Ferroptosis occurs through an osmotic mechanism and propagates independently of cell rupture

Michelle Riegmans, Liran Sagie, Chen Galed, Tom Levin, Noah Steinberg, Scott J. Dixon, Ulrich Wiesner, Michelle S. Bradbury, Philipp Niethammer, Assaf Zaritsky, and Michael Overholtzer

Ferroptosis is a regulated form of necrotic cell death that is caused by the accumulation of oxidized phospholipids, leading to membrane damage and cell lysis. Although other types of necrotic death such as pyroptosis and necroptosis are mediated by active mechanisms of execution, ferroptosis is thought to result from the accumulation of unrepaired cell damage. Previous studies have suggested that ferroptosis has the ability to spread through cell populations in a wave-like manner, resulting in a distinct spatiotemporal pattern of cell death. Here we investigate the mechanism of ferroptosis execution and discover that ferroptotic cell rupture is mediated by plasma membrane pores, similarly to cell lysis in pyroptosis and necroptosis. We further find that intercellular propagation of death occurs following treatment with some ferroptosis-inducing agents, including erastin and C′ dot nanoparticles, but not upon direct inhibition of the ferroptosis-inhibiting enzyme glutathione peroxidase 4 (GPX4). Propagation of a ferroptosis-inducing signal occurs upstream of cell rupture and involves the spreading of a cell swelling effect through cell populations in a lipid peroxide- and iron-dependent manner.

The proper regulation of cell death is important for normal organismal development and the maintenance of tissue homeostasis in adulthood. It was once thought that programmed cell death occurred exclusively through apoptosis, whereas necrotic death resulted only from acute cell stress or injury. However, numerous new cell death modalities have recently been discovered, including programmed forms of necrosis that are regulated by specific and distinct cellular machineries. One form of regulated necrosis—ferroptosis—involves the iron-dependent accumulation of lipid peroxide species in cell membranes. Under physiological conditions, ferroptosis is prevented by antioxidant enzymes that limit the build-up of oxidized lipids, including glutathione peroxidase 4 (GPX4), which uses glutathione as a cofactor to detoxify peroxidation products. Cell death can be triggered by GPX4 inactivation, either through direct inhibition or depletion of cellular glutathione, thereby allowing the accumulation of phospholipid peroxides and cell damage. Recent work has uncovered an additional ferroptosis-preventing mechanism controlled by ferroptosis suppressor protein 1 (FSP1), which catalyses the reduction of the lipophilic antioxidant coenzyme Q10 (CoQ)1,24.

Ferroptosis was previously shown to spread through cell populations, resulting in spatiotemporal patterns of cell death with a wave-like appearance not previously observed in other forms of cell death. It is unknown what mechanism underlies this phenomenon and whether death propagation between neighbouring cells is a consistent feature of ferroptosis or occurs only under certain conditions. Given the emerging links between ferroptosis and degenerative diseases that often involve large, continuous areas of tissue damage, the propagative nature of ferroptosis is important to understand. Furthermore, while factors that affect the accumulation of lipid peroxides and thereby modulate ferroptosis have been elucidated, little is known about how lipid peroxidation leads to plasma membrane permeabilization. Whether cell lysis is involved in the intercellular propagation of ferroptosis is also unknown. Here, we investigate the wave-like nature of ferroptosis, the mechanism of ferroptotic cell rupture and the link between the two processes.

We previously observed wave-like spreading of ferroptosis when cells were treated with ferroptosis-inducing nanoparticles called C′ dots (Fig. 1a,b and Supplementary Video 1), and a similar phenomenon has been reported in mouse renal tubules treated with the ferroptosis-inducing agent erastin. However, the spatiotemporal patterns of ferroptosis have not been investigated systematically. To quantitatively study propagation, we performed live-cell imaging of several cell lines (MCF10A mammary epithelium, MCF7 breast cancer, U937 promonocytic leukaemia, HAP1 chronic myelogenous leukaemia and B16F10 melanoma) in the presence of the cell death inducer sytox Green and different ferroptosis-inducing agents (C′ dots, erastin1, the GPX4 inhibitor ML1622,3, or a combination of ferric ammonium citrate (FAC) and thiol-reactive sulfoxamine (BSO); Extended Data Fig. 1a,b). We then used a bootstrapping approach to quantify potential non-random patterns of cell death. For each video, we calculated the mean time difference between neighbouring cell deaths, and compared this experimental value to a distribution of means derived from computationally generated permutations representing random orders of cell death (Fig. 1c,d). Consistent with wave-like propagation, ferroptosis occurred with non-random spatiotemporal patterns when it was induced by erastin, C′ dots or FAC and BSO, as determined by comparing the experimentally determined pattern accumulation value to the 95th percentile of the random distribution. Interestingly, when ferroptosis was induced by inhibition of GPX4 through treatment

