



Nitric oxide, crosstalk with stress regulators and plant abiotic stress tolerance

Xianrong Zhou¹ · Shrushti Joshi² · Tushar Khare^{2,3} · Suraj Patil² · Jin Shang¹ · Vinay Kumar^{2,3}

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Abstract

Key message Nitric oxide is a dynamic gaseous molecule involved in signalling, crosstalk with stress regulators, and plant abiotic-stress responses. It has great exploratory potentials for engineering abiotic stress tolerance in crops.

Abstract Nitric oxide (NO), a redox-active gaseous signalling molecule, though present uniformly through the eukaryotes, maintain its specificity in plants with respect to its formation, signalling, and functions. Its cellular concentrations are decisive for its function, as a signalling molecule at lower concentrations, but triggers nitro-oxidative stress and cellular damage when produced at higher concentrations. Besides, it also acts as a potent stress alleviator. Discovered in animals as neurotransmitter, NO has come a long way to being a stress radical and growth regulator in plants. As a key redox molecule, it exhibits several key cellular and molecular interactions including with reactive chemical species, hydrogen sulphide, and calcium. Apart from being a signalling molecule, it is emerging as a key player involved in regulations of plant growth, development and plant-environment interactions. It is involved in crosstalk with stress regulators and is thus pivotal in these stress regulatory mechanisms. NO is getting an unprecedented attention from research community, being investigated and explored for its multifaceted roles in plant abiotic stress tolerance. Through this review, we intend to present the current knowledge and updates on NO biosynthesis and signalling, crosstalk with stress regulators, and how biotechnological manipulations of NO pathway are leading towards developing transgenic crop plants that can withstand environmental stresses and climate change. The targets of various stress responsive miRNA signalling have also been discussed besides giving an account of current approaches used to characterise and detect the NO.

Keywords Nitric oxide · Signalling molecule · Abiotic stress · Genetic engineering · Transcription factors · miRNA · Crosstalk

Introduction

Being sessile organisms, plants experience several challenging and stressful external conditions throughout their life cycles that include abiotic (drought, salinity, extreme temperatures, and heavy metals, beside other) and biotic stress factors (bacterial, fungal, viral, and herbivory). Plants respond to these conditions via inducing a series of different molecular, cellular and physiological changes (Yadav et al. 2020). A common consequence and response is generation of reactive chemical species that at low cellular concentrations act as signalling molecules regulating secondary messengers, enzyme activities and expression of key genes (Farnese et al. 2016). However, these reactive species may turn toxic to cells and organelles if their concentrations increase beyond threshold levels (Khare et al. 2020). These reactive species in plants include reactive

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✉ Xianrong Zhou
zxrfy@163.com

✉ Vinay Kumar
vinaymalik123@gmail.com;
vinay.kumar@moderncollegegk.org

¹ School of Life Science and Biotechnology, Yangtze Normal University, Chongqing 408100, China

² Department of Biotechnology, Modern College of Arts, Science and Commerce, Savitribai Phule Pune University, Ganeshkhind, Pune 411016, India

³ Department of Environmental Science, Savitribai Phule Pune University, Pune 411007, India

oxygen species (ROS), reactive nitrogen species (RNS), reactive sulphur species (RSS) and reactive carbonyl species (RCS) (Zhou et al. 2021). Amongst different reactive species, ROS and RNS have gained significant attention because of their prevalent presence and dual roles in cellular metabolism, as signalling molecules or as oxidative burst causing-agents; based on their concentration in cell (Patel et al. 2019). Notable work has been done to understand and explore the roles of ROS in plant cells such as involvement in stomatal guard cell opening (Yamauchi et al. 2019), plasma membrane fluidity (Møller et al. 2007), male-sterility and tapetal programmed cell death (PCD) signalling (Neill et al. 2002; Miller et al. 2010; Suzuki et al. 2011). The RNS biology, however, is emerging as a topic of great interest with linear increase in the reports coming on various aspects of RNS in plants. Members in RNS family include nitric oxide (NO), S-nitrosothiols (RSNOs), S-nitrosoglutathione (GSNO), and peroxynitrite (ONOO⁻). From such diverse members with contrasting characters, the only unifying character among RNS members is their derivation from nitric oxide (NO). Hence, it was necessary to investigate and decipher the NO biochemistry and physiological roles played by it to get better insights on RNS.

NO is a small diatomic gaseous molecule, with a short half-life. In early years, NO was thought to be involved mainly in air pollution and acid rains. It was not earlier than 1987, when role of NO as endothelium-derived relaxation factor (EDRF) in mammalian cells and synthesis of NO from amino acid L-arginine by enzyme NO synthase were identified (Ignarro et al. 1987; Palmer et al. 1987, 1988). Since then, extensive work has been carried out to identify the roles and functions of NO in both animals and plants. During stress, NO affect enzymes and proteins predominantly by covalently attaching to cysteine thiols by S-nitrosylation, along with other modifications such as nitrosylation of metalloproteins and tyrosine residues (Lindermayr 2018). Redox signal of NO by S-nitrosylation has recently emerged as a pioneer signal transmitter of NO bioactivity (Fernando et al. 2019). S-nitrosylation is a selective covalent post-translational modification, through which a nitrosyl group is added to reactive thiol group of a cysteine to ultimately form S-nitrosothiol, a key mechanism in transferring NO-mediated signals (Fernando et al. 2019). NO is also known to be involved in signalling for plant defence (Bellin et al. 2013), antioxidant and inhibitor of PCD (He et al. 2019a), plant-rhizobium infection (Fukudome et al. 2016), plant-fungal and oomycetes interactions (Jedelská et al. 2020), besides, symbiotic associations (Martínez-Medina et al. 2019). Along with this, NO has also been reported to be involved in plant responses to abiotic stresses such as salinity (Hasanuzzaman et al. 2018), drought (Montilla-Bascón et al. 2017), and metal stress (Sharma et al. 2019).

This review discusses the current knowledge and updates on biosynthesis, transport, and scavenging of NO in plants. A critical look on how NO interact with different stress regulators like secondary messengers, reactive chemical species, and phytohormones has been provided. Besides, this review discusses how these interactions can be used for crop improvement especially in creating abiotic stress (or climate) resilient crops using genetic engineering approaches. Recent trends in characterisation of NO along with idea of micro-RNA (miRNAs) signalling and regulations due to NO during stress conditions and adaptations are also discussed.

Nitric oxide: a dynamic entity involved in plant growth and stress responses

For the initial years after its discovery, NO was known as the 'nitrous air' or more commonly the 'laughing gas' and thus was extensively studied in mammals for its potential neurological impacts. The research shifted to plants after the report of NO generation/biosynthesis in plant cells (Kleppe 1975). The NO research in animals got the Nobel Prize when Robert F. Furchgott, Louis J. Ignarro, and Ferid Murad shared the Nobel Prize in 1998 for their discoveries concerning NO as a signalling molecule in the cardiovascular system (Nicholls 2019). Gradually, NO's role in plants shifted from being a reactive species and a signalling molecule, to its involvement in plant defence system, changing the paradigm from cytotoxic to beneficial compound (Sahay and Gupta 2017). The role of NO varies across the strata of the organisms. For instance, it acts as a neurotransmitter in animals and regulates muscular, immunological, and respiratory responses (Esplugues 2002). In plants, it has displayed the involvement in stress response/signalling, plant growth and development and the PCD (Gupta et al. 2011). Evidence is also there that NO may protect some bacteria against oxidative stress (Gusarov and Nudler 2005). Figure 1 depicts the involvement and roles of NO in plant development and plant-environment interactions.

