partly result from increased amounts of ABA.

5 Function of NO in Abiotic Stress in Plants

Nitric oxide is a small, highly diffusible, gaseous molecule with ubiquitous bioactivity. It is synthesized in plant cells either during physiological or stress conditions. Endogenously generated NO and NO-derived compounds referred as reactive nitrogen species (RNS) are involved in many primary physiological processes including seed germination, root growth and development, control of stomatal movements, flowering, regulation of pollen tube growth, ripening of fruits, and senescence (Domingos et al. 2015; Sami et al. 2018). NO and RNS play also an extensive role in triggering the plant defense mechanisms in plant response to abiotic stresses such as salinity, drought, high and low temperature, and heavy metals (Corpas et al. 2011; Sahay and Gupta 2017; Sami et al. 2018).

Biosynthesis of NO occurs in certain cellular compartment including cytoplasm, chloroplasts, mitochondria, peroxisomes, and apoplast space (Tischner et al. 2004; Jasad et al. 2006; del Río et al. 2003; Stöhr and Strelau 2006). Several oxidative and reductive and enzymatic and nonenzymatic pathways for NO biosynthesis have been discovered in plants. Reductive route of NO production involves reduction of nitrite ions and is dominantly associated with catalytic activity of cytoplasmic nitrate reductase (cNR). To date, the numerous studies confirmed the role of cNR in NO production in plants during physiological processes and under abiotic stresses (Kolbert et al. 2010; Hancock 2012; Sun et al. 2014). Further, during anoxia, mitochondrial inner membrane

can also participate in reductive pathways of NO biosynthesis (Tischner et al. 2004; Gupta and Igamberdiev 2011). Xanthine oxidoreductase (XOR), a molybdenum-containing enzyme is responsible for NO production occurring in peroxisomes (Weidert et al. 2014). In root apoplast space, NO_2^- conversion to NO is performed by plasma membrane-associated nitrite-nitric oxide reductase (Ni-NOR), which is closely related to plasma membrane nitrate reductase (PM-NR) (Stöhr and Ullrich 2002; Eick and Stöhr 2012). The oxidative pathways include the enzymatic production of NO from L-arginine. In animals, nitric oxide synthases (NOS) oxidizing L-arginine to NO via L-citrulline are well recognized. Despite many years' explorations, there is no direct molecular evidence for existence of homologous NOS in higher plants, although NO production sensitive to mammalian NOS inhibitors has been detected (Corpas et al. 2009; Santolini et al. 2017).

The action mechanism of nitric oxide in plant response to abiotic stresses can be manifested through affecting the function and activity of numerous proteins due to their gene expression and/or protein posttranslational modifications (PTM). S-nitrosylation of cysteine residue, nitration of tyrosine, and metal nitrosylation are the main ones (Astier and Lindermayr 2012). S-nitrosylation consists a reversible covalent binding of NO moiety to the thiol group of cysteine (Cys) residue of target proteins. This reaction is resulting in the formation of S-nitrosothiol (Astier et al. 2011). S-nitrosylation can alter proteins activity, stability, conformation, subcellular localization, and protein-protein interactions (Farnese et al. 2016). The evidences indicate that this PTM is a crucial component of primary metabolism and photosynthesis and contributes to plant signaling and responses to abiotic stresses (Yu et al. 2014; Sami et al. 2018). To date, several proteins, such as H^+ -ATPase, NADPH oxidase, ascorbate peroxidase glutathione reductase, or nitrosogluthathione reductase, concerned with stress reaction or ROS production and/or detoxification undergo modifications through S-nitrosylation (Tanou et al. 2009; Yun et al. 2011; Romero-Puertas et al. 2013).