1Louis V. Gerstner, Jr. Graduate School of Biomedical Sciences, Memorial Sloan Kettering Cancer Center, New York, NY, USA. 2Cell Biology Program, Sloan Kettering Institute for Cancer Research, New York, NY, USA. 3Department of Software and Information Systems Engineering, Ben-Gurion University of the Negev, Beer-Sheva, Israel. 4Department of Biomedical Engineering, Stanford University, Stanford, CA, USA. 5Department of Materials Science & Engineering, Cornell University, Ithaca, NY, USA. 6Department of Radiology, Sloan Kettering Institute for Cancer Research, New York, NY, USA. 7Molecular Pharmacology Program, Sloan Kettering Institute for Cancer Research, New York, NY, USA. 8E-mail: assafza@bgu.ac.il; overholtz@mail.mskcc.org
Fig. 1 | Ferroptosis exhibits propagative spatiotemporal patterns. a, B16F10 cells treated with C’ dot nanoparticles in amino acid-free (−AA) medium to induce ferroptosis. Images: differential interference contrast (DIC) (left) and SYTOX Green (right). SYTOX-positive cells are dead. Scale bar, 20 μm. Images are representative of five videos from one experiment. b, Nuclei of ferroptotic cells in a, pseudocoloured to indicate the relative timing of cell death, as determined by time-lapse microscopy (Supplementary Video 1). c, Schematic summarizing our method to quantify cell death patterns. Images from time-lapse microscopy (left) are processed to determine the relative timing of neighbouring cell deaths (top right image, ‘experiment’) versus permuted trials (bottom right image, ‘permutation’) to detect potential non-random patterns. The images match the boxed areas in a and b. Scale bar, 10 μm. d, Distribution of time differences between neighbouring deaths (Δt) from the experiment in a–c (blue) versus the averaged distribution of the set of random permutations (orange). The graph shows the fraction of total deaths for the given time differences. e, Spatiotemporal distribution of apoptosis in MCF10A cells treated with TRAIL. Each dot represents a cell; colours indicate relative times of cell death as determined by cell morphology. Data are representative of five fields of view from one experiment. f, Distribution of experimental time differences between neighbouring deaths (Δt) (blue) and averaged distribution of Δt values from the corresponding permuted data in orange. Data belong to the experiment shown in e. g, Ferroptosis, apoptosis and H2O2-induced necrosis show non-random spatiotemporal patterns. µexp Δt is shown versus µperm95 Δt of different cell lines undergoing ferroptosis induced by the indicated treatment (FB = FAC + BSO), apoptosis induced with TRAIL or necrosis induced with H2O2. The dashed line indicates µexp Δt = µperm95 Δt. Each data point represents one video. Data are from two independent experiments for MCF7 + H2O2 and one experiment for all other conditions. h, The spatial propagation index generated from data in g.
with ML162, \( \mu_{\text{exp}} \) was more similar to the 95th percentile of the random permutations (Fig. 1g and Extended Data Fig. 1d,e).

To measure intercellular death propagation in different forms of cell death, we induced necrosis by treatment with hydrogen peroxide (H\(_2\)O\(_2\)) and apoptosis using tumour necrosis factor-related apoptosis-inducing ligand (TRAIL). Although H\(_2\)O\(_2\)-induced necrosis and TRAIL-induced apoptosis displayed no visually obvious wave-like spreading of death (Fig. 1e,f and Supplementary Video 2), they did result in death patterns with non-random spatial features (Fig. 1h). Similarly, H\(_2\)O\(_2\)-induced necrosis and TRAIL-induced apoptosis also do not exhibit propagative features (Fig. 1h). These results demonstrate that the ability to spread in wave-like patterns by propagating between neighbouring cells is a feature of particular forms of ferroptosis.

