

Review

Dual action of the active oxygen species during plant stress responses

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Abstract. Adaptation to environmental changes is crucial for plant growth and survival. However, the molecular and biochemical mechanisms of adaptation are still poorly understood and the signaling pathways involved remain elusive. Active oxygen species (AOS) have been proposed as a central component of plant adaptation to both biotic and abiotic stresses. Under such conditions, AOS may play two very different roles: exacerbating damage or signaling the activation

of defense responses. Such a dual function was first described in pathogenesis but has also recently been demonstrated during several abiotic stress responses. To allow for these different roles, cellular levels of AOS must be tightly controlled. The numerous AOS sources and a complex system of oxidant scavengers provide the flexibility necessary for these functions. This review discusses the dual action of AOS during plant stress responses.

Key words. Abiotic stress; active oxygen species; biotic stress; catalase; oxidative stress.

Introduction

The last decade has seen many new and exciting findings paving the way to a better understanding of plant stress responses. Some of the major areas of focus include the role of antioxidants and active oxygen species (AOS) during biotic and abiotic stresses. Although both types of stress condition are associated with the regulation and potential signaling role of AOS, there has been little overlap between these two fields of research. Understanding the role of AOS during pathogenesis has progressed greatly since the early reports of their involvement in plant disease resistance. It is now clear that AOS play several crucial roles during pathogenesis. They are involved in the hypersensitive response typical of plant-pathogen incompatible

interactions [1–7]. They can also limit the spread of pathogen infection by strengthening plant cell walls and/or by killing pathogens directly [8–12]. Finally, they also act as signaling molecules during local and systemic acquired resistance [5, 9, 13].

The role of AOS during pathogenesis has been extensively reviewed over the years and there are several excellent reviews on the topic [14–18]. As a result, this review will mainly focus on the role of AOS during abiotic stress responses.

The role of AOS during abiotic stress has become a subject of considerable interest given that AOS have been implicated in processes leading to plant stress acclimation [19–24]. High concentrations of AOS can lead to phytotoxicity whereas relatively low levels can be used for acclimatory signaling. This novel finding means that AOS are not simply toxic by-products of metabolism but also function as signaling molecules [24,

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25]. Therefore, the controlled modulation of AOS levels in planta is extremely important and may in part explain complex processes such as cross-tolerance.

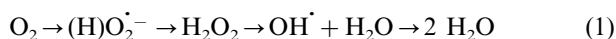
The role of H₂O₂ during plant stress responses may be difficult to discern from that of other AOS. The reactive nature of these molecules makes it extremely difficult to distinguish effects linked to different species. Several methods are available for measuring AOS in planta but the technical difficulty associated with quantifying endogenous levels of these very reactive and short-lived species means that few studies have measured AOS during plant stress. Over the years, most of the evidence for increases in AOS has been provided through the study of antioxidants or other secondary processes such as lipid peroxidation.

Living with oxygen

Biochemical properties of AOS

Oxygen is necessary for plants. The reduction of O₂ to H₂O provides the energy that allows the impressive complexity of higher organisms but its reduction is a mixed blessing. When incompletely reduced, AOS can be extremely reactive and may oxidize biological molecules. All AOS can react with DNA, proteins, and lipids [26].

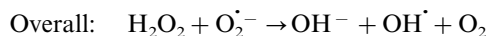
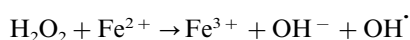
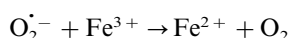
Reduction of molecular O₂ proceeds through four steps, thus generating several O₂ radical species [27]. The reaction chain requires initiation at the first step whereas subsequent steps are exothermic and can occur spontaneously, either catalyzed or not.



The first step in O₂ reduction produces relatively short-lived AOS that are not readily diffusible: hydroperoxyl (HO₂[·]) and superoxide (O₂^{·-}). The half-life for O₂^{·-} is approximately 2–4 μs [28–30]. These oxygen radicals are highly reactive, forming hydroperoxides with enes and dienes [31]. Furthermore, specific amino acids, such as histidine, methionine, and tryptophan can be oxidized by O₂^{·-} [28]. In the cellular environment, O₂^{·-} will also cause lipid peroxidation, thereby weakening cell membranes.

The second O₂ reduction generates hydrogen peroxide (H₂O₂), a relatively long-lived molecule (1 ms) that can diffuse some distance from its site of production [5, 32]. The biological toxicity of H₂O₂ through oxidation of SH groups has long been known and it can be enhanced in the presence of metal catalysts through Haber-Weiss or Fenton-type reactions.

The Fenton or Haber-Weiss reactions are:



The last species generated by this series of reductions is the hydroxyl radical (OH[·]). It has a very strong potential and a half-life of less than 1 μs. As a result, it has a very high affinity for biological molecules at its site of production, reacting at almost diffusion-controlled rates (K > 10⁹ M⁻¹ s⁻¹).

As described above, several AOS can be generated by the reduction of molecular oxygen in plant systems, all with specific biochemical properties making them either extremely reactive or, in contrast, interesting candidates for cell signaling. This dual role for AOS is extremely important and will be discussed in more detail below.

Interactive effects of AOS

From the Haber-Weiss reaction, AOS can obviously react with one another leading to the generation of a third AOS species more toxic than the other two. Similarly, O₂^{·-} may react with other molecules such as lipid peroxides. It can also react with nitric oxide, leading to the formation of peroxynitrite, a species considered less reactive than peroxides [33, 34]. These reactions depend on concentration and on the preferential scavenging capacity of the cell. The plant may favor formation of one or the other species by preferentially scavenging H₂O₂ with antioxidants or, in contrast, accumulate H₂O₂ by activating superoxide dismutase (SOD). In addition, H₂O₂ accumulation may itself lead to higher AOS production. The H₂O₂ disruption of photosynthesis (see below), as well as the probable direct role of H₂O₂ in the activation of an NAD(P)H-dependent oxidase [35–37], suggests that H₂O₂ itself can stimulate AOS accumulation.

Sources of AOS in plants

There are several known sources of AOS in plants. Some of the most common include the leakage of electrons to O₂ from electron transport chains in the chloroplast and mitochondria. Other sources include photorespiration in the peroxisomes, and cell wall oxidases and peroxidases.

