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# Functions of the respiratory burst oxidase in biotic interactions, abiotic stress and development

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The production of reactive oxygen intermediates (ROI) is among the earliest temporal events following pathogen recognition in plants. Initially, ROI were thought to be cell-death executioners. Emerging evidence, however, suggests a broader role for ROI as signals that mediate responses to infection, the abiotic environment, developmental cues, and programmed cell death in different cell types. The *Respiratory burst oxidase homolog (Rboh)* gene family encodes the key enzymatic subunit of the plant NADPH oxidase. Rboh proteins are the source of ROI produced following pathogen recognition and in a variety of other processes.

## Addresses

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## Introduction

The bi-phasic production of apoplastic reactive oxygen intermediates (ROI), the so-called 'oxidative burst', is a hallmark of successful recognition of plant pathogens [1,2,3\*]. Since Doke [4] first reported the generation of superoxide anions ( $O_2^-$ ) during incompatible (plant resistant) interactions between potato and the late blight pathogen *Phytophthora infestans*,  $O_2^-$  and/or hydrogen peroxide ( $H_2O_2$ ), its dismutation product, have repeatedly been detected in successful resistance responses [2,5]. Yet, the diverse functions of ROI, and the mechanism that regulates the initial generation of superoxide were elusive until recently.

ROI can be produced inside the plant cell, in chloroplasts, mitochondria and peroxisomes, as byproducts of metabolic processes such as photosynthesis and respiration [6\*\*]. Several biochemical mechanisms have been proposed to explain ROI production. Apoplastic  $H_2O_2$  pro-

duction can be mediated by cell-wall peroxidases, germin-like oxalate oxidases or amino oxidases [6\*\*,7\*\*]. However, key lines of evidence implicated an NADPH oxidase, analogous to that which generates superoxide during the respiratory burst in mammalian phagocytes, as the source of the ROI detected in plants upon successful pathogen recognition. First,  $O_2^-$  is the primary radical produced in most systems [4,5,8]; second, the rate of ROI production per elicited cell is similar in mammals and plants [9]; and third, ROI generation is inhibited by diphenylene iodonium, a well-characterized inhibitor of the mammalian NADPH oxidase (and flavin-containing enzymes) [8,10].

The mammalian NADPH oxidase, also known as the respiratory burst oxidase (RBO), is a protein complex that is comprised of a membrane-bound NADPH-binding flavocytochrome  $b_{558}$  and cytosolic regulatory proteins. In phagocytes, these regulatory proteins include the  $p47^{phox}$ ,  $p67^{phox}$  and  $p40^{phox}$  phosphoproteins and Rac2, a small GTP-binding protein [11,12]. The membrane cytochrome consists of the glycosylated transmembrane protein  $gp91^{phox}$  and the non-glycosylated  $p22^{phox}$  subunit.  $gp91^{phox}$  contains the entire electron transport chain from NADPH to molecular oxygen to produce superoxide outside the plasma membrane. The NADPH oxidase exists in vesicles that are activated via Rac dissociation, phosphorylation of  $p47^{phox}$  and  $p67^{phox}$  and their subsequent recruitment to the membrane-bound complex [11,12]. In humans, chronic granulomatous disease (CGD) results from mutations that render the NADPH oxidase in phagocytes non-functional, and CGD patients suffer recurrent and prolonged microbial infections [13]. Mutations in  $gp91^{phox}$  account for all cases of X-linked CGD [13]. It was initially thought that direct toxicity of the ROI that are generated by the oxidative burst was responsible for the microbial killing activity of mammalian phagocytes. Surprisingly, superoxide that is generated via NADPH oxidase activity plays a signaling role in microbial killing, leading to  $K^+$  flux that activates specific proteases [14,15]. Additional NADPH oxidases, different from those in phagocytes, have been identified in mammals [11,12]. These exhibit tissue-specific expression and mediate functions from cell proliferation to thyroid hormone biosynthesis. Thus, multiple isoforms of  $gp91^{phox}$  act in NADPH oxidases from different cell types, and perform different functions in mammalian cells.