To examine the mechanism of ferroptotic propagation, we first asked whether iron and lipid peroxidation, two known drivers of ferroptosis, are required for propagation. The addition of the lipid peroxidation inhibitor liproxstatin-1 or the iron chelator deferoxamine (DFO) to cell cultures after the initiation of ferroptosis stopped death from spreading (Fig. 2a–d and Supplementary Video 3), demonstrating that iron and lipid peroxidation are both required for continuous ferroptosis propagation. Because iron and lipid peroxidation are also necessary for ferroptosis to occur in individual cells, these results suggested that the full execution of ferroptosis, including cell lysis, could be required for the spreading of death between cells.
How ferroptosis is executed downstream of lipid peroxidation is not clearly defined. We noted from time-lapse imaging that ferroptotic cells appeared to round and swell before cell death (Fig. 2c). Like cell death, swelling also appeared to spread through cell populations in a manner that was blocked with treatment with liproxstatin-1 or DFO (Supplementary Video 3). Expression of an mKate-tagged version of the zebrafish cPLA2 enzyme, which localizes to the nuclear envelope upon osmotic swelling in HeLa cells, confirmed that ferroptotic cells indeed swell prior to undergoing rupture (Fig. 3d,e). LDH release caused by H$_2$O$_2$-induced death, on the other hand, was not affected by osmoprotectants (Fig. 3c,d).

As ferroptotic cell rupture could be inhibited using osmoprotectants, we sought to examine whether cell lysis is required for ferroptosis propagation. When HAPI1 cells were treated with FAC and BSO in the presence of the osmoprotectant PEG1450, we observed waves of cell rounding that spread through cell colonies and appeared similar to waves of cell death (Supplementary Video 5). However, SYTOX uptake was reduced, consistent with the inhibition of cell rupture. To quantify these waves, we expressed a fluorescent sensor of nuclear calcium (GCaMP6-NLS) in HAPI1 cells, reasoning that pore formation might lead to a spike in intracellular calcium levels that could be used as a readout of cell permeabilization. Live imaging of ferroptotic cells demonstrated that GCaMP fluorescence addition of polyethylene glycols (PEGs) with molecular weights of 1,450 and 3,350 Da (PEG1450 and PEG3350), but not by the smaller osmoprotectants sucrose and raffinose (Fig. 3d,3c). The translocation of cPLA2-mKate to the nuclear envelope was also reduced by PEG1450 and PEG3350 (Fig. 3b), suggesting that ferroptotic swelling and rupture may be caused by the opening of nanoscale pores in the plasma membrane. Induction of ferroptosis with erastin or the GPX4 inhibitors RSL3 and ML162 likewise resulted in cell rupture that was inhibited by treatment with PEG1450 or PEG3350 (Fig. 3d,e). LDH release caused by H$_2$O$_2$-induced death, on the other hand, was not affected by osmoprotectants (Fig. 3c,d).

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Fig. 4 | Ferroptosis spreading involves calcium flux and does not require cell rupture. 

(a) Images show spreading of GCaMP fluorescence (green) before cell rupture marked by SYTOX Orange (red) in HAP1 cells treated with FAC and BSO. The dashed oval shows the origin of death spreading. Note that cells lose GCaMP fluorescence upon cell rupture, probably due to GCaMP efflux. See Supplementary Video 6. Images are representative of three independent experiments. 

(b) Correlation between relative timing of GCaMP fluorescence and SYTOX labelling in HAP1 cells treated with FAC and BSO. Each dot represents a cell and each colour represents a different field of view. Data are from one experiment. Images show spreading of GCaMP fluorescence (green) and SYTOX Orange (red) in HAP1 cells treated with FAC and BSO and PEG1450. The dashed oval shows the origin of death spreading. Note that PEG1450-treated cells maintain GCaMP fluorescence and do not label with SYTOX Orange, unlike the control cells in (a). See Supplementary Video 7. Images are representative of three independent experiments. 

(d) Graph showing $\mu_{\text{exp}}$ versus $\mu_{\text{perm}95}$ of videos of HAP1 cells treated with FAC and BSO and the indicated osmoprotectants, analysed using GCaMP fluorescence. The dashed line indicates $\mu_{\text{exp}} = \mu_{\text{perm}95}$. Each data point represents one video. Data are from one experiment. 

