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EDITED BY

Juan Carlos Mayo,
University of Oviedo, Spain

REVIEWED BY

Toni Petan,
Institut Jožef Stefan (IJS), Slovenia
Pedro Gonzalez-Menendez,
University of Oviedo, Spain

*CORRESPONDENCE

Wan Seok Yang,
✉ yangw@stjohns.edu

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Ferroptosis: whERe is the critical site of lipid peroxidation?

Wan Seok Yang*

Department of Biological Sciences, St. John's University, New York, NY, United States

KEYWORDS

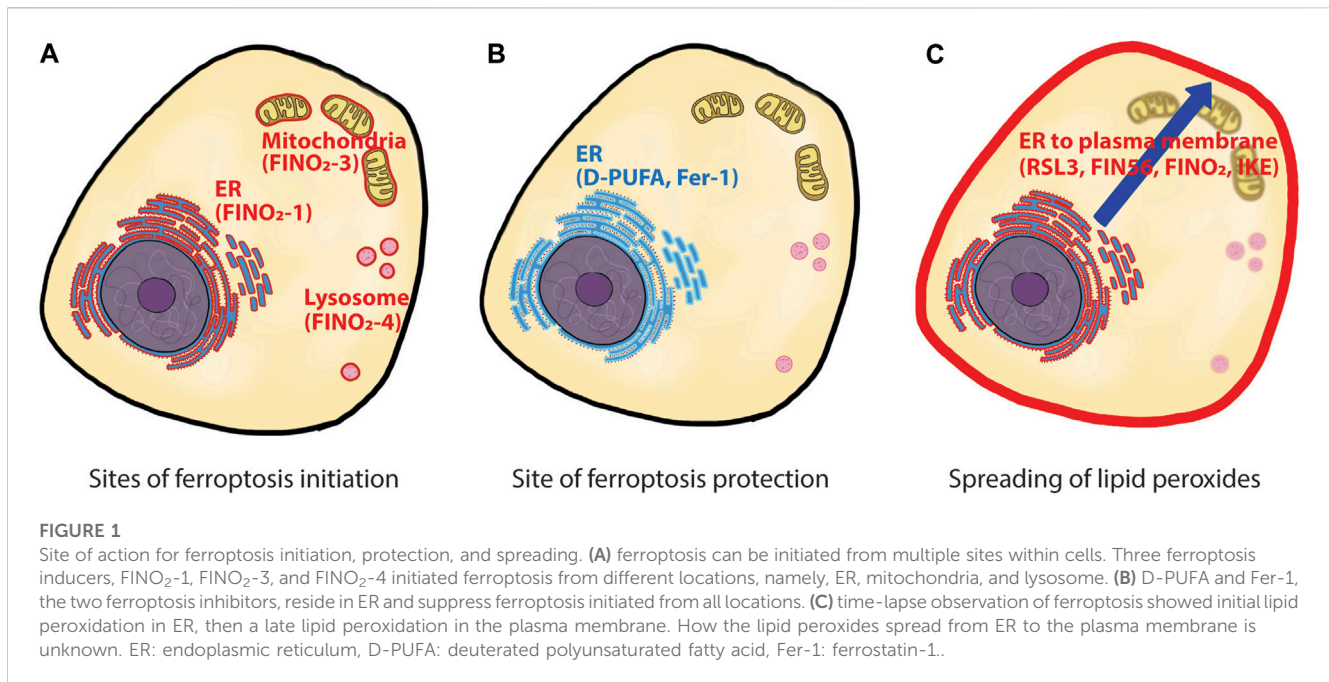
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Ferroptosis is a regulated form of cell death that is driven by a lethal accumulation of lipid peroxides in cell membranes (Dixon et al., 2012). Recent studies have revealed the involvement of ferroptosis in various physiological and pathological conditions (Stockwell, 2022). Understanding the precise mechanism of ferroptosis is necessary to explain this phenomenon in various contexts.

Following the initial observation that lipid peroxides in cell membranes “execute” ferroptotic cell death, researchers have reported a more specific biochemical nature of these peroxides. For example, a genome-wide CRISPR screening identified acyl-CoA synthetase long-chain family member 4 (ACSL4) as an essential gene for executing ferroptosis, revealing that the ACSL4-mediated loading of the PUFAs to the membrane phospholipids is a critical event for executing ferroptosis (Doll et al., 2017). Conversely, unloaded, free PUFAs within cells were found to be neutral in modulating ferroptosis sensitivity. In addition, LC-MS/MS-based lipidomics analysis showed that phosphatidylethanolamines (PEs) containing polyunsaturated fatty acids (PUFAs) were more susceptible to oxidative damage during ferroptosis compared to other phospholipids in cell membranes (Kagan et al., 2017). Later, another genome-wide CRISPR screening discovered ether-phospholipid with PUFAs as a distinct functional lipid class mediating ferroptosis (Zou et al., 2020).

Despite significant progress in identifying and refining lipid classes relevant to ferroptosis, there remained the question of whether specific subcellular locations of lipids that are more susceptible to ferroptosis initiation or play more important roles in ferroptosis execution. A recent study by von Krusenstiern and others (von Krusenstiern et al., 2023) addressed this question using fluorescence and stimulated Raman scattering (SRS) imaging techniques (Figure 1). Compared to fluorescent tags, Raman active tags offer an advantage in monitoring the subcellular location of small molecules, such as drugs or metabolites, due to their relatively tiny size (Shen et al., 2019).