Being an important player in modulating plant growth under stress, NO has been reported to coordinate the activities required for plant organogenesis by participating in the pathways for production of plant hormones like jasmonic acid (JA), and brassinosteroids (BRs) (Raya-González et al. 2019). Auxins (AUX), a class of plant hormones are known to stimulate NO production in plant roots which results in lateral root development (Moni et al. 2018). Another possible mechanism is the methane-induced accumulation of NO, which leads to the increased formation and development of adventitious roots (Qi et al. 2017). Beside these roles, NO plays a vital role as a signalling molecule during the root development, nitrogen fixation, root-fungal interactions and promotion of high microbial diversity in the rhizospheric

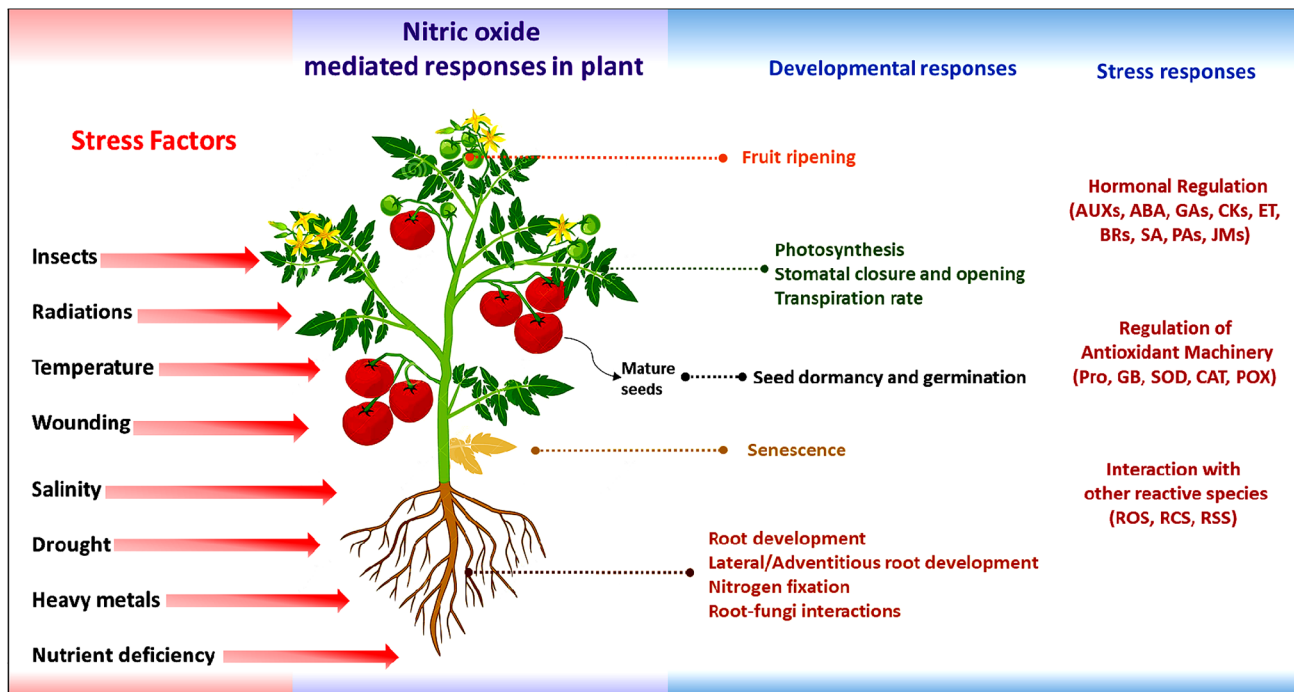


Fig. 1 Nitric oxide (NO) mediated responses during plant development and plant-environment interactions. (AUX auxins, ABA abscisic acid, GAs gibberellins, CKs cytokinins, ET ethylene, BRs brassinosteroids, SA salicylic acid, PAs polyamines, JMs jasmonates, Pro pro-

line, GB glycine betaine, SOD superoxide dismutase, CAT catalase, POX peroxidase, ROS reactive oxygen species, RCS reactive carbonyl species, RSS reactive sulfur species)

environment (Ma et al. 2020). In *Tagetes erecta*, NO donor sodium nitroprusside (SNP) showed indirect organogenesis of shoot and root from callus, showing its role in plant growth (Jafari and Daneshvar 2020).

The role of NO in seed germination has also been described in many reports. The pre-treatment of seeds with SNP have been extensively explored. For instance, exogenous application of SNP to seeds of *Brassica chinensis* exerted positive effects on seed germination rate and seedling growth under saline conditions via enhanced scavenging activity of enzymatic antioxidants including superoxide dismutase (SOD), catalase (CAT) and peroxidase (POX) (Ren et al. 2020). In *Brassica juncea*, SNP pretreated seeds showed improved seed germination rates under copper (Cu) stress due to induced antioxidant defense and increased amylase activity (Rather et al. 2020). Under chromium (Cr) stress, exogenous application of SNP relieves stress-induced impairment via promoting the activities of scavenging enzymes, amylase, and protease resulting in increased seed germination (Khan et al. 2020a). Table 1 enlists some important case studies involving exogenous applications of NO and its donor (SNP) for alleviation of negative stress impacts.

Peroxisomes are thin protrusions from the spherical peroxisomes, produced when the ROS levels are low. The peroxule production in a plant cell is highly dependent on the presence

of NO in the cytoplasm (Terrón-Camero et al. 2020). In *Arabidopsis* under cadmium (Cd) stress, NO production and interaction were observed via electron microscopy during peroxule formation (Terrón-Camero et al. 2020). Exogenous application of NO to fresh fruits and vegetables showed delayed senescence and worked as a freshener or stabiliser during long-term storage of fruits/vegetables (Steelheart et al. 2019). In sweet pepper, the fruit ripening process was found to be NO-dependent (González-Gordo et al. 2019). Authors observed NO-dependent changes in sweet pepper fruit transcriptome, fruit ripening in the absence of NO showed changes in the abundance of 8805 transcripts, and the functional clusters associated with ROS/RNS were significantly modified. The NO treatment induced differential expression of 498 genes within these functional categories. These findings reveal the key roles of NO in the ripening of sweet pepper fruits (González-Gordo et al. 2019).

NO is emerging as a vital player underlying abiotic and/or biotic stress interactions in plants. NO production during such interactions tends to alleviate in such stressful environment. NO has demonstrated to regulate the chlorophyll production and to stimulate the hypersensitive chain reactions during stress interactions (Lazalt et al. 1997; Romero-Puertas et al. 2004). Stress-mediated increase in NO production stimulated the activity of various ATPases, and ion channels for better homeostasis. In maize for instance,

Table 1 A summarised list of plant abiotic stress alleviation via exogenous application of nitric oxide and/or its donor sodium nitroprusside

Plant	Approach	Phenotypic responses	References
<i>Citrus sinensis</i> L.	Exogenous NO donor SNP application under cold stress	Reduced lipid peroxidation levels and hydrogen peroxide contents. NO-induced DPPH radical scavenging activity	Ghorbani et al. (2018)
<i>Brassica napus</i> L.	Exogenous NO and aminolaevulinic acid application under drought stress	Decrease in oxidative damage and improved growth, chlorophyll content and antioxidant enzyme activity	Akram et al. (2018)
<i>Glycine max</i> L.	Exogenous SNP and S-nitroso L-cysteine application under flood stress	Induced glutathione and reduced superoxide anion contents, elevated ABA contents and mitigated the negative effects of short-term flooding stress	Khan et al. (2019)
<i>Hordeum vulgare</i> L.	SNP treatment under Cd stress	Increased expression levels of <i>AOX1a</i> gene, and enhanced Cd stress tolerance	He et al. (2019b)
<i>Brassica juncea</i> L.	NO and SNP pre-treatment under oxidative stress	Advancement in seed germination, alleviation in Cu-induced photosynthetic inhibition	Rather et al. (2020)
<i>Brassica juncea</i> L.	SNP treatment under salt stress	Regulated the photosynthetic performance and induced salt stress alleviation	Jahan et al. (2020)
<i>Triticum aestivum</i> L.	Exogenously-applied NO donor SNP and H ₂ O ₂ as seed primers under drought conditions	Reduction in H ₂ O ₂ and MDA contents and improved drought tolerance	Habib et al. (2020)
<i>Triticum aestivum</i> L.	Exogenous NO donor SNP in presence of TiO ₂ nanoparticles under salt stress	Enhanced seedling dry mass, relative water content, antioxidant enzymes, and proline content, coupled with reduced hydrogen peroxide (H ₂ O ₂) and malondialdehyde (MDA) contents	Faraji and Sepehri (2020)
<i>Cucumis sativus</i> L.	Exogenous NO treatment under low-temperature stress	Improved tolerance to low-temperature stress via enhanced efficiency of the Calvin cycle and the ascorbate–glutathione cycle	Wu et al. (2020)
<i>Solanum lycopersicum</i> L.	SNP treatment under chromium stress	Relieved impairment caused due to chromium by promoting activities of scavenging enzymes, amylase, and protease resulting in increased seed germination	Khan et al. (2020a)
<i>Cucurbita pepo</i> L.	SNP treatment under chilling stress	Enhanced chilling tolerance through S-nitrosylation of proteins and modulation of the antioxidant response	Jiménez-Muñoz et al. (2021)