(e) SPI calculated for the experiments shown in (d). All scale bars, 10 µm. Statistical source data are provided in Source Data Fig. 4.

Ferroptosis spreading involves calcium flux and does not require cell rupture. Indeed increased prior to the uptake of SYTOX, and that GCaMP signals spread through cell populations in a similar manner to SYTOX and cell rounding (Fig. 4a and Supplementary Video 6). We compared the relative timing of GCaMP and SYTOX fluorescence for individual cells and found a high degree of correlation, indicating that GCaMP signals could be used instead of SYTOX uptake to assess propagation (Fig. 4b). When cells were treated with PEG1450 to inhibit rupture, wave-like spreading of GCaMP fluorescence still occurred (Fig. 4c and Supplementary Video 7). We quantitatively examined the spatiotemporal GCaMP patterns, and found that their non-random nature was similar to SYTOX death waves in both the presence and absence of osmoprotectants (Fig. 4d,e), demonstrating that propagation occurs in the absence of cell rupture.

Although treatment with osmoprotectants did not prevent propagation, we wondered if it might affect wave speed. To test this we used U937 cells, which exhibit long-lived, unidirectional waves of ferroptosis that can be imaged by differential interference contrast (DIC) microscopy, even in the absence of SYTOX staining (Fig. 5a). Treatment of U937 cells with PEG3350 inhibited cell lysis (Fig. 5b), yet had no effect on the induction of cell death waves (Supplementary Video 8), consistent with the HAP1 data. However, when we measured the speed of these ferroptosis waves, we found them to be slightly but significantly slower in the presence of PEG3350 (1.66 versus 1.37 µm min⁻¹; Fig. 5c), demonstrating that ferroptosis propagation is faster when cells are able to fully lyse.

Together, these data indicate that wave-like spreading is a feature of specific forms of ferroptosis that require the continuous presence of iron and lipid peroxidation, and involve a signal that propagates upstream of cell rupture. Although ferroptosis propagation has been observed previously, here we quantitatively establish the existence of non-random spatiotemporal patterns of ferroptosis in multiple contexts. Our method allowed us to distinguish two types of ferroptosis: cell-autonomous or ‘single-cell ferroptosis’, observed in response to GPX4 inhibition, and propagative or ‘multicellular ferroptosis’, which is induced by treatments that inhibit the generation of glutathione (erastin, BSO) and/or increase cellular iron concentrations (FAC, C’ dots). Why direct GPX4 inhibition does not induce propagative ferroptosis is important to examine in future studies, and may relate to activities of iron or functions of glutathione that do not directly involve GPX4.

Non-autonomous cell death effects have been described elsewhere, most notably in the radiation-induced bystander effect (RIBE), where damage and death rates are increased in cells adjacent to those exposed to radiation. Although RIBE may increase death frequencies, such phenotypes appear distinct from the wave-like death observed during ferroptosis that, in many cases, leads to the near-complete elimination of a cell population. Further discovery of the underlying molecular mechanisms is required to determine whether death propagation in these different systems involves similar signalling mechanisms. Numerous factors are proposed to mediate RIBE, including gap junctions, transforming...
growth factor-β22, p53 and cyclooxygenase-2 signalling23. Although our U937 data suggest that gap junctions are not involved in ferroptosis propagation, because these cells do not form cell junctions (Supplementary Video 8), whether other RIBE signals could play a role in ferroptosis spreading is not yet known. Our finding that the presence of an osmoprotectant slows propagation could suggest that the release of a spreadable factor is enhanced by cell rupture, although further experiments are needed to test this.