Photosynthesis

The chloroplasts are considered the most powerful source of AOS in plants [38]. Photosynthesis is a combination of successive redox reactions during which light energy is absorbed by the light-harvesting complexes and transferred to the reaction centers of the photosystems. Electrons are then passed on to a final electron acceptor, generally CO₂ (fig. 1). However, because very few plants have rates of CO₂ fixation high enough to

use more than 50% of the light energy absorbed [39], alternative acceptors are used, including oxygen. Conditions limiting CO₂ fixation will inevitably lead to an enhancement of AOS accumulation, as more O₂ molecules will be used as electron acceptors.

The photosynthetic electron transport chain includes a number of enzymes on the reducing (acceptor) side of photosystem I (PSI): Fe-S centers, reduced thioredoxin, and ferredoxin [40]. These electron transport components are auto-oxidizable and under conditions limiting the availability of NADP⁺, O₂^{•-} can be formed [31, 41]. Superoxide can then initiate a chain reaction leading to the production of more toxic radicals. The sum of the Mehler peroxidase reaction is therefore the transfer of electrons from H₂O to O₂, resulting in production of O₂ at PSII, O₂^{•-} at PSI, and the trans-thylakoid proton gradient necessary to drive phosphorylation and photochemistry of PSII [40, 42, 43].

Although production of AOS during photosynthesis is often considered detrimental during stress conditions,

under normal conditions, the ability of O₂ to accept excess electrons prevents overreduction of the electron transport chain, thereby reducing the risk that activated singlet oxygen is formed and minimizing energy exchange between excited chlorophyll and molecular oxygen [40].

Photorespiration

There is another route by which H₂O₂ can be produced during photosynthesis (fig. 2) [44]. During carbon assimilation, ribulose-1,5-bisphosphate carboxylase uses CO₂ to carboxylate ribulose-1,5-bisphosphate. However, ribulose-1,5-bisphosphate carboxylase can also use O₂ to oxygenate ribulose-1,5-bisphosphate [43]. Oxygenation yields two glycolates that are then transported from the chloroplasts to the peroxisomes. There, glycolate oxidation is catalyzed by glycolate oxidase yielding H₂O₂. In addition, the microbodies contain fatty acid β-oxidase and xanthine oxidase as H₂O₂- and O₂^{•-}-producing enzymes, respectively [45].

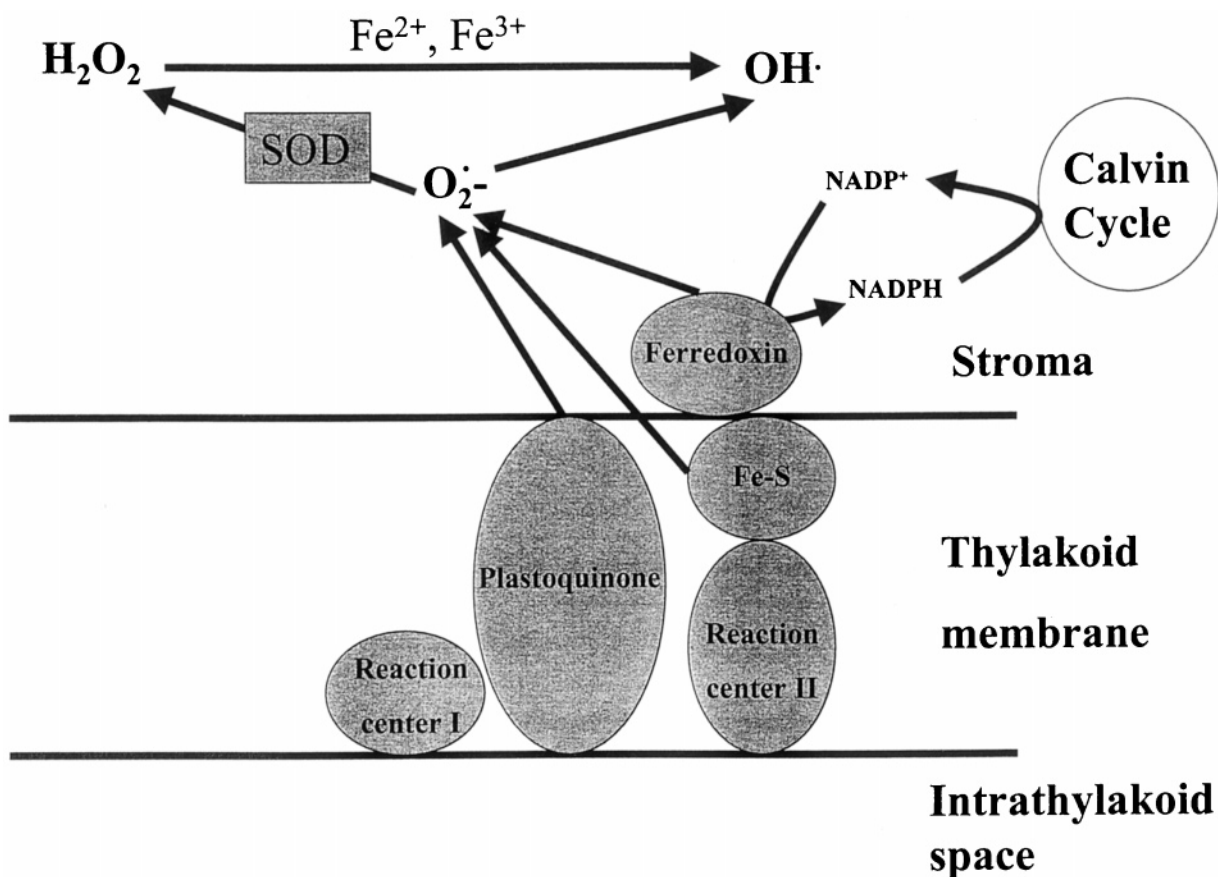


Figure 1. Schematic representation of the chloroplast showing the main sources of oxygen radicals. SOD, superoxide dismutase. Fe-S, iron sulfur cluster.