NO was shown to improve salinity tolerance via regulating the plasma membrane proton pump activities (Zhao et al. 2004). The authors provided the evidence for establishing the involvement of NO in salt tolerance of *Phragmites communis* (Zhao et al. 2004). NO, produced under salt stress, served as a second messenger for the induction of plasma membrane H^+ -ATPase expression, which in turn maintained high K-to-Na ratio for conferring better salt resistance (Zhao et al. 2004). SNP-treatment has been reported to exert several positive impacts like increased seed germination in *Lupinus luteus* (Kopyra and Gwózdź 2003), restored developmental mechanisms in seedlings of sunflower (David et al. 2010), plant growth, increased ionic levels and scavenging enzymes activity in citrus during saline stress (Khoshbakht et al. 2018). Stomatal opening and closing play major roles in conferring plant abiotic stress tolerance. NO being a gas-transmitter, regulates the signalling network involved in the movement of guard cells, causing stomatal opening and closing (Nabi et al. 2019). During heat stress, the heat shock response (HSR) of the plants mediate through biosynthesis and transport of NO and salicylic acid (SA) via calcium signalling (Rai et al. 2020). This acts as a regulator of Heme Nitric oxide/Oxygen (H-NOX) motif of a heme binding protein involved in the regulation of endogenous NO synthases. High concentrations of NO and SA cause conformational changes in non-expresser of pathogenesis related (NPR) genes involved in stress-responsive genes against heat (Rai et al. 2020). These NPR genes act as master regulators of SA signalling under pathogen attack and are important in terms

of providing plants innate immunity (Innes 2018). Thus after several years of extensive research on deciphering the roles and mechanisms of NO in plants, various avenues have been opened up on formulations of new research hypotheses. Owing to its dynamic nature and versatility, NO provides a great scope for its application and exploration in crops for increasing the plant efficiency and tolerance/resilience to stress conditions.

Biosynthesis, transport and scavenging of nitric oxide in plants

NO is endogenously present in the plant cells as a by-product of various catabolic reactions of nitrogen-containing compounds. Besides, it is absorbed by the plants from the surrounding environment and soil. The knowledge and understandings about the NO synthesis in plants was influenced by the findings on NO production in animal systems putting an emphasis on nitrate reductase (NR), the enzyme that usually converted nitrates to nitrites but was later observed to convert the nitrite to nitric oxide (Crawford 2006).

NO has two biosynthetic pathways as depicted in Fig. 2, an oxidative one which includes the production through oxidation of L-arginine or other polyamines (PAs), and second one a reductive pathway which heavily depends on NR and other reductive enzymes present in the mitochondrial and plasma membrane (Gupta and Igamberdiev 2011). The reductive pathway uses enzymes produced in

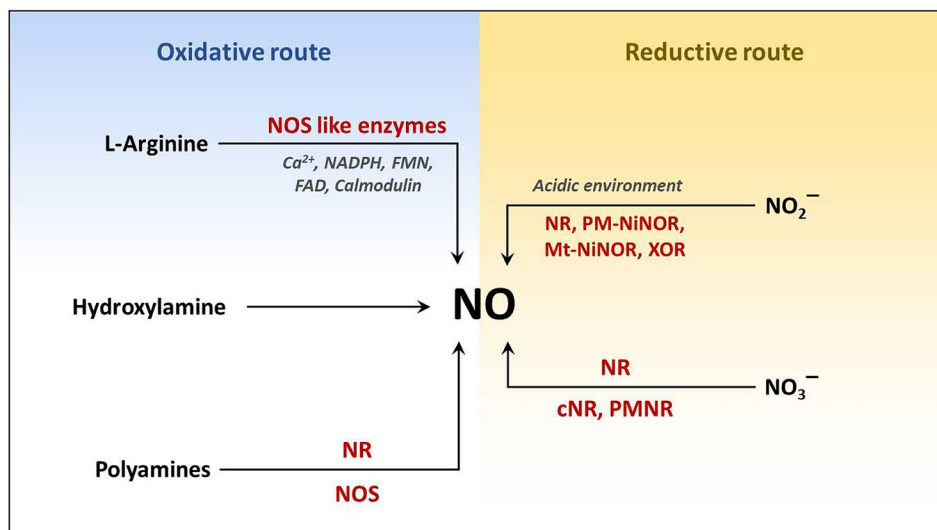


Fig. 2 Oxidative and reductive routes of nitric oxide (NO) biosynthesis. Reductive route (observed in plasma membrane, chloroplasts, apoplast, peroxisomes, cytoplasm, and mitochondria) occur via reduction of nitrate (NO_3^-) or nitrite (NO_2^-) by nitrate reductase (NR). In oxidative route (observed in chloroplasts and leaf peroxisomes) occurs via oxidation of L-Arginine or hydroxylamine. (cNR cyto-

solic nitrate reductase, PMNR plasma membrane-bound nitrate reductase, PM-NiNOR plasma membrane-bound nitrite NO reductase, Mt-NiNOR mitochondrial electron transfer chain-dependent enzymatic nitrite NO reductase, XOR xanthine oxidoreductase, NADPH nicotinamide adenine dinucleotide phosphate, Ca^{2+} calcium ions, FMN flavin mononucleotide, FAD Flavin adenine dinucleotide)

the mitochondrial electron transport chain (ETC), mainly amidoxime reducing compound, molybdenum cofactor dependent reducers (León and Costa-Broseta 2020), and other reductases like xanthine oxidoreductase (XOR), nitrite dependent nitric oxide reductase (Ni:NOR), along with NR (Sahay and Gupta 2017). This pathway is usually dominant under normoxic conditions whereas oxidative pathway requires a hypoxic environment (León and Costa-Broseta 2020).

The production of NO involves the participation of various organelles like mitochondria, chloroplast, peroxisomes, endoplasmic reticulum, apoplast, plasma membrane, and the cytoplasm of the cell (Sahay and Gupta 2017). The mechanism of production varies between organelles depending on their preference to either of the two previously mentioned pathways. The reductive pathway occurs in the plasma membrane where lower pH converts NO_2^- to NO and O_2 and in apoplast releasing the root-specific NO using nitrite substrates present in it. Organelles like chloroplast, peroxisomes, and mitochondria switch the production pathways depending on the availability of substrates and the environmental conditions. Cytoplasm converts nitrate into nitrite in an NADPH-dependent manner for further reduction along with oxidation of PAs. It also produces intermediate NO during GSH to GSNO conversion via a glutathione-based pathway (Nabi et al. 2019).

Exogenous NO is transported inside the cell via nitrogen transporter in the form of nitrate, which then undergoes reductive pathway forming NO by nitrate NR in an NADPH-dependent manner (Zhou et al. 2021). The nitrate homeostasis is maintained by high- and low-affinity nitrate transporters (Chamizo-Ampudia et al. 2017). Along with these, transporters belonging to the chloride channel (CLC) family maintain nitrate concentrations in the cytoplasm via effluxes of various organelles.

NO, like all metabolites in a cell needs to be balanced at a particular concentration, which is done by the cell's metabolic and scavenging activities regulated by specific enzymes and substrates. In animals, proteins like flavohaemoglobin, haemoglobin and reductases degrade the excess NO while in yeast flavoglobin is the main NO degrader (Liu et al. 2000; Gardner 2005; Gupta et al. 2020b). Similarly, plants have their NO scavengers namely phytohemoglobins, that are highly efficient under hypoxic conditions converting NO to nitrate by oxidising itself into metphytohemoglobin (Gupta and Igamberdiev 2011). This is converted back to phytohemoglobin by a reductase mediated reaction co-factored by ascorbate (Hebelstrup et al. 2008; Gupta and Igamberdiev 2011; Gupta et al. 2020b). Reduced glutathione (GSH) reacts with NO to form nitrosoglutathione (GSNO), which can be harnessed to give back NO for nitrosylation of proteins and thus contributing to its cellular concentration maintenance (León and Costa-Broseta 2020; Kalinina and Novichkova 2021).

On the other hand, a further oxidation of GSNO results in production of ammonia and glutathione disulphide by specific reductase (León and Costa-Broseta 2020). When the nitrosylated proteins are non-nitrosylated, they release NO back in the cell affecting the maintained NO-balance (León and Costa-Broseta 2020). This happens when the S-nitrosylation-related signalling is affected during pathogen attack, which triggers plant's inherent immunological response. A thioredoxin-mediated redox signalling was reported to be involved in regulation of S-nitrosylation of proteins, contributing to the homeostasis of NO and involved in plant immunity (Mata-Pérez and Spoel 2019; León and Costa-Broseta 2020). Other regulators of NO degradation are NAD(P)H dehydrogenases-type dehydrogenases, that act by converting the NO into peroxy nitrite (ONOO^-) via forming the superoxide anions, under calcium stimulation and hindered by anoxic conditions by SOD (Gupta et al. 2020a).