## The plant *Rboh*-encoded NADPH oxidase

The first plant NADPH oxidase gene to be identified was the rice gene *OsrbohA*, which is related to mammalian

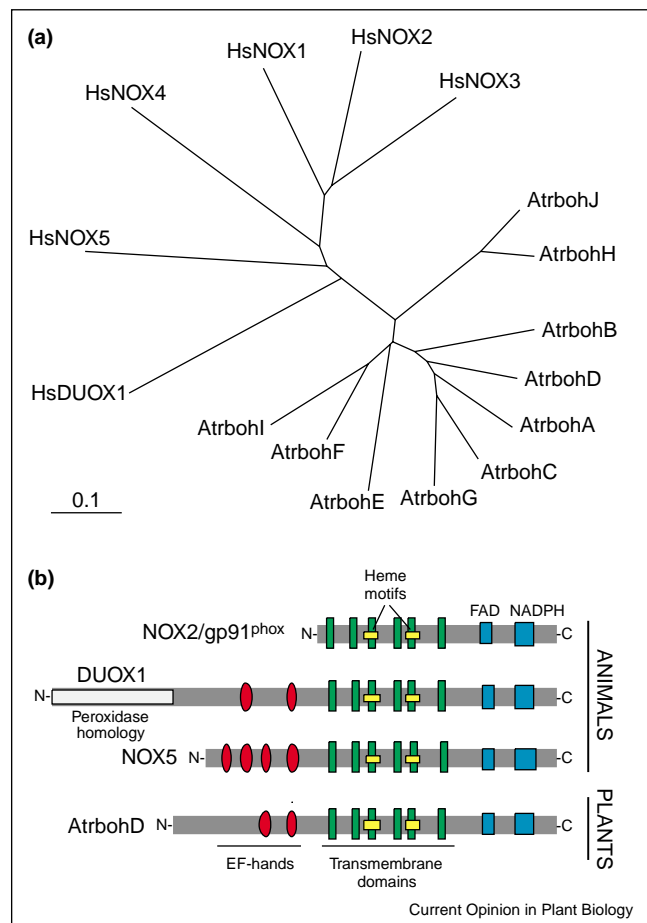
*gp91<sup>phox</sup>* [16]. Subsequent studies documented different *Respiratory burst oxidase homolog (Rboh)* genes in other plant species, including *Arabidopsis*, tomato, tobacco and potato [17–19,20\*,21,22]. Cell fractionation indicated that at least two Rboh proteins are located in the plasma membrane [18,22]. As in the mammalian NADPH oxidase, the rice homologs of human Rac2 can regulate ROI production and HR [23]. But simple comparisons of the regulation and assembly of NADPH oxidases between plants and animals end there. For example, no homologs of the mammalian p47<sup>phox</sup> or p67<sup>phox</sup> were found in the *Arabidopsis* genome [24]; and initial reports [25] that antibodies against the mammalian p47<sup>phox</sup> and p67<sup>phox</sup> proteins cross-reacted with plant proteins of similar size have been dismissed [26].

Plant species contain *Rboh* gene families that have homology to *gp91<sup>phox</sup>*. *Arabidopsis* encodes ten *Atrboh* genes ([17,24]; Figure 1a). All plant *Rboh* genes carry a presumably cytosolic 300-amino-acid amino-terminal extension with two EF-hands that bind Ca<sup>2+</sup> [18], which could account for the direct regulation of these oxidases by Ca<sup>2+</sup>. This is relevant in several functional contexts in which Ca<sup>2+</sup> signaling precedes ROI production. This domain is absent from the mammalian phagocyte *gp91<sup>phox</sup>* but is present in other mammalian NADPH oxidase homologs (Figure 1b). For example, the mammalian NADPH oxidase 5 (NOX5) protein contains four EF-hands in its amino-terminal region and exhibits Ca<sup>2+</sup>-dependent production of superoxide [27]. In contrast to the mammalian *gp91<sup>phox</sup>*, plant Rboh proteins can produce O<sub>2</sub><sup>-</sup> in the absence of additional cytosolic components, and are stimulated directly by Ca<sup>2+</sup> [28]. Thus, a superoxide-generating NADPH oxidase exists in plants, although its precise subunit structure and the regulation of its activation are different than in mammalian phagocytes.