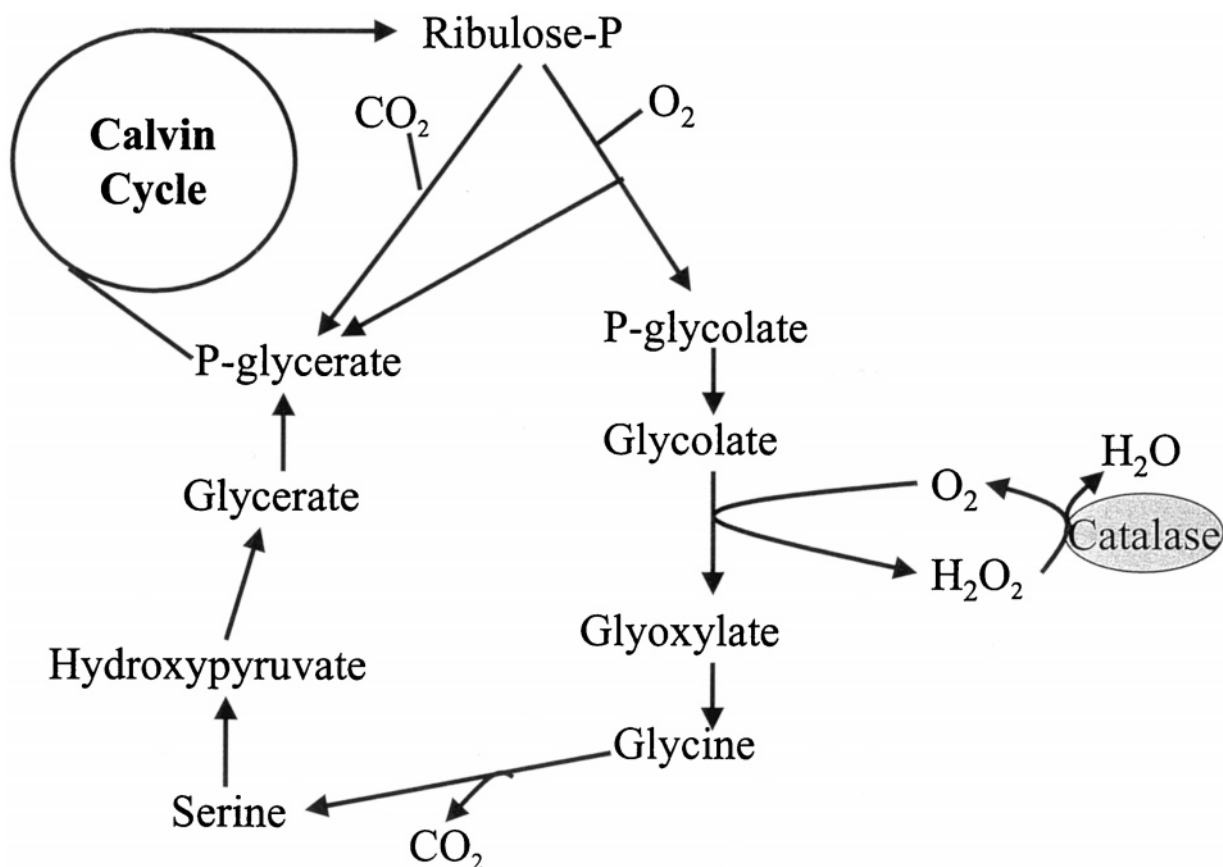


Figure 2. The photorespiratory cycle and the main source of H₂O₂ in the peroxisomes.

Respiration

Although AOS production in plant mitochondria has received little attention in the past, recent evidence suggests that mitochondria are an important source under specific conditions [45]. The mitochondria electron transport chain is made of several dehydrogenase complexes that reduce a common pool of ubiquinone (fig. 3) [46, 47]. The ubiquinone pool is then oxidized by either the cytochrome or the alternative pathway. In general, the main O₂⁻ generators in the mitochondria are the ubiquinone radical and NADH dehydrogenases [26]. Superoxide is formed by the auto-oxidation of the reduced components of the respiratory chain. For example, H₂O₂ production in isolated pea stem mitochondria is mainly formed during substrate oxidation at the level of complex I [48].

Studies on the regulation of the alternative oxidase (AOX) pathway, a substitute for the cytochrome pathway, consisting of a single homodimeric protein, have provided additional evidence for mitochondrial AOS production during stress conditions [for a review, see ref. 49]. Transgenically cultured tobacco cells that are

altered through antisense suppression of AOX accumulate higher level of AOS than wild-type cells [50]. In contrast, overproduction of AOX results in cells with lower AOS abundance. Similarly, use of inhibitors (salicyl hydroxamate and propyl gallate) to block the AOX leads to H₂O₂ accumulation in mitochondria of soybean and pea cotyledons [51]. The observation that AOX can be induced by various environmental stimuli makes mitochondrial AOS production an important component of stress responses [49, 52].

NAD(P)H-oxidase

In recent years, the source of AOS during the oxidative burst of plant-pathogen interactions has been the focus of numerous studies. Doke [53] demonstrated that an NADPH-dependent O₂⁻ production played a part in the oxidative burst in both elicitor-treated potato protoplasts and microsomes infected with *Phytophthora infestans*. Pharmacological and immunological studies confirmed its presence in various plant species, suggesting its widespread role in plant defense [13, 54, 55].

Cross-reactivity of antibodies against components of the mammalian multiprotein complex that are homologous with proteins from various plant model systems helped identify p22^{phox}, p47^{phox}, p67^{phox}, gp91^{phox}, and p21^{rac}-like proteins [56–62]. Furthermore, several of the above plant homologues have been identified as expressed sequence tag cDNA sequences. However, questions remain regarding the precise role and importance of such a system [63]. In focusing on the NADPH-oxidase, other sources, such as pH-dependent peroxidases, may have been overlooked and their importance may become evident in the future. In addition, the potential role of NADPH-oxidase during other stress responses has been suggested [19, 37, 64], although further work is needed.

Other sources

Multiple AOS-producing systems are found in plants, some of which may be developmentally and/or environmentally regulated. However, many still await further characterization.

Cell wall peroxidases produce O_2^- at the expense of NADH in a Mn^{2+} -dependent reaction [45]. The peroxidase-produced H_2O_2 is strongly pH dependent and believed to involve the reduction of the ferrous O_2 conjugate [16]. Most of the H_2O_2 generated during the oxidative burst can be produced by peroxidases [2, 10, 55].

Alternative sources for AOS include a germin-like oxalate oxidase, induced during infection of barley by

Erysiphe graminis [65]. Oxalate oxidase gene expression is also induced by salt stress, salicylate, and methyl jasmonate [66].