Another method of NO scavenging is during reactive species reaction when NO interacts with oxygen generating nitrate and nitrite or when it forms nitro fatty acids via its reactions with lipid peroxyl radical ($\text{LOO}\bullet$) (Hancock 2012; Rubbo 2013; León and Costa-Broseta 2020). Haemoglobins have been reported to contribute in NO scavenging through its reduction to FE(II)Hb which then deoxygenates the NO (Chamizo-Ampudia et al. 2017). NO is metabolised in presence of oxygen by being converted to dinitrogen trioxide (N_2O_3) and NO_2 . Subsequently, in an aqueous environment, they are converted to nitrite and nitrate as part of a nitro-oxidative stress response (Corpas and Palma 2018). Overall, we can conclude that the NR is a key enzyme for nitrogen acquisition by plants. Nitrate, its main substrate, is required for signalling and is widely distributed in diverse tissues in plants. In addition, NR has been proposed as an important enzymatic source of NO (Chamizo-Ampudia et al. 2017). The homeostasis of NO depends on at least two key molybdoenzymes, NR and NO forming NR (NOFNiR) besides dioxygenase activity of haemoglobins.

Current approaches for detection and characterisation of nitric oxide

Characterisation and quantification remain two most important parameters in studying the reactive species in stress biology. Among different reactive species, measurement of NO remains challenging and tricky because of its ubiquitous, gaseous and unstable nature. Another parameter while characterising NO is inter-conversion of NO^- , NO, and NO^+ under physiological conditions; hence care should be taken while designing the experiment and choosing the NO donor to avoid misinterpretation of the results (Yamasaki et al. 2016).

NO detection method based on colorimetric bioassay, mass spectroscopy, chromatography, fluorescent probes; spin traps, electrode-based techniques have been developed (reviewed by Csonka et al. 2015). Each technique has its own advantages and disadvantages. Most frequently used bioassay technique for detection of NO via Griess method. This is a method based on nitrite-derived NO quantification based on the Griess reaction. The Griess reaction is specific for nitrite quantification. Principle behind Griess method is, in presence of O₂, NO gets converted to NO²⁻ which reacts with Griess reagent and is detected spectrometrically at 540 nm using a plate reader (Antoniou et al. 2018). Another version for detection of gaseous NO is using CrO₃ oxidizer column, CrO₃ converts NO to NO₂ which subsequently is trapped using Griess reagent solution to form NO²⁻ and detected using spectrophotometrically (Vitecek et al. 2008). Along NO, detection of S-nitrosothiols has emerged as important player. S-nitrosothiols (RSNO) detection by Saville reaction using mercury chloride and later Griess reagent has been used (Bryan and Grisham 2007).

Real-time monitoring and measuring becomes important in studies when NO is involved in signalling cascade or when formation and conversion takes place in very short time and at low quantity. In such studies bioassay or biophysical techniques are not suitable for real-time monitoring and the real time monitoring can be done using fluorescent probes, chemiluminescent detector or using electrodes. NO-specific probes that are widely used are diaminofluoresceins (DAFs), 2,3-diaminonaphthalene (DAN), 1,2-diaminoanthraquinone (DAQ or DAA), diaminorhodamine (DAR-4M AM), fluorescent nitric oxide cheletropic traps (FNOCT), acridine-TEMPO-DTCS-Fe(II), 8-(3,4-diaminophenyl)-2,6-bis(2-carboxyethyl)-4,4-difluoro-1,3,5,7-tetramethyl-4-bora-3a,4a-diaza-sindacene (DAMBO-PH) (Zhou et al. 2021). Ozone-based chemiluminescent detector (CLD) is another sensitive technique, in which NO reacts with ozone, excited NO₂ emits lights which is detected by photomultiplier (Bryan and Grisham 2007).

Biophysical techniques for measurement of NO are more sensitive. For biological detection of gaseous NO, often detection/quantification is required at very low (ppb) level, and hence high-throughput, ultra-sensitive analytic techniques are needed. NO detection based on Infra-red (IR) spectroscopy at absorbance at 5.3 μm has been reported, accordingly two systems laser photoacoustic detection (LAPD) and quantum cascade lasers (QCL) based on IR spectroscopy have been developed with high sensitivity and specificity (Mur et al. 2011). Mass spectroscopy-based detection is another technique used for detection of gaseous NO present in plant tissues. Two types of MS approaches have been developed for detection of NO, membrane inlet mass spectrometry (MIMS) and restriction capillary inlet mass spectrometry (RIMS) (Conrath et al. 2004). In MIMS/

RIMS system, before MS detection, a prior step for molecule separation is done via membrane (MIMS) or through restriction capillary (RIMS) which allows only small molecules to pass through such as NO. As an advantage over other NO assays, MIMS/RIMS have advantages over other NO assays, as it discriminates nitrogen isotopes and simultaneously measures NO and O₂ (and other gases) from the same sample. This technology thus holds significance and may be used to identify cellular NO sources, besides, it can also be used for elucidating the relationships between primary gas metabolism and NO formation (Conrath et al. 2004).

The most reliable, specific and sensitive technique, however, for detection of NO can be considered as electron spin resonance (ESP) also called as electron paramagnetic resonance (EPR). EPR detects only free radicals, making it specific for NO detection over other radicals (Kleschyov et al. 2007; Mur et al. 2011). It is the most specific method, due to NO-specific spin traps used in it. Major advantage associated with the use of EPR for NO detection is that it only detects the paramagnetic molecules, and the EPR spectrum presents a unique fingerprint of the chemical/electronic structure around the unpaired electron.

Crosstalk of nitric oxide with stress regulators

Crosstalk with signalling molecules

The interaction of NO with signalling molecules such as receptors, secondary messengers, transcription factors (TFs) and enzymes during physiological and environmental responses of the plant cells has led to the study of crosstalk of NO with ions like Ca²⁺, K⁺, and Na⁺. During salinity-induced stress responses, the activities of signalling molecules including ROS, NO and Ca²⁺ are altered in the guard cells leading to the regulatory changes in opening and closing of stomata (Hasanuzzaman et al. 2018). Salinity-driven responses in the mangrove plants revealed the interaction of H₂O₂, ATP, and Ca²⁺ increase the Na⁺ extrusion via NO-induced Na⁺/H⁺ antiporters (Lang et al. 2014). Garcia-Mata et al. (2003) studied the regulation of Ca²⁺ ion channels in *Vicia faba* guard cells mediated through NO by releasing Ca²⁺ from inner cell stores to the cytosol, leading to the controlled guard cell movements and indicating the NO-induced Ca²⁺ activation of cellular signalling. These results confirmed the NO action decisively in one branch of Ca²⁺ signalling pathways engaged by ABA and defined the boundaries of parallel events in control of guard cell movements (Garcia-Mata et al. 2003). Mitogen-activated protein kinase (MAPK) signalling pathway can be activated via phytohormones; however, reports have shown the H₂O₂-NO or NO-mediated cGMP-independent pathway stimulation of

MAPK with the help of secondary messengers during physiological processes and abiotic stress responses (Hasanuz-zaman et al. 2018; Shen et al. 2019). Reports have revealed the crosstalk of NO and MAPK-1/2 via NO generation by NR, the activity of S-nitrosylated glutathione reductase, and activation of MAPK1/2 enzymes during cold stress tolerance in tomato plants (Lv et al. 2017).

An important polyamine spermidine is considered crucial for plant adaptations to the environmental stresses (Alcázar et al. 2006). Besides, spermidine is reportedly involved in the induction of H₂S signalling via increasing the L/D-cysteine desulphydrate (L/D-CD) activities (Li et al. 2019). The TFs such as *bZIP37*, *bZIP107*, *DREB2*, *DREB4*, and *WRKY108715* are regulated by the spermidine during H₂S downstream signalling and crosstalk between NO, H₂S, and Ca²⁺ signalling resulted in water stress tolerance in white clover plants (Li et al. 2019). The WRKY TFs are actively involved in the environmental stress responses. In tomato, *SIWKRY81* reported to repress the transcription of *NR* and to lower the NO accumulation for modulating the stomatal closure and subsequently the drought tolerance (Ahammed et al. 2020). Hajihashemi et al. (2020) studied the crosstalk between NO, H₂O₂ and Ca²⁺ at the germination stages in *Chenopodium quinoa* plants subjected to salinity stress. The findings revealed stimulated amylase activities in the seeds pre-treated either with SNP, H₂O₂ or CaCl₂. This resulted in high starch breakdown rates and thus increased availability of water-soluble sugars; which can be attributed to the improved germination under the saline conditions (Hajihashemi et al. 2020).