The authors first examined the cellular localization of deuterated-PUFA (D-PUFA), a specific ferroptosis inhibitor (von Krusenstiern et al., 2023; Yang et al., 2016). Since deuterium is a Raman active tag, there was no need to modify D-PUFA further to visualize it within cells. SRS imaging revealed that D-PUFAs were primarily located in perinuclear regions and abundant puncta structures in the cells, and subsequent fluorescence imaging with Nile Red confirmed the puncta structures as lipid droplets. Treatment of cells with diglyceride acyltransferase (DGAT) inhibitors removed these lipid droplets. However, the absence of lipid droplets did not affect D-PUFA's anti-ferroptotic activity, suggesting that lipid droplets are not functionally relevant to ferroptosis. Further analysis using fluorescence microscopy with ER-tracker Green dye revealed that the perinuclear region was the endoplasmic reticulum (ER), indicating that the ER is another cellular location where D-PUFAs mainly reside. Unlike lipid droplets, however, there was no straightforward way to alter the surface area of ER using pharmacological or genetic reagents. For example, induction of ER-phagy (Mochida and Nakatogawa, 2022) was unsuccessful in the cell lines used in this study, despite extensive attempts. Instead, the



authors treated cells with pro-ferroptotic PUFAs or anti-ferroptotic MUFAs and determined their cellular locations using SRS imaging. Both fatty acids were mainly located in the ER, with small percentages located in plasma membrane. The authors then overexpressed ACSL4 in the cells to increase the loading of pro-ferroptotic PUFAs or anti-ferroptotic MUFAs to the ER phospholipids. This approach demonstrated a solid correlation between the ER membrane composition and ferroptosis sensitivity, highlighting the ER as a site of ferroptosis modulation.

Next, the authors explored the site of action for FINO₂, a canonical ferroptosis inducer, which directly oxidizes cellular iron and indirectly inhibits GPx4 to induce ferroptosis (Gaschler et al., 2018a). Structurally, FINO₂ is a lipophilic compound that contains an endoperoxide moiety, allowing it to accumulate in cell membranes and induce lipid peroxidation directly at those locations. Taking advantage of this feature, the authors synthesized a series of FINO₂ analogs with fluorescent tags that were distributed to different cellular locations to investigate whether ferroptosis can be initiated by oxidative damage in a particular cell membrane. The original FINO₂ with a fluorescent tag displayed ER distribution and induced ferroptosis. Interestingly, FINO₂ analogs distributed to mitochondria and lysosomes also induced ferroptosis, indicating that ferroptosis can be initiated from various organelles. More importantly, cells treated with mitochondrial and lysosomal FINO₂ analogs were protected by ferrostatin-1 (Fer-1). A previous report demonstrated that Fer-1 primarily acted at the ER site, rather than in mitochondria and lysosomes which were dispensable to Fer-1's anti-ferroptotic activity (Gaschler et al., 2018b). These data suggest that while ferroptosis can be initiated at multiple sites, the ER is the critical site for ferroptosis protection.

Lastly, the authors examined time-dependent changes in the location of lipid peroxidation using BODIPY™-C11 dye and fluorescence microscopy. Upon treatment with ferroptosis inducers, cells underwent initial lipid peroxidation events in

the ER, followed by lipid peroxidation in the plasma membrane. This time-dependent change was observed in all four classes of ferroptosis inducers (RSL3, FIN56, FINO₂, or IKE). These findings complement a previous study, which reported that both erastin2 and RSL3 caused intracellular lipid peroxidation before the onset of lipid peroxidation in the plasma membrane (Magtanong et al., 2022). Altogether, these data strongly point towards the ER membrane as the most critical site for inducing ferroptosis.

In summary, the findings outlined above emphasized the essential role of the ER in ferroptotic cell death. The conclusion from the current research raises several intriguing questions. First, what makes the ER so critical in executing ferroptosis? Does the ER membrane contain a higher level of ferroptosis-relevant lipids compared to other organelles? Are there any microenvironmental factors in the ER that can affect ferroptosis? A refined lipidomics analysis with subcellular fractionation samples may provide answers to these questions. Second, how is the ferroptosis signal initiated from other organelles transduced to the ER? For example, it is unclear how the lipid peroxides in mitochondria or lysosomes spread to the ER when FINO₂ analogs are used. This question also applies to the later stage of ferroptosis. How are the lipid peroxides in the ER transduced or transported to the plasma membrane before the onset of cell death? Could it be the conventional vesicle-mediated transport or some other mechanism? Third, can we find more specific regions within the ER, such as rough ER, smooth ER, ER lumen, or microdomains in the ER membrane, that are responsible for the initiation of ferroptosis? Lastly, what are the ER-specific genes that regulate ferroptosis? Xiaodong Wang's group recently discovered that the ER-resident oxidoreductases POR and CYB5R1 produced initial hydrogen peroxides, which subsequently induced lipid peroxidation and ferroptosis (Yan et al., 2021). Future research focused on the ER should broaden our knowledge about the natural triggers of

ferroptosis and help identify better target proteins for ferroptosis targeted therapy.

Author contributions

The author confirms being the sole contributor of this work and has approved it for publication.

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