Review Article



Ferroptosis in plants: regulation of lipid peroxidation and redox status

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Regulated cell death (RCD) is an essential process that plays key roles along the plant life cycle. Unlike accidental cell death, which is an uncontrolled biological process, RCD involves integrated signaling cascades and precise molecular-mediated mechanisms that are triggered in response to specific exogenous or endogenous stimuli. Ferroptosis is a cell death pathway characterized by the iron-dependent accumulation of lipid reactive oxygen species. Although first described in animals, ferroptosis in plants shares all the main core mechanisms observed for ferroptosis in other systems. In plants as in animals, oxidant and antioxidant systems outline the process of lipid peroxidation during ferroptosis. In plants, cellular compartments such as mitochondria, chloroplasts and cytosol act cooperatively and coordinately to respond to changing redox environments. This particular context makes plants a unique model to study redox status regulation and cell death. In this review, we focus on our most recent understanding of the regulation of redox state and lipid peroxidation in plants and their role during ferroptosis.

Ferroptosis: an oxidative, iron dependent type of cell death

Regulated cell death (RCD) is a ubiquitous process in living organisms that implicates integrated signaling cascades and precise molecular-mediated mechanisms. Unlike accidental cell death (ACD), which is an uncontrolled biological process, RCD is actively mediated through complex signaling pathways that are triggered in response to a specific exogenous or endogenous stimulus. RCD may occur in multiple forms in response to different stresses, physiological processes or developmental clues. Although for several years the prevailing assumption was that apoptosis was the only form of regulated cell death, other cell death programs have emerged, such as necroptosis [1], pyroptosis [2], parthanatos [3], entotic [4] and NETotic cell death [5], autophagy-mediated cell death [6] and ferroptosis [7] among others [8].

In particular, ferroptosis is an oxidative, iron-dependent type of RCD characterized by lipid peroxidation and altered plasma membrane permeability. The term ferroptosis was first used in 2012 by Dixon et al. [7], who described a new cell death pathway triggered by erastin in tumor cells that was distinct from apoptosis and necrosis. Erastin-induced cell death was characterized by mitochondrial atrophy and reduced mitochondrial cristae, while cells do not produce apoptotic bodies and nuclei seemed unaffected [7]. Since its discovery, the ferroptotic pathway has been found in several species, including humans, other vertebrates [9], plants [10, 11] and cyanobacteria [12].

The ferroptosis pathway requires the availability of redox-active iron and is characterized by impairment of the cellular antioxidant machinery and the accumulation of toxic lipid hydroperoxides. Lipid hydrogen peroxides can be produced by the action of lipoxygenases or by oxygen free radical mediation, in which the resultant lipid hydroperoxides generate alkoxy radicals under the catalysis of Fe^{2+} ,

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initiating a chain reaction. The accumulation of lipid hydroperoxides and the depletion of polyunsaturated fatty acids containing phospholipids (PUFA-PLs) change the fluidity and permeability of the plasma membrane, eventually leading to cell death [13, 14].

As demonstrated in mice, tumor cells and neuronal cells, Glutathione peroxidase 4 (GPX4) is one of the key regulators of ferroptosis [15–18]. This role of GPX4 is based on its ability to reduce complex hydroperoxides, interrupting the lipid peroxidation chain reaction. GPX4 converts glutathione (GSH) into oxidized glutathione (GSSG), reducing the cytotoxic lipid hydroperoxides (L-OOH) to the corresponding alcohols (L-OH). Thus, the inhibition of GPX4 activity generally leads to the accumulation of lipid peroxides, which is a marker of ferroptosis. Accordingly, down-regulation of GPX4 expression sensitizes cells to ferroptosis, while up-regulation of GPX4 expression is enough to inhibit this process [18]. In addition, as the activity of GPX4 is dependent on GSH, the source of GSH is an important regulatory condition for GPX4 action.

On the other hand, lipid peroxidation is also caused by an increase in the activity of lipoxygenases (LOXs). LOXs are iron-containing enzymes that catalyze the deoxygenation of lipids that have been proposed to contribute to the buildup of toxic lipid peroxides that kill the cell through ferroptosis [19]. Although current work in animal cells suggests that LOXs activity is not essential for the execution of ferroptosis, LOXs may participate in its initiation by building up the pool of lipid hydroperoxides that promote lipid autoxidation [19]. The subcellular location of lipid peroxidation is still unknown, although the plasma membrane is considered a likely source together with mitochondrial membranes, as their morphology is compromised during ferroptosis.