Crosstalk with reactive species

The inevitable production of the reactive species during the stress interactions in plants is well-established. The evidences have suggested the interplay between NO and the ROS (such as H₂O₂) as well as RSS (such as H₂S). Aerobic metabolic processes in subcellular-compartments of various organelles results in ROS generation in mitochondria, chloroplasts, apoplasts, and plasma membrane, which is formed by partial reduction of oxygen. Amongst these, the hydrogen peroxide (H₂O₂), superoxide anion (O₂^{•-}) and hydroxyl radical (OH) are the major reactive species (Kaur et al. 2019; Kalia et al. 2017). ROS are major signalling components involved in plant growth, development, and abiotic/biotic stress responses, which interplays with the RNS generation and NO signalling (Hancock et al. 2002; Wrzaczek et al. 2013). NO and ROS play vital roles in biological- and cross-talk processes involved in plant responses to environmental stresses (Kohli et al. 2019). In plants, NO acts as iron ligand in heme-containing enzymes which leads to their activation/inhibition. During stress-interactions, the ROS generation induces NO signalling, where NO binds

with the heme group of the enzymatic ROS scavengers (like CAT). The highly-reactive, lipophilic NO binds to the thiolic- and metallic-groups of enzymes, which may modulate the enzyme activities (Arora et al. 2016). Further, at low NO concentration, the scavenging of OH confers the antioxidant properties to NO. Besides, NO is also known as the strong inhibitor of protein oxidation and lipid peroxidation (Fancy et al. 2017).

Several reports have established melatonin (MET) as one of the amphiphile multi-functional signalling molecules involved in physiological responses, abiotic responses, ROS and RNS scavenging, cell antioxidant activities, and oxidative-stress improvement (Debnath et al. 2019). The MET levels can regulate the ROS concentrations in two ways; firstly, by chemical interaction with ROS leading to the ROS inactivation, or secondly, by MET induced activities of enzymatic antioxidants including SOD, CAT and POX (Arnao et al. 2019; Khan et al. 2020a, b). Exogenous MET associate with CAND2/PMTR1 (CANDIDATE G-PROTEIN COUPLED RECEPTOR 2/Phytomelatonin Receptor 1) receptors, resulting in triggering of responses against stressors and regulation of ROS and RNS (Pardo-Hernández et al. 2020). Liu et al. (2020) reported reduced trehalose-mediated lipid-peroxidation levels in tomato leaves treated with H₂O₂ and NO scavengers, dimethyl thiourea (DMTU) and 2-(4-carboxyphenyl)-4, 4, 5, 5-tetramethylimidazole-1-oxyl-3-oxide (cPTIO). The results indicated that the trehalose might act as H₂O₂ and NO inducer that modulates the expression of genes from antioxidant defense system for conferring cold tolerance. On similar lines, the interactions between H₂O₂ and NO were studied during the BR-induced stress responses in *Medicago truncatula* by Arfan et al. (2019). The results shown the participation of H₂O₂ in NO generation, and NO was found key in BR-induced alternative oxidase capacity, which protected the photosystem under cold stress (Arfan et al. 2019).

Chemical priming is emerging as an effective approach for conferring stress tolerance in plants. For instance, *Citrus aurantium* L. roots pre-treated with SNP (NO donor), enhanced the NO- and H₂O₂-induced salinity and drought acclimation in plants (Molassiotis et al. 2016). Priming of wheat roots (using polyethylene glycol 6000) stimulated the production of H₂O₂ and NO which was attributed to the priming-induced drought tolerance (Wang et al. 2020c). The report also suggested the direct relation of H₂O₂ biosynthesis with respiratory burst oxidase homologs (RBOH) and the downstream working of NO during the H₂O₂ mediated accumulation of proline and glycine betaine (Wang et al. 2020c).

The nutrient recycling event of the leaf senescence is also linked with the redox balance in plants (Jing et al. 2008). During the senescence, there is marked increase in the H₂O₂ level in leaf tissues. Apart from the damaging role of the macromolecular oxidisation that promotes the cell death,

H₂O₂ also functions as a signal to induce the expression of senescence related genes (Cui et al. 2013). On the other hand, NO can provoke or impede the senescence process depending on its subcellular localization and concentration. The toxicity of ROS may get alleviated by the NO via enhancing the activity of antioxidant enzymes (Liu and Guo 2013). Similarly, the intracellular NO and nitrosothiol levels act as vital mediators during the H₂O₂-induced leaf cell death (Lin et al. 2012; Wang et al. 2013a). Collectively, both NO and H₂O₂ can induce the cell death in leaves during which there is observable interplay between NO and H₂O₂ (reviewed by Wang et al. 2013b).

Besides H₂O₂, the H₂S (RSS) is also a vital signalling molecule during seed germination, photosynthesis and environmental stress responses (Chen et al. 2015a, b). The NO and H₂S jointly regulate the key events during the plant growth and development (seed germination, root development, stomatal movement and fruit ripening) and stress tolerance (reviewed by Mukherjee and Corpas 2020). Studies have revealed the interactive effects of NO and H₂S during Cd-induced stress interactions in wheat, which resulted in the reduced oxidative stress and Cd uptake, attributable to the activated antioxidant defence and uptake of essential minerals like Zn and Fe (Kaya et al. 2020). Further, the interaction between NO and H₂S resulted into the generation of persulfide (intermediate product) which regulate the ROS and RNS contents in plant cells (Lisjak et al. 2013). The seedlings of sweet potato exposed of sodium hydrosulfide (NaHS, H₂S donor) displayed the increased levels of H₂S and NO (Zhang et al. 2009). The plants also displayed H₂S-stimulated adventitious root formation mediated through NO, which authors attributed to H₂S activity upstream of the NO signal-transduction pathways (Zhang et al. 2009). However, further studies are still required to establish a deeper understanding of the crosstalk between NO and H₂S (Chen et al. 2016).

Crosstalk with phytohormones

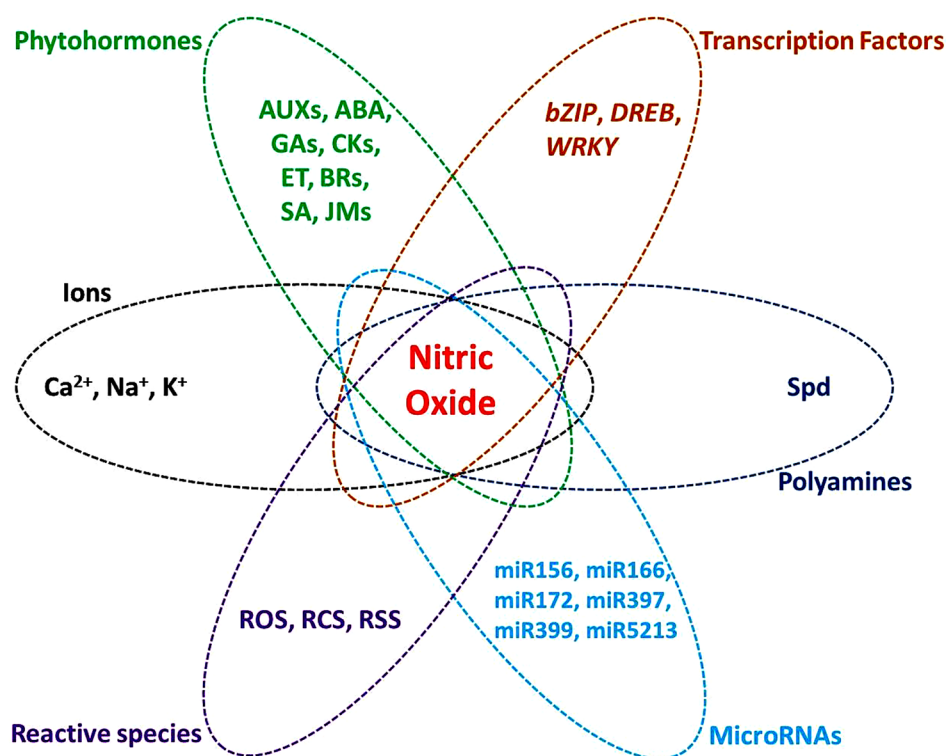
Like the reactive species and secondary messengers, NO also interacts with the phytohormones such as ABA, AUX, cytokinins (CKs), ethylene (ET), gibberellins (GAs), BRs, salicylic acid (SA), γ -aminobutyric acid and jasmonates during different plant metabolic and physiological conditions and abiotic stress responses (Asgher et al. 2017). Studies have shown the signalling pathways of NO and AUX being inter-related with each other during abiotic stress responses and during the root growth and development (reviewed by Sanz et al. 2015). NO improved Cd-tolerance in *M. truncatula* roots via reducing the oxidative damage, enhancing the ion uptake (K⁺ and Ca⁺) and maintaining the AUX equilibrium (Xu et al. 2010). Santos et al. (2020) studied ABA-NO-AUX interactions in roots of *Solanum lycopersicum*, and observed