Tyrosine (Tyr) nitration is another NO-mediated protein modification which involves addition of nitro group ($-\text{NO}_2$) transferring from peroxynitrite (ONOO^-) to the aromatic ring of Tyr residue (Corpas et al. 2009). Tyr nitration leads to structural and functional changes in target proteins and leads to altered cell and tissue homeostasis. Several stress-related proteins, such as some antioxidant enzymes, were identified to undergo Tyr nitration (Mata-Pérez et al. 2016). Metal nitrosylation consists of reversible NO binding to transitions metal, e.g., iron, zinc, and copper, in metalloproteins to form metal-nitrosyl complexes (Astier and Lindermayr 2012; Farnese et al. 2016). This PTM induces changes of protein conformation that affects the activity or reactivity of target proteins (Astier and Lindermayr 2012). Besides posttranslational proteins modifications, NO can affect gene expression of several protein related to and involved in plant response to abiotic stresses. Other studies showed that exogenous NO induces the expression of plasma membrane H^+ -ATPase in plants under salt stress (Siddiqui et al. 2010).

Many studies have documented the essential role of NO for the plant's tolerance to elevated salt concentrations. Generally, it was shown that during salt treatment, NO improved biomass of shoots and roots (Egbichi et al. 2014), enhanced relative

water content and chlorophyll level, decrease leakage of electrolyte, and protect from and reduces membrane injury and lipid peroxidation (Liu et al. 2016). Plant's exposure to raised NaCl concentration leads to accumulation of NO in cucumber tissues (Janicka et al. 2018). Salt-induced NO synthesis is associated with NR-dependent and/or NOS-like-dependent NO biosynthesis pathway (Zhao et al. 2007; Reda et al. 2018). Moreover, NO enhances K^+/Na^+ ratio by increasing activity of plasma membrane and tonoplast proton pumps and Na^+/H^+ antiport (Zhang et al. 2006; Kabała and Janicka-Russak 2012). In plants, both extremely high and low temperatures lead to accumulation of *S*-nitrosothiols and rise of protein nitration (Corpas et al. 2011; Farnese et al. 2016; Sami et al. 2018). To date, extreme temperatures usually promote a rapid increase in the endogenous NO and other RNS levels (Corpas et al. 2011; Li et al. 2013; Yu et al. 2014; Janicka et al. 2018). Moreover, exogenous application of NO to plants subjected to high temperature stress enhances the activity of several antioxidant enzymes (Sami et al. 2018). It was shown that activation of plasma membrane H^+ -ATPase occurring during low temperature acclimation is NO mediated (Janicka et al. 2018). Additionally, action of NO on plasma membrane proton pump was tightly associated with H_2O_2 (Janicka et al. 2018). A growing body of evidence shows changes in NO generation in various plants subjected to heavy metals. However, intensity of NO production seems to be strictly dependent on form and concentration of metal used (Chmielowska-Bąk et al. 2014). Exogenously applied NO alleviates Cd toxicity by increasing pectin and cellulose content in cell walls leading to increasing Cd deposition in root cell walls (Xiong et al. 2009). NO can also promote incorporation of metals into vacuole by increasing phytochelatin synthesis (Arasimowicz-Jelonek et al. 2011) or metal removal from the cell by enhancing the activity of plasma membrane proton pump (Janicka-Russak et al. 2012a, b).

6 Conclusion

Recent studies show that NO and H_2O_2 interact to alleviate the negative influence of abiotic stress on plant growth, metabolism, and development. Both NO and H_2O_2 are generated in plant cells to activate different resistance mechanisms. As the plasma membrane is the main place of the signal perception related to unfavorable conditions, the PM-bound H^+ -ATPase, responsible for proton gradient generation, seems to be a key enzyme involved in plant adaptation to changing environment. The activity, phosphorylation status, and gene expression of plasma membrane proton pump are modulated under abiotic stress conditions. It was demonstrated that enhanced PM H^+ -ATPase functioning is one of the main adaptive mechanisms enabling plant survival under salinity, heavy metals, and low and high temperatures. Recently, NO and H_2O_2 were recognized as signaling molecules, which can affect PM proton pump action. Therefore, it is postulated that both NO and H_2O_2 signaling pathways are involved in the regulation of this enzyme and improvement of stress-induced cell disorders (Fig. 1).