Acyl-CoA synthetase long chain family member 4 (ACSL4), which is part of the Acyl CoA synthase (ACS) family, converts long-chain fatty acids to acyl-CoA. As ACSL4 plays an essential role in the synthesis of long-chain PUFA-CoA, their overexpression has been linked to the accumulation of long-chain PUFA-CoA and in the accumulation of oxidized phospholipids on the plasma membrane surface, sensitizing cancer cells to ferroptosis [20].

Another source of free radical oxidation is the NADPH oxidase (NOX) oxidation reaction. NOX enzymes consume NADPH producing reactive oxygen species (ROS) during processes related to host defense and signal regulation. However, when deregulation of NOX activity occurs as a result of an overwhelming stress, ROS can accumulate in the cell, which can also promote the sensitivity to ferroptosis, as observed in tumor cells [7].

In plants, ferroptosis has been first shown to be triggered by heat stress, displaying some of the characteristics described in mammalian cells: iron-dependent ROS accumulation, lipid peroxidation and GSH depletion, as well as many morphological hallmarks: mitochondria shrinkage, cytoplasm retraction, formation of small vacuoles and normal nuclei. It was also shown that ferroptosis can be prevented by using canonical ferroptosis inhibitors such as Ferrostatin-1 (Fer-1, a lipophilic antioxidant), and Ciclopiroxolamine (CPX, an intracellular iron chelator, Figure 1) [11].

Since then, additional studies in plants have shown the importance of ferroptosis as a mechanism of RCD in other processes. *Nicotiana benthamiana* plants infected with a highly infectious tobacco mosaic virus (24A + UPD strain) showed accelerated death related to high intracellular iron and ROS concentrations. Plants were protected from 24A + UPD induced death when intracellular iron concentration was decreased or after applying ferroptosis inhibitors. Also, silencing *NbGPX* results in higher ferroptotic cell death caused by 24A + UPD infection [21]. Hypersensitive response (HR) cell death is a plant immune response that restricts pathogen invasion. In rice (*Oryza sativa*), the interaction with the fungus *Magnaporthe oryzae* results in a HR that was characterized as an oxidative iron-dependent type of cell death. When this cell death was prevented with ferroptosis triggered by *M. oryzae* infection. *osmek2* knock-out plants do not show iron and ROS accumulation, and overexpression of Osmek2 induces cell death, which can be prevented by ferroptosis inhibitors [22]. Interestingly, the involvement of MAPK cascades has been also recently related to ferroptosis in tumor cells [23, 24] Another study suggests that acrolein, a carbonyl species derived from lipid peroxides, might also have a role in ferroptosis. Accordingly, acrolein induced cell death was prevented by ferroptosis inhibitors (Figure 1) [25].

In addition, ferroptosis has been also described in cyanobacteria: after heat shock, *Synechocystis sp* undergoes a cell death process that can be prevented by canonical ferroptosis inhibitors. As the described pathway shares many of the pivotal characteristics of ferroptosis, the process was called c-ferroptosis and it was suggested that this mechanism might be an ancient and conserved cell death program [12].

ROS-mediated lipid peroxidation is the key step that drives ferroptosis in all species described so far. The accumulation of ROS and the consequent oxidative damage results from an imbalance between the generation





Figure 1. Major mechanisms of oxidative damage and antioxidant defense proposed to work in plant ferroptosis. Ferroptosis can be induced by biotic or abiotic stresses. ROS can be produced by three main sources: (i) ROS produced by membrane associated NOX enzymes (ii) ROS produced in mitochondria, and (iii) Fenton reactions. Lipid peroxidation of PUFAs can occur through enzymatic or non-enzymatic processes. MAPKs can phosphorylate WRKY transcription factors which in turn induce NOX expression, leading to ROS accumulation. GPX detoxifies lipid peroxides using thioredoxin (TRX) as a reductant agent. RSL3 inhibits GPX, leading to lipid peroxides accumulation. Lipid ROS can be degraded to reactive carbonyl species (RCS), such as acrolein, which are related to cell death. Acrolein addition induces cell death, which can be prevented by glutathione (GSH). Treatment with CPX, Fer-1, DPI, LIP-1 (liproxstatin-1) and D-PUFA inhibits ferroptosis. Pro-ferroptotic pathways are shown in orange, anti-ferroptotic pathways are shown in blue. Dashed lines indicate indirect evidence.

of free radicals and the antioxidant system ability to neutralize or eliminate their harmful effects. In this review, we will focus on the regulation of redox status in plants and its implications during ferroptosis.