ABA and molybdenum dependent enzymes as decisive factors for the salt stress driven NO production. The study also highlighted the integration of NO in the ABA–indole acetic acid (IAA) signalling network during salt stress response along with the ABA mediated and NO-dependent activity of the antioxidant system (Santos et al. 2020). Cd and As treatments decreased the endogenous NO concentration in roots of *Oryza sativa* along with high rate of H₂O₂ formation and altered AUX biosynthesis (Piacentini et al. 2020). The SNP-formed NO proved to mitigate the Cd/As toxicity by reducing the H₂O₂ production and induced expression of *AUX1* (IAA–influx carrier) (Piacentini et al. 2020). Application of SNP in the root systems of *Oryza sativa* under Cd/As stress resulted in the alleviation of IAA and reduced H₂O₂ generation, leading to the increased lateral root production and NO generation, ultimately mitigating the toxic impacts of these toxic heavy metals (Piacentini et al. 2020). Figure 3 highlights the interplay between the NO and phytohormones, TFs, miRNAs, reactive species and ions.

GAs are known to be involved in various physiological and/or developmental processes in plants including seed germination, leaf expansion, stem elongation, initiation of flowering and development of fruits. Several reports have supported the possible interplay between GAs and NO (Asgher et al. 2017). In *Arabidopsis thaliana*, antagonistic interaction between the NO and the GA resulted in the regulation of root growth during fluctuating levels of phosphorous (Wu et al. 2014). Lozano-Juste and León (2011) studied the crosstalk between NO and GA involving the phytochrome-interacting factors (PIFs) and GA-regulated DELLA proteins; where NO synchronise with PIF genes, increases the DELLA proteins content, and promotes photo-morphogenesis. Beside GAs, CKs also play important roles in the plant developmental processes (such as chloroplast biogenesis, leaf senescence and cell division) (Fahad et al. 2015). Both synergistic and antagonistic interactions have been observed during NO and CKs crosstalk (Asgher et al. 2017). The effectiveness of CKs in photosynthesis processes was studied under drought stress in *Zea mays*, which revealed the CK-stimulated conversion of excitation energy to electron transfer due to the probable interaction between NO and plastoquinone. Results also indicated CK-induced NO formation apparently via NR (Shao et al. 2010). Further, Liu et al. (2013) investigated the interactions of CK and NO in *Arabidopsis* under metal stress, where CK suppressed the action of NO probably via direct interaction between them, resulting in the overall reduced levels of endogenous NO levels.

It is well-established that ABA is involved in seed germination, dormancy and abiotic stress responses. The ABA mediated enhancement in NO biosynthesis in guard cells was observed, indicating the strong interplay between ABA and NO for provoking the stomatal closure whenever required (Joudoi et al. 2013). NO regulated the stomatal closure via

Fig. 3 Interplay between the nitric oxide (NO) and phytohormones (*AUXs* auxins, *ABA* abscisic acid, *GAs* gibberellins, *CKs* cytokinins, *ET* ethylene, *BRs* brassinosteroids, *SA* salicylic acid, *JMs* jasmonates), transcription factors, Polyamines (*Spd* spermidine), microRNAs, reactive species (*ROS* reactive oxygen species, *RCS* reactive carbonyl species, *RSS* reactive sulfur species) and ions



modulating the K^+ and Ca^{2+} in guard cells (reviewed by Asgher et al. 2017). ABA-induced NO synthesis pathway was reported in *Z. mays* leaf with increased concentrations of H_2O_2 inhibited by the NADPH oxidase inhibitor, called diphenylene iodonium (DPI), and activation of NO mechanisms for the maintenance of cell homeostasis (Tossi et al. 2009). Various studies have been done in *Arabidopsis* to study the crosstalk between NO and ABA or ET. In *Arabidopsis*, MoCo-sulfurase ABA3 enzyme converts the de-sulfo MoCo form (cofactor of NR enzyme) to sulfo MoCo form (cofactor of aldehyde oxidase) for the production of NO and ABA in the plants (Arc et al. 2013). H_2O_2 is produced endogenously via ABA signalling, leading to the generation of NO for the regulation of stomata movements (Bright et al. 2006). Increased UV-B treatment can lead to the production of ROS and RNS and induction of various signalling pathways during the stress responses in plants; increased production of ABA causes the generation of NO and H_2O_2 , leading to the maintenance of cellular homeostasis and cellular damage alleviation (Prakash et al. 2019).

ET is known to play major roles in fruit ripening, flower senescence, and adventitious root hair formation where crosstalk of NO and ET is involved antagonistically during stress conditions (Wei et al. 2020). During abiotic stresses, NO signalling is recognised by S-nitrosation regulated by the production of ET, leading to the antagonism in various plant developmental and physiological processes such as Cd-induced cell death, stomatal opening, and UV-B

induced stomatal closure (Kolbert et al. 2019). The S-adenosyl-methionine (S-AdoMet), precursor of ET and polyamine; is involved during the negative feedback regulation between ET and polyamine dependent synthesis of NO (Arc et al. 2013); ABA signalling and NO synthesis is regulated by the Cu-amine oxidase enzyme. In *Arabidopsis thaliana* and *Cucumis sativus*, up-regulation of genes such as *AtFIT* (TF), *AtFRO2* (ferric reductase), and *AtIRT1* (iron transporter) during cross-talk of NO and ET was observed in acquisition of iron (García et al. 2010). SA is widely known for its involvement in the immune responses under various abiotic stress responses showing both synergistic and antagonistic effects (Asgher et al. 2017). Key roles played by SA and NO signalling in plant growth and physiological processes beside environmental stress responses, have always attracted the scientific community. The non-expresser of pathogenesis related (NPR) gene signalling is mediated by SA and NO during the stress interactions via switching the SA and NO biosynthesis through Ca^{2+} signalling (Rai et al. 2020). SA/NO-mediated heat stress resistance and stress memory has been studied using epigenetically modified plants for the regulations of cellular immunity, leading to the activation of various heat stress-responsive TFs and genes for the better understanding of signalling cascades (Rai et al. 2020). In *Vigna angularis*, the exogenous application of SA triggered the interactions between SA and NO, which resulted into the enhanced photosynthesis, coupled with upregulation of amino acid and carbohydrates as well as

antioxidant enzymes (SOD, CAT), and ultimately improved the salt stress tolerance (Ahanger et al. 2020). Wang et al. (2020a) observed SA-induced stomatal closure, controlled by ROS and NO generation, which was mediated by ET.

Similarly, NO also interact with BRs and play key role in plant stress responses. Zhou et al. (2018) observed that the BR treatment increased NO accumulation which resulted the reduction of virus accumulation in Arabidopsis plants. Furio et al. (2019) in an attempt to evaluate the protective effects of 24-epibrasinolide (EP24) and a formulation based on a brassinosteroid spirostanic analogue DI-31 (BB16) against the fungal pathogen *Colletotrichum acutatum*, the causal agent of anthracnose disease, and the treatment with both the BR compounds induced a defence response in strawberry plants which was mediated through enhanced NO levels. In another interesting study, BR treatment of water-stressed *Z. mays* leaves induced the NO levels and upregulated the expression levels of ABA pathway gene, which enhanced the water stress tolerance (Zhang et al. 2011). Further, Arfan et al. (2019) reported that H₂O₂ and NO crosstalk mediates BR-induced cold stress tolerance in *Medicago truncatula*.