Copper-containing amine oxidases can also catalyze the oxidation of various amines yielding NH_3 and H_2O_2 [16]. Flavin-containing oxidase and cell wall amine oxidase and peroxidase produce AOS in epidermal cells of tobacco following induction by cryptogein, a fungal elicitor [67].

An efficient system to regulate AOS

The antioxidant system

The omnipresence of O_2 in the environment and the various cellular locations where AOS are produced render oxidant scavengers necessary for plant growth and survival. The capacity for AOS to serve as signals adds to the importance of antioxidants to specifically regulate different AOS in various cellular locations. Plants have several antioxidant enzymes and metabolites located in different plant cell compartments, the main ones being SODs, a family of metalloenzymes catalyzing the dismutation of O_2^- to H_2O_2 [for reviews, see refs 68–70], catalases (CATs), which are heme proteins that catalyze the removal of H_2O_2 [for reviews, see refs 12, 71], and the enzymes and metabolites of the ascorbate-glutathione cycle that are involved in the removal of H_2O_2 [for a review, see ref. 40].

The majority of enzymes of the ascorbate-glutathione cycle [ascorbate peroxidase (APX), glutathione reduc-

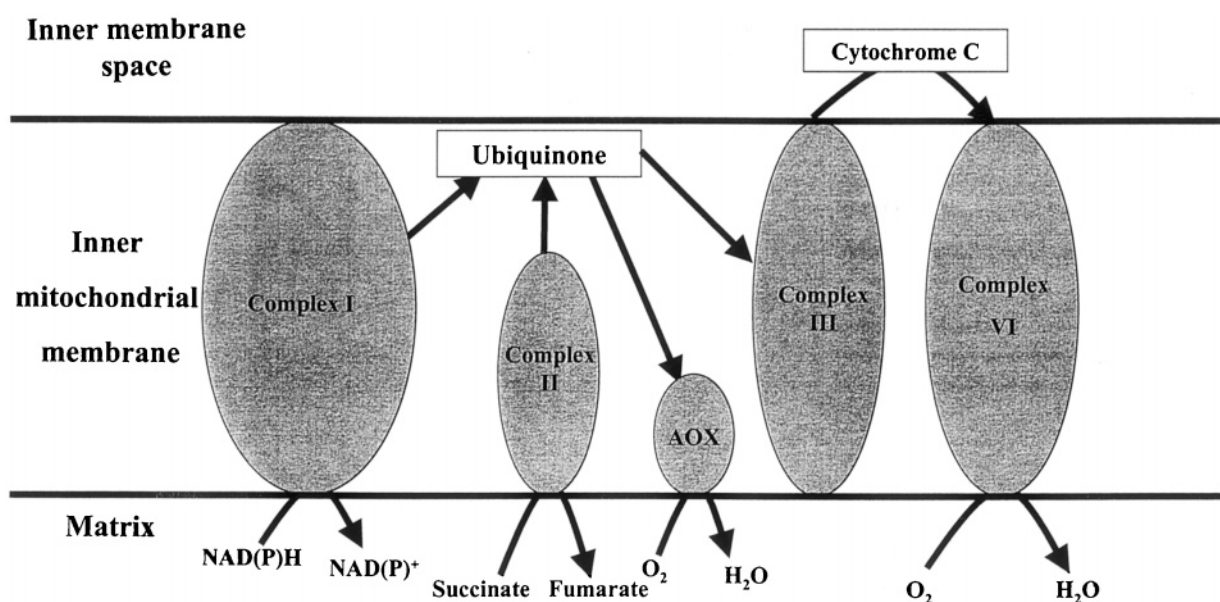


Figure 3. Schematic representation of the plant mitochondria complexes and the electron transport chain. AOX, alternative oxidase.

tase (GR), and dehydroascorbate reductase (DHAR)] have been found in chloroplasts, cytosol, mitochondria, and peroxisomes [40, 68, 72–75]. Catalase and the ascorbate-glutathione cycle are important in H₂O₂ scavenging. Although their properties and requirements are different, they function effectively in parallel. Catalase does not require reducing power and has a high reaction rate but a low affinity for H₂O₂, thereby only removing the bulk of H₂O₂ [32]. In contrast, APX requires a reductant (ascorbate) and has a higher affinity for H₂O₂, allowing for the scavenging of small amounts of H₂O₂ in more specific locations.

Antioxidant capacity is very much dependent on the severity of the stress as well as the species and its developmental stage [76–81]. The capacity of foliar antioxidants also varies with the physiological age of the leaf [82, 83] and its position in relation to irradiance interception [82].

Catalase

Catalases are tetrameric heme-containing enzymes that convert 2 H₂O₂ → O₂ + 2 H₂O, thus protecting the cell from the damaging effects of H₂O₂ accumulation [84]. Catalases have been mainly associated with the removal of H₂O₂ in microbodies [12] and can catalyze either the direct decomposition of H₂O₂ or the oxidation by H₂O₂ of substrates such as methanol, ethanol, formaldehyde, formate, nitrite, or elemental mercury [85]. Catalases are present in peroxisomes, glyoxysomes, and related organelles where H₂O₂-generating enzymes, such as glycolate oxidase, are found. There are three main isoforms: CAT1, CAT2, and CAT3 [32, 71, 85–88]. A basic nomenclature has been adopted to avoid confusion between the isozymes in the different species studied (table 1). The class I catalases include the SU2 of cotton [89], *Cat1* of *Nicotiana plumbaginifolia* [88, 90], *CAT2* of *Arabidopsis thaliana* [91], and *CAT-2* of *Zea mays* [71]. They are highly expressed in leaves, are light dependent, and involved in the removal of H₂O₂ during photorespiration. Class II catalases include the *Cat2* of *N. plumbaginifolia*, *CAT2* of castor bean [92], *CAT-3* of maize, a tomato catalase [93], *Cat2St* from potato [94], and *CAT1* from *A. thaliana* [91]. They are mainly found in vascular tissues. Finally, the class III catalases include *SU1* of cotton, *Cat3* of *N. plumbaginifolia*, *CAT1* from castor bean, *CAT-1* from maize, and *CAT3* from *A. thaliana* [91]. They are mainly involved in the removal of H₂O₂ from glyoxysomes and are highly abundant in seeds and young seedlings [12].