Regulation of redox status and cell death in plants

Redox chemistry is a key feature of life on earth. Like animals, plants are obligate aerobic organisms and they require oxygen for mitochondrial respiration. However, as green tissues continuously produce oxygen through photosynthesis during day time, plant cells are exposed to much higher concentration of oxygen than animal



cells. The incorporation of the photosynthetic electron transport chain resulted in a large production of ROS, such as superoxide, hydrogen peroxide and singlet oxygen. ROS are very reactive and have the potential to modify essential biomolecules like proteins, lipids and DNA. This increase in oxidative species arising from photosynthesis was accompanied by a plethora of mechanisms to monitor redox status and redox regulatory networks that enable plants to sense and respond to changes in redox homeostasis [26, 27]. Accumulation of ROS is tightly regulated by the antioxidative system. This system consists of large pools of antioxidants and specific enzymes like Ascorbate Peroxidase (APX) and Glutathione peroxidase (GPX) that control the lifetime and the specificity of ROS signaling. In addition to GSH, plants synthetize additional antioxidants like ascorbate and tocopherols. The balance of ROS formation and ROS reduction enables cells to survive the inevitable ROS production avoiding excessive damage.

However, ROS can also play a key role during plant life acting as signal molecules. ROS signaling could induce acclimation to different stresses, for example by modulating stomatal closure, root hair growth and hormones responses [28]. However, ROS can also take part of signal transduction pathways that end in RCD processes in response to several environmental and developmental triggers. ROS production occurs in distinct subcellular compartments, such as cell wall, apoplast, chloroplast, mitochondria, and peroxisomes. Cell compartments can differ in their redox status, as they diverge in their pool of antioxidants and sources of ROS (reviewed in [29]). As different types of ROS are produced at different subcellular sites, the outcomes and the integration of such signals are very specific. Redox regulation occurs independently in these subcellular compartments and ROS bursts induced in specific locations will only trigger a signaling route available in such space. Also, as ROS buffering systems also vary, lifetime of the signal might also be completely different [30]. Although many aspects of ROS production and signaling remain to be elucidated, these facts might explain why ROS can regulate distinct types of RCD [31-34]. One example of ROS signaling as part of a developmental triggered RCD constitutes the process of tapetum RCD, which is essential for pollen development. Aberrant tapetum RDC (either premature or delayed) due to a disturbed ROS balance produced by the aberrant up-regulation of Superoxide Dismutase (SOD), Catalase (CAT), APX, and GPX during early pollen development [35] results in sterility. Xie and coworkers [33] reviewed how ROS signaling is regulated at different levels to guaranty tapetum RCD at a specific time during another development. For instance, a complex transcriptional network regulates NOX expression in tomato, rice, arabidopsis and tobacco tapetum cells. Enzymes involved in lipid homeostasis or antioxidant system are also regulated [33].

During biotic stresses, specific types of plant-pathogen interactions elicit the HR, a form of RCD that occurs at the site of pathogen entrance that prevents pathogen spread. Available data show that during the HR displayed in different plant-pathogen systems, ROS production arises after NOXs activation, which has been largely recognized as a crucial event to induce cell death [36, 37]. Additionally, down-regulation of antioxidant enzymes, like CAT or APX, occurs in parallel to ROS increase [38, 39]. Abiotic stresses like drought, heat, light, ozone, among others, also induce ROS-dependant RCD [32]. A nice example of the complex processes that regulate cell death is heat stress, as it has been reported that different temperatures could trigger diverse signaling pathways [22, 39, 40]. Upon heat shock, NOXs are involved in ROS production, which is required to induce heat shock transcription factors (HSFs) and heat shock proteins (HSPs), MAPKs, and vacuolar processing enzymes (VPEs) (reviewed in [32]). Interestingly, Suzuki and Katano [40] reviewed evidence from different laboratories and proposed a model that suggests how ROS signaling could integrate development, biotic stress and heat response through calcium and NOX-dependent ROS signals.

Oxidative damage and the antioxidant system in plant ferroptosis

ROS-mediated lipid peroxidation is a key step that drives ferroptosis. In animal cells, ferroptosis is initiated by ROS accumulation as a result of three main sources/processes: (i) ROS produced by membrane associated NOX enzymes (ii) ROS produced in mitochondria, and (iii) Fenton reactions (discussed in the next section).