Our data also indicate that ferroptosis is an osmotic process, as it involves cell swelling (Fig. 5d), and can be blocked by the addition of large osmoprotectants. The ability of osmoprotectants to block lysis following induction of necroptosis or pyroptosis, both in culture and in vivo, and the observed size dependence of the protective effects of different osmoprotectants (Fig. 3), have been interpreted previously as evidence for the existence of pore-like structures that trigger lysis in these forms of necrosis22,24. This is indeed known to occur during pyroptosis, in which the caspase-dependent cleavage of gasdermin D triggers its oligomerization in the plasma membrane25,26. Similarly, necroptosis may involve plasma membrane permeabilization mediated by the pseudokinase MLKL27,28. Our data thus suggest that ferroptotic rupture is mediated by the formation of plasma membrane pores of a few nanometres in diameter and that cell permeabilization during ferroptosis could be a regulated process. Intriguingly, lipid peroxidation has been proposed to lead to conformational changes in lipid domains and plasma membrane regions29,30, raising the possibility that pore formation could occur through a lipid-based mechanism rather than by activation of a pore-forming protein. Ferroptotic pore formation could regulate not only cell death execution but also the potential release of pro-inflammatory cytokines or DAMPs, which is known to occur during pyroptosis31. We have previously shown that ferroptosis induction in mouse xenografts leads to tumour regression and a concomitant immune response, implying that ferroptosis-inducing agents may be promising cancer therapies32,33. Ferroptosis is also
implicated in cell death resulting from ischaemia reperfusion injury during stroke or myocardial infarction, as well as in acute kidney injury, all of which result in the formation of large zones of necrotic tissue, possibly indicating a role for ferroptosis propagation in these diseases\(^1\). Intriguingly, the paper by Katikaneni et al. published in this issue shows large waves of cellular deformation occurring in intact zebrafish larvae following microperfusion of arachidonic acid\(^1\). As arachidonic acid is a known driver of ferroptosis, this finding suggests that wave-like propagation of ferroptosis may also occur in vivo, causing widespread tissue damage\(^1\). Uncovering the molecular mechanisms that regulate ferroptosis execution and propagation through cell populations will ultimately further our understanding of how modulators of ferroptosis may be leveraged for therapeutic benefit.

**Online content**

Any methods, additional references, Nature Research reporting summaries, source data, extended data, supplementary information, acknowledgements, peer review information; details of author contributions and competing interests; and statements of data and code availability are available at [https://doi.org/10.1038/s41556-020-0565-1](https://doi.org/10.1038/s41556-020-0565-1).

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**References**

1. Stockwell, B. R. et al. Ferroptosis: a regulated cell death nexus linking metabolism, redox biology and disease. *Cell* **171**, 273–285 (2017).

2. Dixon, S. J. et al. Ferroptosis: an iron-dependent form of nonapoptotic cell death. *Cell* **149**, 1060–1072 (2012).

3. Ros, U. et al. Necroptosis execution is mediated by plasma membrane nanopores independent of calcium. *Cell Rep.* **19**, 175–187 (2017).

4. Fink, S. L. & Cookson, B. T. Caspase-1-dependent pore formation during pyroptosis leads to osmotic lysis of infected host macrophages. *Cell Microbiol.* **8**, 1812–1825 (2006).

5. Degterev, A. et al. Chemical inhibitor of nonapoptotic cell death with therapeutic potential for ischemic brain injury. *Nat. Chem. Biol.* **1**, 112–119 (2005).

6. Cookson, B. T. & Brennan, M. A. Pro-inflammatory programmed cell death. *Trends Microbiol.* **9**, 113–114 (2001).

7. Linkermann, A. et al. Synchronized renal tubular cell death involves ferroptosis. *Proc. Natl Acad. Sci. USA* **111**, 16886–16891 (2014).

8. Kim, S. E. et al. Ultrasound nanoparticles induce ferroptosis in nutrient-deprived cancer cells and suppress tumour growth. *Nat. Nanotechnol.* **11**, 977–985 (2016).

9. Dolma, S., Lessnick, S. L., Hahn, W. C. & Stockwell, B. R. Identification of genotype-selective antitumor agents using synthetic lethal chemical screening in engineered human tumor cells. *Cancer Cell* **3**, 285–296 (2003).

10. Yang, W. S. et al. Regulation of ferroptotic cancer cell death by GPX4. *Cell* **156**, 317–331 (2014).

11. Galluzzi, L. et al. Molecular mechanisms of cell death: recommendations of the Nomenclature Committee on Cell Death 2018. *Cell Death Differ.* **25**, 486–541 (2018).

12. Seibt, T. M., Proneth, B. & Conrad, M. Role of GPX4 in ferroptosis and its pharmacological implication. *Free Radic. Biol. Med.* **133**, 144–152 (2019).