### The Rboh-NADPH oxidase and biotic interactions

ROI were initially proposed to orchestrate the establishment of the plant defense response and the hypersensitive response (HR) cell death that often accompanies successful pathogen recognition [29,30]. Several groups have reported that *Rboh* genes are transcriptionally upregulated by pathogens or fungal elicitors [21,22,31]. Genetic proof of the function of *Rboh* in the pathogen-induced oxidative burst came from analysis of *Rboh* mutants and antisense lines. Infection of *Arabidopsis* mutants that lack functional *AtrbohD* and *AtrbohF* demonstrated that *AtrbohD* is responsible for nearly all of the ROI produced in response to avirulent bacterial or oomycete pathogens, whereas *AtrbohF* is important in the regulation of HR [32]. Antisense reduction of *NtrbohD* indicated that the Rboh encoded by this gene is responsible for the production of ROI after treatment of tobacco cells with the fungal elicitor cryptogein [22]. Silencing of

Figure 1



Relationship and domain structure of the NADPH oxidases.

(a) Phylogenetic tree comparing the mammalian NADPH oxidases with the *Arabidopsis* *Atrboh* proteins: *AtrbohA* (At5g07390), *AtrbohB* (At1g09090), *AtrbohC* (At5g51060), *AtrbohD* (At5g47910), *AtrbohE* (At1g19230), *AtrbohF* (At1g64060), *AtrbohG* (At4g25090), *AtrbohH* (At5g60010), *AtrbohI* (At4g11230), *AtrbohJ* (At3g45810). The *rbohA* described in [18] is identical to *AtrbohF* in [17,32] and is represented here as *AtrbohF*. Only the carboxy-terminal with homology to *gp91<sup>phox</sup>*/NOX2 (excluding the EF hands) was used in the alignment. (b) Schematic representation of the NADPH oxidase proteins with their functional domains. Mammalian NOX1, NOX3 and NOX4 are similar to *gp91<sup>phox</sup>*/NOX2. The plant Rboh are similar to *AtrbohD*. FAD, flavin adenine dinucleotide, DUOX1, Dual oxidase1.

*NtrbohA* and *NtrbohB* in *Nicotiana benthamiana* plants led to less ROI production and reduced resistance to infection by *Phytophthora infestans* [20\*]. There is also evidence of functional overlap between different Rboh proteins. For example, in *Arabidopsis*, the phenotypes of the individual *atrbohD* and *atrbohF* mutants are accentuated in the double mutant *atrbohD|atrbohF* [32,33\*\*]. These data suggest a complex signaling network with interactions between different members of the *Rboh* family.

Interestingly, the downregulation or elimination of *Rboh* leads to variable effects on pathogen growth and HR. For

example, *Nbrboh*-silenced plants are more susceptible to normally avirulent *P. infestans* and HR is suppressed in these plants [20<sup>•</sup>]. By contrast, the *Arabidopsis atrbohF* mutant is more resistant to a weakly virulent strain of the oomycete *Peronospora parasitica*, and actually expresses enhanced HR [32]. Thus, although the Rboh proteins are required for pathogen-induced ROI production, these ROI might serve different signaling functions in disease resistance and HR.

Interactions with other plant defense regulators might account for the divergent outcomes of NADPH-dependent ROI signaling. Exogenously applied ROI can act synergistically in a signal-amplifying loop with salicylic acid (SA) to drive HR [34,35]. However, our recent studies indicate that ROI that are produced via *AtrbohD* actually antagonize SA to stop the spread of cell death beyond the site of HR (MA Torres, JDG Jones, JL Dangl, unpublished results). Also, coordinated levels of ROI and nitric oxide (NO) are required to produce HR [36], and both can mediate abscisic acid (ABA)-induced stomatal closure [37<sup>•</sup>]. These surprising results underscore how ROI that are produced via *Atrboh* proteins can mediate different functions in different contexts, and suggest that we have some way to go to explain the function of ROI in response to infection.

Characterization of plant Rac2 homologs (called Rop for Rho-like proteins [38<sup>•</sup>]) also tied plant NADPH oxidases to ROI production during defense signaling and cell death [23,39<sup>•</sup>]. These analyses also suggested that the combination of specific Rac isoforms with specific Rboh isoforms might have differential regulatory effects. For example, *Osrac1* is a positive regulator of ROI production and cell death [40], whereas *Ntrac5* acts as a negative regulator of *NtrbohD*-dependent ROI production after elicitation with cryptogein [41<sup>•</sup>]. Rac/Rop signaling is also involved in oxygen deprivation, potentially via an NADPH oxidase, and RopGAP4 acts as a rheostat to control ROI production in this context [42]. Comparisons among these datasets are complicated by the use of different plant species to study combinations of *Rac* and *Rboh* family members. These disparate datasets highlight the necessity of careful analysis of the functions of all *Rboh* family members in a single model system and in a variety of developmental and environmental contexts.