In plant ferroptosis, ROS accumulation is one of the earliest recognized biochemical events that take place soon after RCD induction. This cell death pathway, triggered by biotic o abiotic stresses, could be prevented by pretreatment with diphenyleneiodonium (DPI), a NOX inhibitor. Upon heat stress, cytosolic ROS increases shortly after treatment and can be consistently measurable from 15 min to 3 h after HS. Pretreatment with DPI not only prevents cytosolic ROS accumulation, but also cell death [11].



Additionally, Dangol and coworkers [22] proposed that ferroptosis triggered by pathogen infection, which is also accompanied by a massive production of ROS, activates MAPKs that might phosphorylate WRKY transcription factors to subsequently induce NOXs expression (Figure 1).

Mitochondrial ROS production mainly occurs during oxidative phosphorylation in the electron transport chain at the inner membrane. Electrons leakage from complex I and complex III results in a partial reduction in oxygen to form superoxide anion $(O_2^{\bullet-})$, which is rapidly converted into hydrogen peroxide (H_2O_2) . Overall, $O_2^{\bullet-}$ and H_2O_2 produced during this process are called mitochondrial ROS. The accumulation of mitochondrial ROS induces a decrease in the transmembrane potential ($\Delta\Psi$ m), a phenomenon observed both in animal and plant cell death [41, 42]. The involvement of mitochondrial ROS during ferroptosis in human cells seems to be specific depending on the system and cell death triggers [7, 43–45].

Mitochondrial ROS have been largely associated with plant cell death pathways. Cell death progression during the HR can be inhibited by supplying antioxidant enzymes to the mitochondria and perturbation of alternative oxidase (AOX) leads to altered cell death rates [46–48]. However, so far there is not clear evidence indicating a role for mitochondria on ROS production during plant ferroptosis. The use of mitoSOX (a mitochondrial superoxide sensitive fluorescent probe) was not able to detect an increment of fluorescence in Arabidopsis seedlings undergoing ferroptosis after a heat stress [11].

Additionally, chloroplasts might also contribute to a large increase in ROS upon heat stress, [46–48]. Chloroplasts constitutively express antioxidant defense mechanisms to maintain redox homeostasis that prevent the oxidative damage to biological macromolecules. This antioxidant capacity is essential for heat stress adaptation and the acquisition of thermotolerance. Furthermore, chloroplasts ROS can work as plastid signals that are sensed in the nucleus to activate the expression of genes that allow an efficient adaptation to environmental stresses [49].

It is also possible that chloroplastic ROS contribute to the oxidative burst that follows heat stress in plants undergoing ferroptosis [11]. When the aerial parts of Arabidopsis seedlings were exposed to high temperatures, plants exposed to 43°C were observed to die at higher rates in the light than in the dark, suggesting that active chloroplasts are contributing to cell death in leaves [11]. As this cell death pathway triggered by heat stress was prevented by ferroptosis inhibitors, it was postulated that chloroplasts might be involved in this pathway, although the basis of this contribution is still a matter of study. Also, it has been proposed that mitochondria and chloroplasts might work cooperatively during the induction of plant cell death. This is supported by several reports in which light is required for efficient execution of cell death, that is also dependent on mitochondrial ROS production [50–53]. In addition, plants that carry a mutation in Deficiency in MOSAIC DEATH 1 (MOD1), a plastid-localized enoyl-ACP reductase, accumulate ROS and undergo cell death, which is suppressed by mitochondrial complex I mutations. Remarkably, both chloroplastic DICARBOXYLATE TRANSPORTER 1 (DiT1) and mitochondrial MALATE DEHYDROGENASE 1 (mMDH1) can rescue the ROS accumulation and RCD phenotypes in the *mod1* background, indicating a communication from chloroplasts to mitochondria via the malate shuttle during cell death [54].

Regulation of lipid peroxidation and ferroptosis

As we have mentioned before, lipid-ROS production is a hallmark of ferroptosis, which mainly occurs when PUFAs are oxidized, via enzymatic or non-enzymatic processes (reviewed in [55]). In plants, the inhibition of PUFA peroxidation blocks heat stress-induced ferroptosis [11]. However, the mechanism responsible for such oxidation process, subcellular localization and direct lipid-ROS target are yet to be identified.