Nitric oxide and miRNA signalling

Non-coding small RNAs, particularly miRNAs play important regulatory roles at transcriptional and/or post-transcriptional levels in plant responses and adaptation to biotic and abiotic stresses (Shriram et al. 2016; Xu et al. 2019; Zhou et al. 2020). The miRNAs are being extensively investigated owing to their direct effect on sequence complementary based degradation of target mRNAs that often encode TF families. The roles of both miRNAs and NO under environmental stresses have been observed in hormone-dependent pathways (Singh et al. 2017). During stress, NO usually targets mitochondrial and chloroplast complexes and the proteins present in them by participating in signal transduction, regulation of Ca²⁺ and TFs (Singh et al. 2017). Recent studies have shown the interaction between biosynthesis of miRNAs and actions of post-transcriptional modifications (Iki et al. 2018; Szweykowska-Kulinska and Jarmolowski 2018; Wang et al. 2018). Chromatin remodelling factor CHR2 acts on a post-transcriptional level in *Arabidopsis* to inhibit formation of miRNA (Wang et al. 2018). The primary-miRNAs (pri-miRNAs) have been observed to undergo post-transcriptional splicing. Factors like stabilized1 (STA1) and GLYCINE-RICH RNA-BINDING PROTEIN (GRP7), etc. in *Arabidopsis* and barley are involved with the splicing of pri-miR168a, pri-miR162a, pri-miR842 (Stepien et al. 2017).

During drought stress, ABA signalling is heightened which results in activation of SnRK2 leading to inactivation of channels regulating efflux in the cell, and triggering

H₂O₂-mediated activation of NO. ABA also mediates inositol 1,4,5-trisphosphate (IP₃) to increase Ca²⁺ influx causing NO production. As seen earlier NO plays an important role in regulating protein kinases like CDPKs, MAPKs leading to an indirect control on genes responsive to ABA signalling. These genes have an interactive network with the miRNAs as seen with miR393, miR160, miR168 from *Arabidopsis*, poplar, rice and *Brachypodium* plants and that play important roles in stress responses via upregulating the expression of their targets, ABA-related genes via auxin response factors (ARFs) and auxin receptor transport inhibitor response I (TIR1), involved in cellular homeostasis (Bright et al. 2006; Courtois et al. 2008; Neill et al. 2008; Singh et al. 2017; Prakash et al. 2019). These are involved in AUX signalling, down-regulation of stress-related genes, and regulating plant growth and development (Bright et al. 2006; Courtois et al. 2008; Neill et al. 2008; Singh et al. 2017; Prakash et al. 2019). In Alfalfa, it was observed that exogenous application of NO down-regulated miRNAs like mtr-miR156a, mtr-miR399a, mtr-miR399c, mtr-miR399q, and mtr-miR5213-5p, targeting SPL genes (SQUAMOSA promoter binding like proteins genes), phosphate transporter genes (miR399a, c, q) and disease resistant protein gene, respectively, playing key role in drought tolerance showing a positive regulation and increasing plant growth and development under drought conditions (Zhao et al. 2020).

NO production and synthesis depend on the temperature of the environment in which the plant coexists. During cold, NO production is stimulated in some plants by activation of ion channels leading to sequential activation of inducer of CBF expression (ICE) protein and C-repeat binding factor/Dehydration-responsive element binding-1 (CBF/DREB1) TFs. The TFs are known to be regulated by different miRNA candidates like miR397, miR166, miR172, and some ABA signalling regulators like miR168 (Kumar 2014). The CBF/DREB factors are shown to work via binding to C-repeat elements/low-temperature response elements (CRT/LTRE) regulating the cold-responsive genes involved in giving cold stress tolerance to plants (Kumar 2014). In plants including *Arabidopsis*, *Brachypodium*, *O. sativa*, *Lotus japonica*, *Pisum sativa*, and poplar, these miRNAs (miR166, miR168, miR172, miR397) were found to be NO-responsive and were attributed for conferring cold tolerance to these plants (Kumar 2014; Singh et al. 2017; Prakash et al. 2019).

The interplay between oxidative stress and miRNAs are well known. The most common miRNA with their targets involved during stress are miR171—GRAS TF (Huang et al. 2017), miR396—Growth Regulating Factor (GRF) (Chen et al. 2015a, b; Yuan et al. 2019), miR159—SIMYB TF genes (López-Galiano et al. 2019), miR474—proline dehydrogenase (PDH) (Wei et al. 2009), miR528—polyphenol oxidase (PPO) (Zhu et al. 2020), observed in various plants like *Arabidopsis*, *Z. mays*, *O. sativa* and are up-

down-regulated. It has been observed that cellular ROS production and accumulation leads to enhanced NO-production, contributing to the activation of stress-responsive genes and ROS scavenging enzymes. The specific miRNA like miR398 regulating SOD is involved in the down-regulation of copper chaperone during Cu^{2+} stress (Singh et al. 2017; Prakash et al. 2019). The miRNA family belonging to miR399 is also involved in the regulation of stress-responsive genes during mineral deficient conditions like phosphate starved environment inhibiting UBC24 in roots of *Arabidopsis* (Fujii et al. 2005; Aung et al. 2006; Bari et al. 2006; Park et al. 2018). The absence of above-mentioned miRNAs contributes to NO production, as they regulate the oxidative stress. During nutrient stress like lower concentrations of phosphate, sulphate, ferrous, etc., the plant cell undergoes anaerobic conditions due to imbalances in the catabolic and anabolic pathways causing increased production of NO, which sequentially activate genes responsive to stress-causing feedback activation of such miRNAs that regulate these genes. (Singh et al. 2017; Prakash et al. 2019).

Biotechnological manipulations of NO and its interactions for engineering abiotic stress tolerance in plants

Biotechnological manipulations of metabolite pathways and their applications have opened new vistas of research in plant abiotic stress tolerance and crop improvement. The studies aimed on interactions of NO with other signalling molecules, phytohormones, and reactive species have revealed the crosstalk mechanisms in plant growth and physiological developments. Studies on the exploration of NO interactions during stress responses and tolerance in crop and model plants have suggested the use of various mutants for the production of stress-tolerant plants. Lechón et al. (2020) developed *cue1* NO overproducer mutants with enhanced arginine content in the roots of *A. thaliana*. Similarly, *cue1-1*, *cue1-5*, *cue1-6*, and *nox1* alleles resulted in the accumulation of NO during seed germination, primary root elongation, and salt stress resistance revealing potential roles NO play in plant physiological developments. When plants are exposed to external environmental stimuli, they synthesize/upregulate peptides for maintaining plant growth and development. The CLAVATA3/Embryo Surrounding Region-Related (CLE) peptides are essentially involved in the process of closing stomata. Zhang et al. (2019) showed that CLE9 acts as an essential regulator in the induction of stomatal closure. Overexpression and loss of function of CLE9 resulted in enhanced resistivity and sensitivity in *Arabidopsis thaliana* during drought stress via increased NO/ H_2O_2 generation and ABA signalling (Zhang et al. 2019). Further, CLE9 peptides upregulated the H_2O_2 and

NO synthesis associated with stomatal closure, which was stopped in the NADPH oxidase-deficient mutants or NR mutants, respectively. Collectively, these results indicated the role of CLE9 in the regulation of stomatal apertures, and in turn stress acclimatisation of plants (Zhang et al. 2019).

The haemoglobin 1 (Hb1) proteins play an essential role in the plant environmental stress responses. The overexpression of *Nicotiana tabacum* non-symbiotic class 1 haemoglobin gene (NtHb1) in *A. thaliana* resulted in NO-scavenging, reduction in ROS/NO accumulation, and increase in antioxidant enzyme activity, Cd export transporters (PDR8), and $\text{Ca}^{2+}/\text{H}^+$ exchangers (CAXs) during Cd stress (Bahmani et al. 2019). Nitric reductase enzyme is primarily involved in the NO signalling mechanisms, following NO crosstalk with anthocyanin, Li et al. (2020) studied NO generation in lyceum fruits by cloning a *Lyceum barbarum* NR gene leading to inhibition of anthocyanin synthesis and enhancement of proanthocyanin accumulation. During salt stress, modulations in 1-aminocyclopropane-1-carboxylic acid (ACC)-oxidase enzyme activity in sunflower seedling resulted in the regulation of lateral-root formation via ACC-oxidase-NO complex formation and reduced ethylene biosynthesis, leading to the development of enhanced salt stress tolerant plants (Singh and Bhatla 2018). S-nitrosoglutathione reductase (GSNOR) enzyme plays a putative role in NO homeostasis and is primarily involved in plant biotic and abiotic stress responses. The overexpression of *GSNOR* in tomato plants resulted in increased sodic alkaline stress tolerance via *GSNOR* regulated NO signalling activation and ROS scavenging efficiency (Gong et al. 2015). Suppression of *GSNOR1* enzyme in plants by S-nitrosylation via NO generation revealed the NO scavenging modulation of *GSNOR1* activity (Frunghillo et al. 2014). Induction of Delta-1-pyrroline-5-carboxylate synthase 1 (P5CS1) gene resulted in NO-dependent signalling between proline and PAs during abiotic stress responses (Shi and Chan 2014). Two mutants, *nia1nia2* (nitrate reductase [NR]-defective double mutant) and *Atnoa1/rif1* (nitric oxide associated1/resistant to inhibition by fosmidomycin1) resulted in overproduction of NO, upregulation of P5CS1, and down-regulation of proline dehydrogenase (ProDH) genes during cold stress and acclimation (Zhao et al. 2009). These results thus clearly demonstrated that NR-dependent NO production play critical role in cold stress tolerance mediated through proline accumulation in *Arabidopsis*.