13. Bersuker, K. et al. The CoQ oxidoreductase FSP1 acts parallel to GPX4 to inhibit ferroptosis. *Nature* **575**, 688–692 (2019).

14. Doll, S. et al. FSP1 is a glutathione-independent ferroptosis suppressor. *Nature* **575**, 693–698 (2019).

15. Riegman, M., Bradbury, M. S. & Overholtzer, M. Population dynamics in cell death: mechanisms of propagation. *Trends Cancer* **5**, 558–568 (2019).

16. Yang, W. S. & Stockwell, B. R. Ferroptosis: death by lipid peroxidation. *Trends Cell Biol.* **26**, 165–176 (2016).

17. Feng, H. & Stockwell, B. R. Unsolved mysteries: how does lipid peroxidation cause ferroptosis? *PLoS Biol.* **16**, e2006203 (2018).

18. Rittker, J. A. et al. in *Probe Reports from the NIH Molecular Libraries Program* (Bethesda, MD, National Center for Biotechnology Information, 2010); [https://www.ncbi.nlm.nih.gov/books/NBK55069/](https://www.ncbi.nlm.nih.gov/books/NBK55069/).

19. Enyedi, B., Jelic, M. & Niethammer, P. The cell nucleus serves as a mechanotransducer of tissue damage-induced inflammation. *Cell* **165**, 1160–1170 (2016).

20. Cao, J. Y. et al. A genome-wide haploid genetic screen identifies regulators of glutathione abundance and ferroptosis sensitivity. *Cell Rep.* **26**, 1544–1556 (2019).

21. Zhou, H. et al. Mechanism of radiation-induced bystander effect: role of the cyclooxygenase-2 signaling pathway. *Proc. Natl Acad. Sci. USA* **102**, 14641–14646 (2005).

22. Iyer, R., Lehner, B. E. & Svensson, R. Factors underlying the cell death–related bystander responses to alpha particles. *Cancer Res.* **60**, 1290–1298 (2000).

23. Chen, X. et al. Pyroptosis is driven by non-selective gasdermin-D pore and its morphology is different from MLKL channel-mediated necroptosis. *Cell Res.* **26**, 1007–1020 (2016).

24. Sborgi, L. et al. GSDMD membrane pore formation constitutes the mechanism of pyroptotic cell death. *EMBO J.* **35**, 1766–1778 (2016).

25. Wang, H. et al. Mixed lineage kinase domain-like protein MLKL causes necrotic membrane disruption upon phosphorylation by RIP3. *Mol. Cell* **54**, 133–146 (2014).

26. Zhang, Y., Chen, X., Gueydan, C. & Han, J. Plasmin membrane changes during programmed cell death. *Cell Res.* **28**, 9–21 (2018).

27. Agmon, E., Solon, J., Bassereau, P. & Stockwell, B. R. Modeling the effects of lipid peroxidation during ferroptosis on membrane properties. *Sci. Rep.* **8**, 5158 (2018).

28. Runas, K. A., Acharya, S. J., Schmidt, J. J. & Malmstadt, N. Addition of cleaved tail fragments during lipid oxidation stabilizes membrane permeability behavior. *Langmuir* **32**, 779–786 (2016).

29. Evavold, C. L. et al. The pore-forming protein gasdermin D regulates interleukin-1 secretion from living macrophages. *Immunity* **48**, 35–44 e36 (2018).

30. Katikaneni, A. J. M., Gerlach, G., Ma, Y., Overholtzer, M. & Niethammer, P. Lipid peroxidation instructs long-range wound detection through 5-lipoxygenase in zebrafish. *Nat. Cell. Biol.* (in the press).