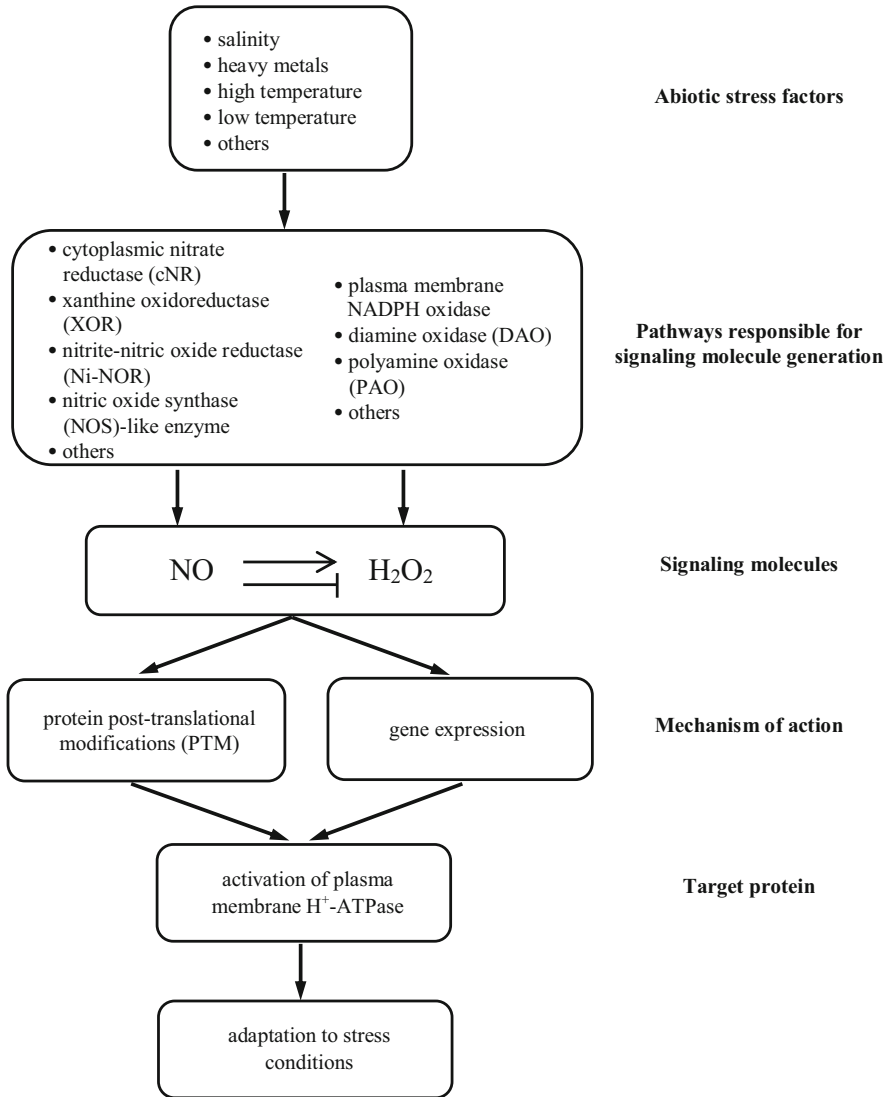


Fig. 1 Model of NO and H₂O₂ signaling pathways, induced under abiotic stress conditions, responsible for activation of plasma membrane H⁺-ATPase involved in plant stress tolerance. Abiotic stresses, including salinity, heavy metals, and low as well as high temperatures, are known to induce NO and H₂O₂ generation. Increase in NO and H₂O₂ levels occurs as a result of enhanced activity of different enzymatic systems existing in plant cells. Additionally, NO can affect H₂O₂ accumulation, elevating or inhibiting it. Both NO and H₂O₂ act as signaling molecules which modulate the function and activity of numerous proteins due to their gene expression and/or posttranslational modifications (PTM), such as reversible phosphorylation, S-nitrosylation, or Tyr-nitration. The PM-bound H⁺-ATPase, responsible for proton gradient generation, is a key enzyme involved in plant adaptation to environmental stresses. NO as well H₂O₂ can activate PM proton pump at both genetic and protein levels and in this way participate in plant tolerance

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