Beside abiotic stress responses and tolerance attributed to NO manipulations, plant-insect interactions are being studied to explore the NO interactions for conferring biotic stress tolerance in plants. Xu et al. (2020) reported that NO boosted the *Bemisia tabaci* performance via suppressing the JA signalling pathway in tobacco plants. The results suggested that the NO signalling got activated by *B. tabaci* infestation, NO was found to be involved in

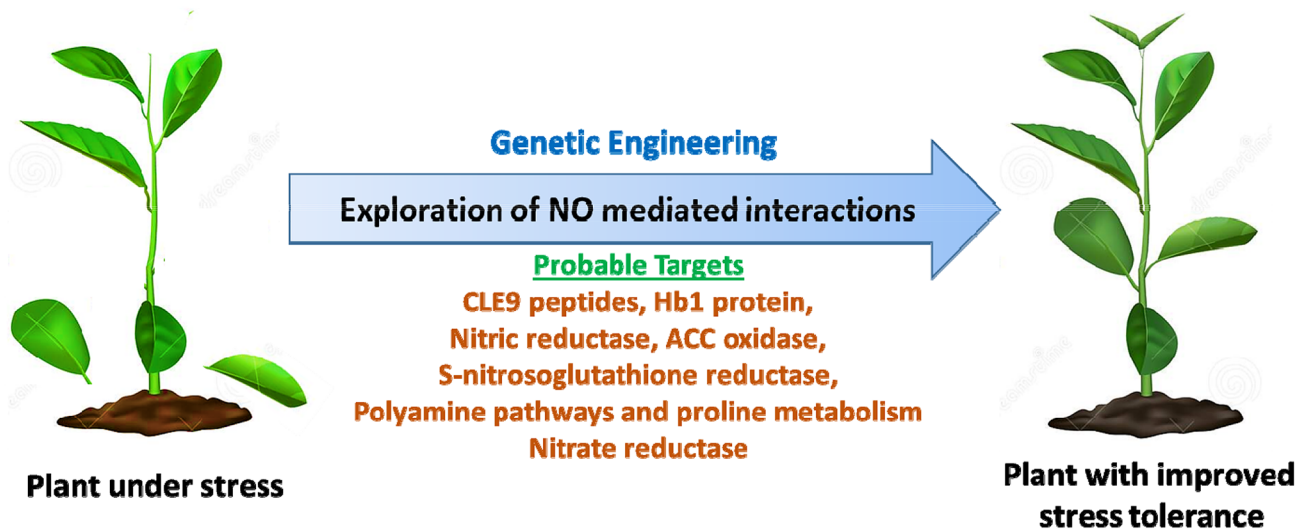


Fig. 4 Exploration of nitric oxide (NO) mediated interactions in for engineering stress tolerance in plants. (*CLE9* CLAVATA3/ESR (CLE)-related protein 9, *Hb1* hemoglobin 1, *ACC* 1-aminocyclopropane-1-carboxylic acid) [The probable targets were presented based

on the literature: Zhang et al. (2019), Bahmani et al. (2019), Li et al. (2020), Singh and Bhatla (2018), Gong et al. (2015), Rodríguez-Ruiz et al. (2019), Frungillo et al. (2014), Shi and Chan (2014), Zhao et al. (2009)]

the suppression of JA-dependent plant defence, and, consequently, NO improves *B. tabaci* performance on host plants (Xu et al. 2020). *S*-nitrosogluthathione reductase (GSNOR) was silenced in *Arabidopsis thaliana* for investigating the plant–pathogen interactions, and it resulted into the increased expression levels of the NO and *S*-nitrosothiol (Rustérucchi et al. 2007; Rodríguez-Ruiz et al. 2019). Figure 4 represents the approaches for exploration of NO-mediated interactions in plants for developing stress-tolerant plants, while Table 2 enlists events of developing transgenic plants via manipulating the NO pathways for enhanced abiotic stress tolerance.

Conclusion

After 40 years from identification of NO in plant cells, extensive research on this dynamic reactive and signalling molecule has opened up various functional capabilities in plant growth and development. Being so dynamic and versatile, NO provides a great scope for its application in crops for various purposes using different approaches to increase the plant's efficiency and tolerance under stress

conditions. NO crosstalk between different signalling molecules has given us glimpse on involvement of NO in process such as stomata closure, salinity tolerance, cGMP pathway, seed germination, beside others. Interactions between NO and plant hormones like GA, AUX, CKs and BRs have shown us influence of NO in photo-morphogenesis, leaf senescence, root growth and photosynthesis, although deeper and more comprehensive research is needed for interpreting whether it is because of direct involvement of NO or RNS and to know the exact mechanism of action. Interaction of NO with different signalling molecules, and stress regulators have shown potential role of NO pathway in crop improvement and creating stress resilient crops. Accordingly, overexpression studies have shown promising results for future crop improvement programme. Interplay between NO and miRNA has recently gained attraction among researchers. Feedback activation and regulation of genes controlled by miRNA under presence of NO has potential for keeping cellular homeostasis in cell and ultimately using it crop improvement.

Table 2 A list of transgenic plants developed via manipulating NO pathways for enhanced abiotic stress tolerance

Gene of interest	Source organism	Transgenic plant	Description	References
No synthase (<i>NOS</i>)	Rat	<i>Oryza sativa</i>	Overexpression resulted in increased NO levels, higher salt and drought tolerance	Cai et al. (2015)
<i>NOS</i> -like gene	<i>Ostreococcus tauri</i>	<i>Arabidopsis</i>	Transgenic plants showed higher NO content and increased tolerance to stresses including salinity, drought, and oxidative stress	Foresi et al. (2015)
<i>PRMT5C125S</i> transgene, carrying a non-nitrosylatable mutant <i>Cys-125</i>	<i>Arabidopsis</i>	<i>Arabidopsis prmt5</i> mutant	Rescued the developmental defects, but not the stress hypersensitive phenotype and the responsiveness to NO during stress responses	Hu et al. (2017)
<i>MdHY5</i>	<i>Malus hupehensis</i>	Orin' apple	Increased nitrate reductase activities, anthocyanin accumulation and nitrate contents	An et al. (2017)
S-adenosylmethionine decarboxylase	<i>Cynodon dactylon</i>	<i>Eremochloa ophiuroides</i>	Improved cold tolerance through involvement of H ₂ O ₂ and NO signalling	Luo et al. (2017)
Glutathione synthetase (<i>OsGS</i>)	<i>Oryza sativa</i>	<i>Oryza sativa</i>	Improved stress tolerance and productivity in transgenic rice	Park et al. (2017)
<i>OrNOS</i>	<i>Ostreococcus tauri</i>	<i>Nicotiana tabacum</i>	Accumulation of higher NO levels compared to siblings transformed with the empty vector, and displayed accelerated growth in different media containing sufficient nitrogen availability	Nejamkin et al. (2020)
S-nitrosogluthathione reductase	<i>Spinacia oleracea</i>	<i>Nicotiana tabacum</i>	lower malondialdehyde content, reactive oxygen species and higher activities and transcript level of superoxide dismutase, catalase, peroxidase under nitrate stress	Wang et al. (2020b)
<i>S1WRKY81</i>	<i>Solanum lycopersicum</i>	<i>Solanum lycopersicum</i>	Transcription factor inhibited the stomatal closure by attenuating nitric oxide accumulation in the guard cells	Ahamed et al. (2020)
Glucose-6-phosphate dehydrogenase (<i>G6PD</i> or <i>G6PDH</i>)	<i>Glycine max L</i>	<i>Arabidopsis thaliana</i>	Increased salinity tolerance by decreasing ROS accumulation	Jin et al. (2021)
<i>NR1</i> and <i>NR2</i>	<i>Arabidopsis</i>	<i>Arabidopsis</i>	Increased NO content	Costa-Broseta et al. (2021)

Author contribution statement XZ and VK contributed substantially to the conceptualization and designing of the review. SJ, SP, TK, JS drafted the text and final version, XZ and VK edited and approved the final version. All the authors contributed significantly, seen and approved the manuscript.

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Declarations

Conflicts of interest The authors declare no conflict of interest.

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