Catalase and plant stress

The expression of catalase genes is not only developmentally regulated but it is also sensitive to various

Table 1. Nomenclature of plant catalases [adapted from ref. 12].

Species	Class I	Class II	Class III
<i>Gossypium hirsutum</i> (cotton)	SU2		SU1
<i>Nicotiana plumbaginifolia</i>	Cat1	Cat2	Cat3
<i>Ricinus communis</i> L. (castor bean)	CAT2	CAT1	
<i>Zea mays</i> (maize)	CAT-2	CAT-3	CAT-1
<i>Arabidopsis thaliana</i>	CAT2	CAT1	CAT3
<i>Lycopersicon esculentum</i> (tomato)		TOMCAT1	
<i>Solanum tuberosum</i> (potato)		Cat2St	

environmental signals [71, 90, 95–97]. The mRNA levels of *CAT2* and *CAT3* of *A. thaliana* [98], *Cat1* of *N. plumbaginifolia* [88], and *CAT-3* of *Z. mays* [99] are regulated by circadian rhythm, whereas others, such as *Cat2* and *Cat3* of *N. plumbaginifolia* and *CAT1* of *A. thaliana*, are not [91].

Catalase is photoinactivated in moderate light under conditions to which plants are adapted [100, 101]. Although they are involved in H₂O₂ scavenging, their turnover being continuous, their steady-state level can be rapidly lowered under any stress conditions in which translation is inhibited or degradation enhanced [100]. The various catalases are, however, differentially responsive to light. *CAT1* and *CAT2* mRNA abundance increases upon illumination, whereas *CAT3* shows a rapid and transient decline [102]. *Cat1* expression in tobacco leaves is unaffected by light [12, 32].

Differential sensitivity of catalases to stress is also observed during temperature extremes. In maize, *CAT-3* is more sensitive to temperature than *CAT-1* [85]. Generally, catalase activity is inhibited under low temperature [103], and heat shock [22, 23, 104, 105]. For example, catalase photoinactivation in rye seedlings was induced after 4 h at 40 °C [104]. *Cat1* and *Cat2* transcript abundance in *N. plumbaginifolia* decreased during a 5-h heat shock at 37 °C, but recovered rapidly after return at 22 °C. In contrast, *Cat3* expression was unchanged by these treatments [12, 88]. Catalases also exhibit differential responses to thermal extremes in maize and *Arabidopsis* [71, 102], and induction of *CAT-3* during cold acclimation in maize is correlated with chilling tolerance [106, 107].

Changes in catalase in response to other stress conditions have also been reported. For example, the three catalases of tobacco all responded to environmental challenges with UV-B, O₃, and SO₂ [88]. As a consequence of its sensitivity to environmental conditions, early loss of catalase was suggested as a signal for antioxidant defenses [100]. However, further studies are needed to confirm such a role.

The importance of catalase inhibition during pathogenesis has been the center of much debate. Catalase inhibition by salicylic acid (SA) was originally proposed as one of the mechanisms by which SA may mediate defense responses in tobacco [108, 109]. Identification and characterization of an SA-binding protein (SABP) with a cDNA sequence highly similar to catalase in tobacco [108, 110–112], cucumber, tomato, *Arabidopsis*, and rice [84, 113] was at the origin of this proposed model. Recently, an additional SABP (SABP2) has been identified and characterized in tobacco [114]. However, several studies have questioned the ability of SA to specifically inhibit catalase in vivo [115–118]. Because the SA concentrations that are required for in vitro inhibition of catalases are extremely high, this mechanism may only occur in restricted subcellular compartments [119]. Cat1- and Cat2-deficient transgenic tobacco plants are not altered in their pathogenesis-related (PR) response under normal growth conditions, which implies that catalase inhibition is not sufficient to generate a signal for activation of PR responses [120]. The Cat1-deficient tobacco plants are also more susceptible to infection [121].

The above discussion clearly demonstrates that the AOS-scavenging system in plants is an essential component of the protective mechanisms against biotic and abiotic stress. Corroborating the correlation between antioxidant potential and stress tolerance [40], free-radical scavenging capacity often increases with plant acclimation [64, 122, 123]. Acclimation to elevated AOS is generally associated with enhanced activity of antioxidant enzymes [124, 125]. This increase in stress tolerance has been exploited. Plants have been transformed to overproduce components of the antioxidant system [126–131]. However, results have not always been conclusive. SOD overproduction, though, shows more encouraging results [132–140]. Are these results related to the capacity of SOD to modulate endogenous levels of $O_2^{\cdot-}$ and H_2O_2 , rather than to scavenge AOS like other antioxidants? Enhancement of the AOS-scavenging system may be sufficient to protect plants from some stress, but such transformation may also be detrimental. The fact that H_2O_2 and $O_2^{\cdot-}$ are involved in mediating cell defense responses is one possible explanation [4, 9, 22, 24, 25, 141]. Enhancing the antioxidant capacity may protect plants under specific conditions but it will also interfere with the signaling cascade involved in plant adaptation. Only recently, several research groups have realized that suppressing AOS accumulation in plants may be a futile approach. Instead, close control of AOS levels in various cellular compartments may offer greater possibilities, but the antioxidant mechanisms involved during stress responses and their complex interactions are only just being elucidated.

Generation of AOS under adverse conditions

Environmental stress is a major factor that limits plant productivity. Although AOS are formed in normal cell metabolism and their regulation is a common cellular event [24, 142], oxidative damage is often associated with plant stress. Most, if not all, abiotic stress conditions are exacerbated by the effect of AOS accumulation [24, 30].

Exposure to high light intensities

High-light exposure is one of the most common sources of oxidative stress in plants. High light-induced AOS generation arises principally when it is combined with additional stress factors in which CO_2 fixation is limiting [38]. Under normal photosynthesis, chloroplasts generate approximately 150–250 μmol of H_2O_2 per milligram of chlorophyll per hour [143]. Under excess light conditions, enzymatic processes for CO_2 fixation become rate limiting and, as a result, photosynthesis produces more NADPH and ATP than necessary. This accumulation of redox and energy equivalents will reduce the plastoquinone pool and/or inhibit the water-splitting complex, inevitably leading to PSII inactivation, so-called photoinhibition [144–147]. An increase in the rate of O_2 photoreduction with high irradiance has been reported for *Vinca major*, *Schefflera arboricola*, and *Mahonia repens* [125]. Exposure of *Arabidopsis* seedlings to excess light ($> 2000 \mu\text{mol photons m}^{-2} \text{s}^{-1}$) for 1 h resulted in photoinhibition, accompanied by H_2O_2 accumulation [148]. When exposed to high light ($> 300 \mu\text{mol photons m}^{-2} \text{s}^{-1}$), leaves of catalase-deficient (Cat1AS) tobacco plants become bleached as a result of H_2O_2 accumulation in the peroxisomes [32]. These results clearly demonstrate that a decline in catalase combined with high light results in oxidative stress.