It has been reported that ferroptosis in animal cells is dependent on lipid peroxidation mediated by LOX enzymes [17, 56, 57]. Lipidomic studies have determined that PUFA-PEs are more prone to undergo peroxidation in this type of RCD [57]. This might be due to the fact that PEs are capable of develop a non-bilayer arrangement that might facilitate oxidation during ferroptosis [58]. Recently Wenzel et al. [59] described a phosphatidylethanolamine-binding protein 1 (PEBP1), a small scaffolding protein, that seems to be responsible for the substrate specificity observed in LOX lipid peroxidation. PEBP1 can bind to two human LOX isoforms (15LO1 and 15LO2). This protein family is present in all eukaryotes. Particularly in plants, this family can be subdivided in three subfamilies: TERMINAL FLOWER1 (TFL1)-like, FLOWERING LOCUS T (FT)-like and MOTHER OF FT AND FTL1 (MFT)- like [60]. However, there is no evidence of the involvement of these plant PEBPs in cell death.

LOX enzymes in plants are classified as 9-LOX or 13-LOX, depending on the position of the carbon atom at which the oxygen is added [61]. Particularly in Arabidopsis there are 6 isoforms of LOX: LOX 2, 3 and 4 are



9-LOX, while LOX6 is a 13-LOX [62]. Lipid peroxidation mediated by LOX in plants has been proven to be a hallmark in cell death triggered by biotic and abiotic stresses. A lipidomic analysis revealed that cell death triggered during the interaction between Arabidopsis and *Pseudomonas syringae* is preceded by plastid lipid peroxidation of galactolipids and triacylglyceride species [63]. Additionally, the activity of LOX2 increased significantly during the infection, which was crucial for the enzymatic lipid peroxidation induced by the interaction. Silencing of 9-LOXs or 13-LOXs in *Nicotiana benthamiana* was shown to reduce the rate of RCD induced by the infection with the Potato Virus X-Potato Virus Y (PVX-PVY) or with Tomato spotted Wilt Virus (TSWV) [64]. Also, the transient expression a 9-LOX of pepper (CaLOX1) was enough to induce defense responses and cell death [64]. Similarly, overexpression of CaLOX1 in Arabidopsis conferred a greater resistance to a diverse number of infections [65]. On the other hand, silencing of CaLOX1 reduced lipid peroxidation, ROS levels and salicylic acid accumulation, as well as the expression of defense-related genes in response to a bacterial infection [65].

In lentil (*Lens culinaris*), root protoplasts exposed to oxidative stress by H_2O_2 undergo RCD, which is characterized by early membrane lipid peroxidation and an increase in LOX levels and activity. Furthermore, two LOX products (9- and 13-hydroperoxy-octadecadienoic acids and their reduced alcohol derivatives), were able to induce RCD in the lentil root protoplasts [66]. Interestingly, Christensen et al. [67] showed that the activity of 9-LOX on linolenic acid produces 10-oxo-11-phytotenoic acid (10-OPEA) as well as 12- and 14-carbon cyclopente(a)nones which are named 'death acids' (DAs) and regulate the expression of genes related to defense and promote cytotoxicity.

Lipid peroxidation also regulates ferroptosis in plants. Pretreatment of Arabidopsis root cells with deuterated PUFAS (D-PUFAs), which contain deuterium at bis-allylic carbons, inhibits oxidation and prevents heat-stress induced ferroptosis [11]. In addition, it has been recently shown that a reactive lipid peroxide derived named acrolein induces ferroptosis in Arabidopsis cell cultures, which is prevented by treatment with canonical ferroptosis inhibitors and GSH (Figure 1) [25].

Lipid hydroperoxides can also be produced by non-enzymatic lipid peroxidation. Non-enzymatic lipid peroxidation occurs when oxygen radicals react with double bonds in a PUFA, giving a lipid-peroxyl radical. Oxygen radicals (HO•) are produced through Fenton reaction, according to the following equation:

$$Fe^{2+} + H_2O_2 \rightarrow Fe^{3+} + HO \cdot + HO^-$$

HO• reacts then with PUFAs resulting in a lipid radical. These radical lipids are able to: (i) react with other lipids in a chain reaction, (ii) produce ROS or (iii) react with molecules like proteins or DNA, (Figure 1, reviewed in [68]). Supporting a contribution of Fenton chemistry in plant ferroptosis, accumulation of Fe³⁺ and ROS was found associated to cell death sites in rice after the infection with an avirulent strain of *M. oryzae.* This suggests an iron-dependent ROS burst may mediate HR cell death in rice [10]. As the Fenton reaction requires labile iron, iron associated to FeS clusters, heme groups or attached to iron storage proteins does not participate directly in non-enzymatic lipid-ROS production [55]. Regulation of labile iron is essential to preserve homeostasis. Although iron is required for vital metabolic processes, an excess of labile iron is potentially harmful because of its tendency to participate in oxidation–reduction reactions that generate free radicals. Labile iron pools increase during ferroptosis in animal cells [69, 70]. Similarly, abiotic stresses in plants, like high temperatures, can induce the release of 2Fe-2S clusters [71, 72], possibly sensitizing plant cells to ferroptosis as occurs in animal systems. Furthermore, iron can also bind to Poly rC Binding-Proteins (PCBPs), which are capable to transfer iron to lipoxygenases implicated in ferroptosis [73]. Supporting a similar role for these proteins in plants, strong induction of a PCBP protein has been found in *Nicotiana Benthamiana* during ferroptosis induced by a viral infection [21].

Lipid ROS could persist and diffuse through lipid bilayers, or can be degraded to aldehydes (e.g. malondialdehyde (MDA), 4-hydroxynonenal (4HNE), acrolein) or hydroxi-acids, which are also reactive molecules [55]. These species are known as reactive carbonyl species (RCS) and have been largely related to cell death in plants. Hajdinak and coworkers [25] showed that Arabidopsis cells treated with acrolein endured cell death, which can be prevented by cotreatment with ferroptosis inhibitors and GSH. This cell death pathway was also associated to an activation of caspase-like proteins [74], which have been also linked to heat-induced ferroptosis [11] and RSL3- induced cell death [25].



Because RCS are comparatively more stable than ROS, they have been proposed as suitable signals modulating a wide range of physiological aspects of plants (including cell death), not only inside the cell but also between cells (Figure 1) [75]. Interestingly, RCS might contribute to propagate a ferroptotic signal between cells, as has been reported for hydroperoxides in tumor cells [76].

Concluding remarks

Ferroptosis in plants shares all the core molecular mechanisms described for ferroptosis in other systems. As described above, it occurs through antioxidant depletion, ROS accumulation and iron-dependent lipid peroxidation. In this context, redox homeostasis regulation is crucial, as multiple oxidative and antioxidant systems shape the process of lipid peroxidation during ferroptosis. The oxidative damage is limited by the activation of an antioxidant system, which is finally overwhelmed in cells entering the ferroptotic pathway. Although several aspects of plant ferroptosis are similar to the processes described in animal systems, plants show specific characteristics. Active chloroplasts seem relevant for this pathway, as light is a determinant factor in plant cells undergoing ferroptosis. However, the basis of chloroplast contribution is still unknown, as their putative concerted role with mitochondria producing pro-cell death signals and ROS. Additionally, although lipid peroxidation is known to play an essential function in plant ferroptosis, the ultimate executors of cell death are still unknown. Interestingly, lipid ROS derivatives appear as possible new players of this pathway that might act not only within the cell, but also promoting cell death in distant tissues through a cell-non-autonomous mechanism. This is an intriguing and interesting idea that deserves further research attention, as it might provide with new and exciting data to explain how this cell death pathway is regulated and executed.

Competing Interests

The authors declare that there are no competing interests associated with the manuscript.

Author Contributions

Conceptualization: A.M.D., G.A.L., E.Z. and G.C.P.; Funding Acquisition: G.C.P., E.Z. and A.M.D.; Project Administration: G.C.P. and A.M.D.; Supervision: A.M.D. and G.C.P.; Writing: A.M.D., G.A.L., V.B. and G.C.P.; Writing – Review and Editing: A.M.D., G.A.L., V.B. and E.Z.

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Abbreviations

ACSL4, Acyl-CoA synthetase long chain family member 4; APX, ascorbate Peroxidase; CAT, catalase; CPX, ciclopiroxolamine; DPI, diphenyleneiodonium; GPX, glutathione peroxidase; HR, hypersensitive response; LOXs, lipoxygenases; MAPKs, mitogen activated protein kinases; NOX, NADPH oxidase; PEBP1, phosphatidylethanolamine-binding protein 1; RCD, regulated cell death; RCS, reactive carbonyl species; ROS, reactive oxygen species.

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