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Methods

Cell culture. HT1080 cells (ATCC), HeLa cells (ATCC), HAP1 chronic myelogenous leukemia cells (the kind gift of J. Carrete, Stanford School of Medicine), and MCF7 breast cancer cells (Lombard Cancer Center, Georgetown University) were cultured in high-glucose Dulbecco’s Modified Eagle’s Medium (DMEM) (MSKCC Media Preparation Facility) supplemented with 10% heat-inactivated fetal bovine serum (FBS) (F2442; Sigma-Aldrich) and penicillin/streptomycin (30-002-CL; Mediatech). B16F10 melanoma cells (ATCC) and U937 cells expressing MSH-tagged Cre (H-0888; Sigma-Aldrich), 100 ng/mL cholera toxin (C-8052; Sigma-Aldrich), and GCaMP6-NLS were generated using the Sleeping Beauty transposase system. HAP1 cells were transfected with pSb-cMV-MCS-Puro GCaMP6-NLS and pS8-cag-100x-Transposase, using Lipofectamine 3000 (L3030-015; Thermo Fisher) as recommended by the manufacturer. Amino acid-free culture medium was prepared by dialyzing FBS or HBS in PBS (MSKCC Media Preparation Facility) using MWCO 3500 dialysis tubing (21-152-9; Fisherbrand) and adding it to amino acid-free base media (MSKCC Media Preparation Facility). These media were used in combination with FBS and BSO to induce ferroptosis in MCF10A and MCF7 cells, and with nSMase-tagged Cα dots to induce ferroptosis in B16F10 cells.

Reagents. Ferric ammonium citrate (FAC) (F879; Sigma-Aldrich), l-buthionine sulfoximino (BSO) (B2515; Sigma-Aldrich), and deferoxamine (DFO) (D9533; Sigma-Aldrich) were dissolved in water, and stock solutions were stored at 20 °C. FAC and BSO were used at 400 µM and DFO at 200 µM. SuperKillerTRAIL (ALX-201-115-C100; Enzo) stock solution was diluted to 100 µg/mL in killerTRAIL dilution buffer (20 mM HEPES, 300 mM NaCl, 0.01% Tween-20, 1% sucrose, 1 mM DTT), aliquoted, and stored at −80 °C. This was used at a final concentration of 30 ng/mL. Hydrogen peroxide (216763; Sigma-Aldrich) was first diluted in water and then added to cell culture media at a final concentration of 1 mM. C11-BODIPY581/591 (D3681; Molecular Probes) was dissolved in DMSO, stored at −20 °C, and diluted to a final concentration of 5 µM prior to use. Sucrose (S57903; Sigma-Aldrich), raffinose (R0514; Sigma-Aldrich), polyethylene glycol (PEG) 1450 (P7181; Sigma-Aldrich), and PEG3550 (P3640; Sigma-Aldrich) were dissolved directly into cell culture media at a concentration of 20 mM. All other compounds were prepared as stock solutions in DMSO and stored at −20 °C. Eritest (E7781, used at 7.5 µM with HAP1s and 2 µM with HT1080s), ferrostatin-1 (SML0583, used at 4 µM), and Trolox (238813, used at 100 µM) were from Sigma-Aldrich; RSL3 (S8155, used at 1 µM) and liproxstatin-1 (S7699, used at 200 µM) were from SelleckChem. ML162 was synthesized by Acme Bioscience (Palo Alto, CA) and was used at 4 µM. nSMase-tagged Cα dots were synthesized as described previously25 in water, stored at 4 °C, and used at a concentration of 5 µM. SYTOX Green (S7070; Molecular Probes) was used at a concentration of 10 nM and SYTOX Orange (S11368; Molecular Probes) at a concentration of 50 nM for all experiments.

Analysis. Single-cell time of death annotation: we used a custom MATLAB (R2016A) script to determine the xy position at points where each locus corresponds to a cell nucleus’s xy location. Each point on the plane belongs to a single region, and each region consists of all the closest points to the associated nuclear locus. Neighbors were defined as regions sharing a border, which allowed us to identify all pairs of neighboring cells. Note that neighboring cells in the tessellation do not necessarily share cell–cell junctions in the spatial component to the observed experimental cell–cell propagation. This was achieved by using a non-parametric permutation test to reject the null hypothesis that the cells’ time of death is independent of their neighbors’ time of death. For every field of view, the following procedure was repeated for 1,000 iterations. The recorded time of death was randomly permuted between the cells (i.e., each cell was assigned a random time of death, with the same number of deaths at each time point as in the experimental data), and the mean difference in time of death between all neighboring cell pairs was recorded. The P value was calculated as the fraction of iterations in which the mean time difference between neighbors using the permuted (“random”) cell death was faster than the experimental measurement. We considered a P value of 0.05 as statistically significant.