### The Rboh-NADPH oxidase in abiotic stress and development

ROI that are generated by plant Rboh-NADPH oxidases also regulate abiotic stress responses and development. ROI function as intermediates in ABA signaling during stomatal closure [43]. H<sub>2</sub>O<sub>2</sub> induces the activation of Ca<sup>2+</sup> channels to mediate the increase in cytosolic Ca<sup>2+</sup> concentration in intact guard cells [44]. *AtrbohD* and *AtrbohF* are expressed in guard cells and are tran-

scriptionally induced in response to ABA. The *Arabidopsis atrbohD/atrbohF* double mutant is impaired in ABA activation of Ca<sup>2+</sup> channels and stomatal closure, whereas exogenous application of H<sub>2</sub>O<sub>2</sub> restored Ca<sup>2+</sup> channel activation and partial stomatal closure in *atrbohD/atrbohF* [33<sup>••</sup>]. This study provided genetic evidence that *Atrboh* proteins mediate ABA signaling in stomata, and suggested that the same *Atrboh* isoforms might mediate different ROI-dependent functions in different cellular contexts.

In this light, it is important to note that a third family member, *AtrbohC* regulates cell expansion during root-hair formation. Analysis of the *atrbohC* mutant (also called *root hair defective 2* [*rhd2*]) revealed that ROI that are produced by *AtrbohC* activate hyper-polarization Ca<sup>2+</sup> channels that are responsible for localized cell expansion during the root-hair formation [45<sup>••</sup>]. In this case, the very reactive hydroxyl radical (OH<sup>-</sup>), which is generated from H<sub>2</sub>O<sub>2</sub> in the presence of transition metals (such as Cu<sup>2+</sup> or Fe<sup>2+</sup>), is the specific ROI implicated.

Thus, ROI that are generated via the NADPH oxidase and the activation of Ca<sup>2+</sup> channels might represent a signaling link that is common to many plant responses. In the defense response, the oxidative burst has also been implicated in activating Ca<sup>2+</sup> influx [46]. Yet, Ca<sup>2+</sup> is also required for ROI production both after pathogen infection and following ABA treatment [47,48,49<sup>•</sup>], and Ca<sup>2+</sup> can activate an Rboh protein *in vitro* [28]. Therefore, Ca<sup>2+</sup> fluxes appear to function both upstream and downstream of ROI production, indicating a complex spatio-temporal Ca<sup>2+</sup> regulation of these signaling networks [46]. Fungal elicitors can induce both ROI and, through the activation of Ca<sup>2+</sup> channels in guard cells, stomatal closure [50]. Thus, the NADPH-oxidase-dependent activation of Ca<sup>2+</sup> channels might represent a cross-talk point between ABA and defense signaling.

Phosphorylation might also play important roles both upstream and downstream of Rboh function (see [6<sup>••</sup>] for review). Yoshioka *et al.* [20<sup>•</sup>] reported that MAPK-induced cell death might be mediated by ROI produced by *NbrbohB*, and that the same MAPK cascade acts to increase *NbrbohB* levels. Particularly relevant in this regard is the identification of Oxidative signal inducible 1 (OXI1)/AGC2-1 (AGC2-1 for protein related to cAMP-dependent and Cgmp-dependent protein kinases, and protein kinase C 2-1), a kinase involved in phospholipid signaling that is upregulated by stimuli that produce ROI [51<sup>••</sup>,52<sup>••</sup>]. Interestingly, the *oxi1* knockout displays alterations in root-hair formation, suggesting that the *OXI1* gene is linked to *AtrbohC* function.

ROI have been implicated in the regulation of cellular responses to stresses other than pathogen infection. For example, H<sub>2</sub>O<sub>2</sub> acts as a second messenger for the induc-

tion of defense genes in response to systemin and jasmonate during wound responses [53]. Using anti-sense transgenics, Sagi *et al.* [54<sup>\*</sup>] showed that ROI that are produced by an Rboh are required for the expression of certain wound-response genes. Reduced expression of *Rboh* additionally induced multiple pleiotropic developmental effects, resulting in the ectopic expression of flower-specific homeotic genes and an altered redox-related metabolism. These findings suggest that ROI that are produced by Rboh influence metabolic balances acting in several hormone signaling pathways. It is difficult to know, however, how many different isoforms of Rboh were affected in these anti-sense transgenics.