Expression analysis of antioxidants has provided evidence for the accumulation of AOS during high-light exposure. Alteration in antioxidants after high-light treatment often infers enhancement of AOS accumulation. Tobacco plants exposed to high light showed a strong induction of chloroplastic FeSOD (FeSOD_{chl}) transcripts [149]. The FeSOD_{chl} mRNA increase was not light dependent but directly responsive to oxidative stress, because 3-(3',4'-dichlorophenyl)-1,1-dimethylurea infiltration, which blocks re-oxidation of the primary electron-accepting quinone of PSII, abolished the induction of FeSOD_{chl} mRNA. Activities of most antioxidant enzymes increased in pea seedlings transferred from low to high light [124].

Additional evidence for an involvement of AOS during high-light treatment is provided by the degradation of glutamine synthetase, phosphoglycolate phosphatase,

the large subunit of ribulose-1,5-bisphosphate carboxylase/oxygenase, and an increased content of carbonyl groups in stromal proteins of pea [150]. All these processes result from oxidative damage.

Drought and salt stress

Drought-induced inhibition of photosynthesis leads to increased AOS production in the chloroplasts [30, 151, 152]. Numerous studies report increased oxidative stress during water deficit [153–158]. The accumulation of AOS during such conditions originates mainly from the decline in CO₂ fixation, leading to higher leakage of electrons to O₂. Accordingly, thylakoid membrane electron leakage to O₂ increased in sunflower [159] and in wheat [41] after drought.

Other AOS-dependent changes are often associated with water deficit. For example, lipid peroxidation, a commonly used indicator of oxidative stress, increases in various tissues during drought [81, 160–162]. Changes in antioxidants are also correlated with water deficit [161, 163, 164]. Chloroplastic and cytosolic Cu/ZnSOD activity increase in pea subjected to water deficit [81]. Activities of cytosolic and chloroplastic Cu/ZnSOD and cytosolic APX rise during drought of pea plants [78, 79], and osmotic stress enhances Mn-SOD transcript abundance in maize [165]. Drought-tolerant *Sorghum bicolor* has higher antioxidant capacity during water deficit than drought-susceptible varieties [166]. Drought stress can therefore alter the oxidative balance of the cell and acclimation to drought is generally correlated with keeping AOS levels relatively low through the antioxidant system.

Plant responses to salt stress often include drought-mediated symptoms. As a result, it is not surprising that oxidative stress participates in salt damage. The generation of AOS during salt stress is probably similar to that during drought and is mainly attributed to increased leakage of electrons to O₂, ensuing from a decline in CO₂ fixation. Concentrations of H₂O₂ in shoot tissue of rice significantly increased upon salt stress [167]. Tobacco plants deficient in catalase (Cat1AS) were also more susceptible to salt stress [32].

Further evidence for the involvement of oxidative stress during salt stress comes from the study of antioxidants. Mitochondrial and chloroplastic SOD and APX isozymes increase during salt stress in pea [168, 169]. Various antioxidant enzymes are upregulated upon salt treatment of cotton cell calli [170] and in citrus cells [171]. NaCl treatment activates *Cat2* and *Cat3* genes in hydroponically grown tobacco [172]. Other salt-induced responses include AOS-mediated lipid peroxidation in citrus [173].

Low-temperature exposure

Low temperature is one of the most studied abiotic stresses in plants. Direct evidence for accumulation of AOS during chilling has been reported for several plant species. The chloroplast is a major source of AOS during chilling through inhibition of CO₂ fixation, although, recently, mitochondrial AOS accumulation during chilling has been demonstrated [174–176]. Early results by Okuda et al. [177] showed that cold treatment of winter wheat induces a transient but significant increase in endogenous H₂O₂. Similarly, chilled cucumber plants [103] and maize seedlings [20] accumulate H₂O₂. Exposure to 4°C of callus tissue of *A. thaliana* leads to conditions of oxidative stress [178]. Upon chilling, cellular H₂O₂ increases followed by lipid peroxidation. A correlation between H₂O₂ accumulation and chilling damage has been reported in *A. thaliana* callus tissue [178], cucumber, winter wheat [103, 177], rice [179], and maize [20, 180].

High-temperature exposure

Similar to low temperatures, high-temperature exposure involves AOS accumulation. The mechanisms by which AOS can accumulate under such conditions are diverse, but the prime site of production is probably the chloroplast, through leakage of electron following a decline in CO₂ fixation. Alternatively, an NAD(P)H-dependent oxidase might be activated upon heat treatment [19, 64]. Doke [181] showed that within 15 min after heat shock, potato leaf tissues produced an oxidative burst. More recently, exposure of whole tobacco seedlings to 40 °C for 1 h in the light induced a significant increase in H₂O₂ [24, 182]. A similar accumulation of H₂O₂ after a heat treatment was measured in mustard seedlings [23]. Although these latter studies show direct evidence of H₂O₂ accumulation in planta, other studies have reported changes in the antioxidants during high-temperature treatment.

SOD levels rise in tobacco under high-temperature treatment [149]. Heat protection of wheat is improved by keeping APX and GR activities high [183], and GR activity is significantly enhanced during heat-induced thermoprotection in mustard seedlings [64]. Furthermore, exposure of alfalfa to supra-optimal temperature leads to increases in APX and catalase [184].

Heavy metals

Although stress-related metal toxicity has received little attention in the past, it has become a major issue in reclamation of industrial sites. Copper, aluminum,

cadmium, zinc, and iron damage have all been linked to oxidative stress.