The response to ozone (O<sub>3</sub>) also induces an oxidative burst that has similarities to the pathogen-induced oxidative burst [55<sup>\*</sup>]. In *Arabidopsis*, ozone exposure drives both the spatial and the temporal progression of ROI signaling, commencing with the elevation of ROI levels in guard-cell chloroplasts and membranes, and spreading to neighboring cells [56<sup>\*\*</sup>]. *AtrbohD* and *AtrbohF* have been implicated in the intercellular signaling and ultimate cell death that arises from O<sub>3</sub> exposure [56<sup>\*\*</sup>]. Interestingly, heterotrimeric G proteins control this ROI production and sensitivity to O<sub>3</sub>. Heterotrimeric G protein was also implicated in the regulation of ROI production through activation of extracellular calmodulin during stomatal closure [57<sup>\*\*</sup>]. Also, heterotrimeric G protein functions upstream of *Osrac1* in the regulation of the defense response and cell death in rice [58]. Together, these studies suggest that heterotrimeric G proteins might be a common intermediate for ROI signaling in responses to different stimuli.

For ROI to exert a signaling function, the production and removal of ROI have to be tightly controlled and compartmentalized. In addition to the enzymatic production of ROI, chloroplasts (and also mitochondria and peroxisomes) can generate ROI during metabolic processes [6<sup>\*\*</sup>]. Plant cells possess a battery of scavenging systems, including ascorbate peroxidases, glutathione, superoxide dismutases and catalases (see the excellent review by Mittler *et al.* [7<sup>\*\*</sup>]), that maintain ROI homeostasis. The downregulation of scavenging systems in response to stress can lead to an oxidative burst and cell death that is phenotypically similar to HR [59<sup>\*</sup>]. Davletova *et al.* [60<sup>\*\*</sup>] showed that cytosolic ascorbate peroxidase APX1 provides cross-compartment protection to the chloroplast during light stress, and that *AtrbohD* might be required for an ROI-dependent signal that maintains high APX1 levels during light stress. Trienoic fatty acids from the chloroplast might mediate this activation [61<sup>\*</sup>]. In tobacco, *NtrbohD* was upregulated in catalase-deficient plants in response to elevated levels of H<sub>2</sub>O<sub>2</sub> that were produced under high light [59<sup>\*</sup>], suggesting that ROI produced by the Rboh-NADPH oxidase also modulate the response to light stress.

## Conclusions and future directions

Like mammals, plants have an *Rboh* gene family, and recent data suggest that specific *Rboh* genes function in different cellular contexts. The identification of the *Arabidopsis* RbohD-NADPH oxidase as the enzyme responsible for the production of apoplastic ROI in response to pathogens answered a question first posed 25 years ago [1,4]. The *Arabidopsis* Rboh protein family has a diversified functional portfolio. *AtrbohC* appears to have a very specific function in root-hair development, whereas *AtrbohD* and *AtrbohF* display pleiotropic functions following pathogen recognition and during ABA signaling. The accentuated phenotypes of the double mutant, compared to that of either single mutant, indicates functional overlap between *AtrbohD* and *AtrbohF* function, suggesting that interactions between different family members might help fine tune both ROI production and the response to that production. Further characterization of the other Rboh family members and the identification of knockout/knockdown lines for these genes should define additional functions for plant NADPH oxidases and the ROI that they produce.

Analyses of Rboh function suggest that ROI act in complex signaling networks that operate in responses to developmental cues or to the environment. Particularly relevant are the links between Rboh-dependent ROI production and Ca<sup>2+</sup>, and the regulation of Rboh function by Rac/Rop (in common with mammalian gp91<sup>phox</sup>) by heterotrimeric G proteins or by phosphorylation. Also, SA and NO are important defense-response regulators that interact with ROI signaling in response to pathogens. Thus, ROI that are produced by the Rboh-containing NADPH oxidase are part of many signaling pathways and provide a crucial link in the cross-talk to defense responses.

Plants have to cope with excess ROI produced during metabolism, and have evolved sophisticated ways to use the reactive properties of these ROI to modulate cellular signals by tightly regulating ROI production and compartmentalization. We are just beginning to dissect the functions of plant Rboh proteins and to understand the varied functions of the ROI they ultimately produce.

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