When present in excessive amounts, copper, as other heavy metals, causes uncontrolled redox reactions in the cell resulting in the formation of AOS and lipid peroxidation [185–188]. The differential activation of SODs in tobacco [189] and soybean [190] after Cu treatment suggests AOS accumulation during Cu stress. Other copper/AOS-mediated effects include the expression of PR10 protein in birch roots and leaves [191].

When present in high amounts, heavy metals, such as aluminum, cause AOS-mediated lipid peroxidation [192]. Induction of various AOS-related defense responses has also been reported. Aluminum induces glutathione-*S*-transferase (GST), peroxidase, β -1,3-glucanase, PR2, and phenylalanine ammonia lyase in *Arabidopsis* [193], and several antioxidant transcripts in soybean [194].

Zinc phytotoxicity is exacerbated by AOS. Zinc treatment of *Phaseolus vulgaris* increases the levels of H_2O_2 in roots and lipid peroxidation in primary leaves [195]. Interestingly, zinc deficiency symptoms also implicate AOS. Photo-oxidation of thylakoid constituents and impairment of detoxifying mechanisms are responsible for zinc deficiency symptoms, such as leaf chlorosis [196].

Iron excess can also result in oxidative stress [186]. Iron toxicity is in part due to its potential for reacting with H_2O_2 and $O_2^{\cdot-}$ to generate the more toxic radical OH^{\cdot} . Increased activity of catalase and APX in tobacco seedlings after iron excess supports this model [197]. Iron-induced oxidative stress in maize was proposed as a possible trigger for ferritin mRNA and protein accumulation [198].

UV radiation

By exciting electrons by electronic transition in appropriate photosensitizers, UV radiation induces production of $O_2^{\cdot-}$ [199]. SOD can then dismutate the produced $O_2^{\cdot-}$ to H_2O_2 . Accordingly, H_2O_2 accumulation peaks 60–90 min after UV-C irradiation of *Rosa damascena* cell suspensions [199]. AOS are also implicated in the regulation of transcript levels of photosynthetic and acidic PR proteins in *Arabidopsis* exposed to UV-B [200].

Exposure of *N. plumbaginifolia* to UV-B strongly increases Cat2 and glutathione peroxidase (GPX) mRNA [90]. In contrast, APX and SOD transcript levels remain relatively unaltered. Similarly, SOD transcript levels of *Pisum sativum* decrease after UV-B exposure, whereas those of GR are several-fold higher [201].

Air pollutants

Ozone enters plant tissues through the open stomata, where it reacts with components of the cell wall and plasma membranes to form various AOS [202]. Mehlhorn et al. [203] clearly showed formation of AOS during ozone exposure by using electron spin resonance. Bluegrass and ryegrass [204] and *Arabidopsis* [37] accumulated $O_2^{\cdot-}$ and H_2O_2 , respectively, upon ozone treatment. Recently, O_3 was shown to trigger an oxidative burst by activating an NADPH-dependent oxidase in *Arabidopsis* [37]. Analysis of antioxidants also clearly demonstrates a mediating role for AOS during ozone stress.

Expression of the cytosolic *APX1* gene of *Arabidopsis* increases after low-level exposure to ozone [205]. Similarly, cytosolic APX, Cu/ZnSOD, and GST mRNA increase with the onset of visible injury in tobacco, *Arabidopsis*, and bean [90, 206–208]. In contrast, FeSOD_{chl} and GR decrease in tobacco and *Arabidopsis* during ozone treatment [90, 208]. Additional evidence for an involvement of AOS was provided by the overlap in AOS-dependent signaling induced by ozone exposure and pathogen infection [209–214].

Ozone is not the only air pollutant linked to AOS formation and damage. Once it enters the leaf, sulfur dioxide (SO_2) is hydrated to form HSO_3^- and $SO_3^{\cdot-}$ which can affect carbon metabolism [215]. The presence of sulfite and bisulfite leads to a free radical chain reaction initiated by the interaction of sulfite with $O_2^{\cdot-}$ from the Mehler reaction. Exposure to SO_2 may also enhance the level of AOS in plant tissues. In tobacco exposed to SO_2 , Cat2 and GPX transcript levels rise whereas other antioxidants are either repressed or unaltered by the treatment [90]. Increased SOD activity correlates with SO_2 tolerance in bean [215, 216] and poplar [217]. In *Arabidopsis*, APX and guaiacol peroxidase activities are enhanced [205], whereas in *Cassia siamea*, SOD and guaiacol peroxidase activities increase [218].

Mechanical and physical stress

Several pathways are involved in physical and mechanical wounding responses. One of these clearly involves AOS accumulation. The source of AOS during such conditions is believed to be NAD(P)H dependent, because wound-induced AOS accumulation in various species is inhibited by diphenyleneiodonium, an NADPH-oxidase inhibitor [219]. Wounding stimulates H_2O_2 generation systemically in tomato leaves [219]. AOS accumulate after slicing tissues of various species [220], mechanical wounding of mesocarp tissue of winter squash [221] or potato tissues [19, 181], and mechanical stress in red macroalgae [222].

Pathogen attack

Several excellent reviews on the role of AOS during plant-pathogen interactions have been published recently [63, 223–225]. A very brief overview is given of the most relevant findings that implicate AOS.

Generation of AOS in plants can be detected within minutes of elicitor or microbial treatment [14, 226–228]. The production of AOS occurs in two distinct phases: an initial, non-specific phase follows within minutes of pathogen addition, and a secondary phase, dependent on recognition of incompatible pathogens by the host begins 1–3 h after the initial burst [1, 5, 228–230]. The exact role of these two AOS bursts is still unclear, but potentiation effects of SA, defense gene activation, and programmed cell death probably require different H_2O_2 kinetics for induction. The initial burst probably originates from an NADPH-dependent oxidase, although as noted above, other candidate sources exist. The produced $O_2^{\cdot-}$ is dismutated by SOD to H_2O_2 and will activate benzoic acid 2-hydroxylase [231] leading to SA accumulation. This rise in SA, in combination with H_2O_2 , is necessary for potentiating defense gene activation and plant cell death [232]. The second, longer-lasting burst is involved in activating defense responses, albeit thought to be NAD(P)H dependent. Its role is supported by results obtained through direct injection of either H_2O_2 or pathogen elicitors. The artificial oxidative burst obtained induces the expression of PR genes [110, 233], phytoalexins [226, 234, 235], cross-linking of the hydroxyproline-rich cell wall glycoproteins [11, 236], and benzoic acid and SA accumulation [115, 231, 237].

A large body of evidence substantiates the damaging effects of AOS during biotic and abiotic stress. Many stress-related symptoms are exacerbated, if not directly dependent on AOS. However, evidence also suggests that AOS may play another role during stress responses, related to their capacity to serve as secondary messengers.

AOS: friends or foes?

Although plant signaling responses and possible cross-talk between different signaling cascades are increasingly well-understood, the mode of action of H_2O_2 remains unknown [63]. It may function as a secondary messenger at low concentrations but still be phytotoxic at higher concentrations [238]. Early evidence in the field of pathogenesis clearly supported a signaling role for AOS during plant-pathogen interactions. In recent years, specific induction of defense responses has been obtained with H_2O_2 treatments and corroborates the idea that H_2O_2 directly regulates defense responses.

The specific induction of GPX and GST by 2 mM H_2O_2 in cell suspension cultures of soybean and the cross-membrane trafficking of H_2O_2 were the first demonstration of H_2O_2 -inducible gene expression in plants [5]. Use of transgenic technology has helped to characterize the H_2O_2 signaling role. Transgenic potato overexpressing a fungal glucose oxidase gene targeted into the apoplast contains constitutively elevated H_2O_2 levels, accumulates SA, and increases production of acidic chitinase and anionic peroxidase transcript levels [8, 239]. Similarly, tobacco deficient in catalase (Cat1AS), which accumulates H_2O_2 under high light, was used to investigate the role of H_2O_2 during systemic and locally acquired resistance. Acidic PR proteins accumulate in both local and distal leaves protected from high light-imposed H_2O_2 stress, clearly demonstrating that enhanced H_2O_2 levels in the exposed part can activate the expression of defense proteins [9]. In both the potato and tobacco system, disease resistance was enhanced.

Although the field of pathogenesis led the way in oxidative stress signaling in plants for many years, several recent studies have implicated H_2O_2 in a more general signaling role. Transcript levels of cytosolic APX are significantly increased by H_2O_2 or paraquat treatment in cell suspension cultures of rice [240]. Addition of diethyldithiocarbamate (a SOD inhibitor resulting in lower H_2O_2) reduces the induction of APX, whereas inhibition of catalase or APX (resulting in H_2O_2 accumulation) increases APX mRNA levels [240]. These results suggest that H_2O_2 is part of the signaling cascade leading to cytosolic APX induction. This study was the first thorough analysis of APX induction by H_2O_2 , although the regulation of APX by paraquat [78, 80, 241] or H_2O_2 [241, 242] had already been reported.

Additional evidence for a signaling role for AOS was also provided by recent findings in the field of plant stress acclimation. Nodal potato explants subcultured from H_2O_2 -treated microplants are resistant to a 15-h heat shock at 42 °C [22, 24]. Small heat shock proteins (HSPs), including mitochondrial HSP22, accumulate in cell suspension cultures of tomato after addition of 2 mM H_2O_2 , but not in response to $O_2^{\cdot-}$ -generating agents [243]. Increased levels of H_2O_2 in Cat1AS plants exposed to high light lead to HSP17.6 accumulation [9]. Injection of H_2O_2 in *Arabidopsis* leaves induces protection from high light-induced photobleaching [21]. Maize coleoptiles treated with H_2O_2 are protected from subsequent chilling stress [20, 106]. Treatment of crowns of winter wheat with various levels of H_2O_2 and a catalase inhibitor lead to concentration-dependent synthesis of various polypeptides, similar to those found when the plants are exposed to low temperature [244]. Additionally, chilling or H_2O_2 treatment of tobacco induces a similar transient burst of calcium [245, 246]. AOS, and more particularly H_2O_2 , can clearly function as a signaling molecule during abiotic stress responses.

For some time, signaling properties of AOS were considered exclusive to H_2O_2 . However, increasing evidence points to differential signaling amongst AOS. Tomato cells treated with digitonin or xanthine oxidase are able to trigger the accumulation of a set of extensin transcripts [247]. One of these is specifically induced by $O_2^{\cdot-}$ and not H_2O_2 or H_2O_2 -generating compounds, such as glucose oxidase. Similarly, phytoalexin accumulation in cell suspension cultures of parsley and lesion formation in lesion-stimulated disease resistance mutants of *Arabidopsis* are induced by $O_2^{\cdot-}$ rather than H_2O_2 [4, 141]. A cDNA sequence with high homology to 1-aminocyclopropane-1-carboxylate synthase (WSACS2) is preferentially expressed after $O_2^{\cdot-}$ treatment [221].

In addition to the direct evidence of AOS signaling during plant stress adaptation, tolerance to a specific environmental stress can sometimes improve the ability to withstand one or more other stresses. Cross-tolerance is often attributed to the fact that various stresses produce a similar effect at the cellular level, including oxidative stress [68]. Ozone or UV exposure of *A. thaliana* and *N. tabacum* induces resistance against virulent *Pseudomonas syringae* strains and tobacco mosaic virus, respectively [210, 248]. Pre-exposure of mung bean (*Vigna radiata* L.) and pea (*P. sativum* L.) to ethylene makes them more resistant against further exposure to ozone [249], and cotton plants exposed to water deficit are more resistant than irrigated plants against paraquat treatment [250]. Finally, drought treatment induces chilling acclimation in various plant species [251, 252]. The capacity for H_2O_2 -induced protection and reports of cross-tolerance suggest that AOS play a common mediator role under these conditions. AOS, and more particularly H_2O_2 and $O_2^{\cdot-}$, may therefore be involved in a general stress resistance signal transduction pathway. Although the stress conditions described above involve the accumulation of individual AOS, AOS levels were not monitored in these studies. The role of AOS in cross-tolerance is therefore still not clearly demonstrated.

Conclusion

This review has summarized the wealth of data on the importance of oxidative stress during many biotic and abiotic stress responses. Besides exacerbating cellular damage, AOS are capable of inducing defense gene expression. H_2O_2 and/or $O_2^{\cdot-}$ can induce different genes, in combination or separately, thereby giving more flexibility to the AOS signaling function. In addition, the evidence emerging from abiotic stress responses clearly indicates that AOS may also play a more general signaling role during plant stress adaptation.

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