We devised the spatial propagation index to quantify the contribution of the spatial component to the observed experimental cell–cell propagation. This measure was defined as the deviation of the experimental mean propagation (\(\mu_{\text{exp}}\)) from the 95th percentile of the mean randomly permuted death times (\(\mu_{\text{perm}}\)) normalized to \(\mu_{\text{perm}}\): \(\frac{\mu_{\text{exp}} - \mu_{\text{perm}}^{0.95}}{\mu_{\text{perm}}}.\) This measure can be interpreted as the fraction of the \(\mu_{\text{perm}}\) needed to reconstruct back the spatial information in the experimental data.

Quantifying ferroptosis propagation from DIC: to quantify the distance covered by the ferroptosis wave in each field of view, lines delineating live and dead cell regions were drawn manually in NIS elements AR (Nikon, 3.22.15) at 1-h intervals, starting from a timepoint at which smaller initiation points, if present, had converged into larger waves. For each interval, the area between two lines was measured and divided by their average length to obtain the mean distance traveled during a given interval in a given field of view. Information from different fields of view in the same condition was then combined to plot the mean distance covered in each condition, and a linear regression was performed to calculate the speed of the corresponding ferroptosis wave.

Live-cell imaging. Cells were seeded on glass-bottom plates (P24G-1.5-13-F; Mattek) and treated in fresh culture media the next day. For amino acid-free conditions, cells were washed with PBS twice prior to treatment. Imaging was performed in live-cell incubation chambers maintained at 37 °C and 5% CO2. Images were acquired every 5 to 30 min for 24–48 h using a Nikon Ti-E inverted microscope attached to a CoolSNAP charge–coupled device camera (Photometrics) and NIS Elements AR software (Nikon, 3.22.15). For experiments with ML162, time-lapse imaging was performed on plastic tissue culture plates (3527; Corning) on a Zeiss microscope using ZEN software (Zeiss). Images were quantified manually in NIS Elements AR (Nikon) or ZEN (Zeiss) and processed using ImageJ (2.0.0) and Adobe Photoshop (CS6 13.0.5).

LDH assays. LDH assays were performed using the Pierce LDH CytoToxicity Kit (88954; Thermo Fisher) following the manufacturer’s instructions. Briefly, cells were seeded on 96-well plates and treated in triplicate the next day. At the indicated time, 50 µL media was transferred from each well to a well containing 50 µL assay buffer, and plates were incubated at RT for 30 min. At this point 50 µL stop solution was added to each well, bubbles were removed using a syringe, and the absorbance was read on a BioTek Synergy H1 Hybrid Reader at 490 nm and 680 nm wavelengths. The 490 nm absorbance was subtracted from the 680 nm absorbance, and background from cell culture medium was subtracted from all values. Data were averaged across technical replicates and normalized to the indicated treatment without osmoprotectants to calculate percent LDH release. For suspension cells, cells were treated in 24-well tissue culture plates containing 1 mL media per well. At the indicated time, 200 µL media was taken from each well and spun down to remove dead cells, and the supernatant was used to perform the assay as described.

Crystal violet assays. Cells were seeded on 24-well tissue culture plates (3527; Corning) and treated in triplicate the next day. After 24 h, when most control cells had died, cells were washed twice with PBS, then fixed in 4% paraformaldehyde (15710-S; Electron Microscopy Sciences) in PBS for 15 min. After washing with water, cells were stained with 0.1% crystal violet (61135; Sigma-Aldrich) in 10% ethanol for 20 min, then washed again with water until clear and allowed to air dry. After drying, crystal violet stain was dissolved in 10% acetic acid by shaking the plate at room temperature for 30 min. This solution was diluted 1:4 with water, and absorbance was measured at 590 nm on a BioTek Synergy H1 Hybrid Reader. Background absorbance from wells containing only tissue culture medium was subtracted from all readings, and values were averaged across technical replicates and normalized to the untreated controls to obtain percent viability.

Confocal imaging. Cells were plated on glass-bottom dishes and treated the next day. For C11-BODIPY583/591 imaging, cells were washed twice with PBS, then fixed in 4% paraformaldehyde (15710-S; Electron Microscopy Sciences) in PBS for 15 min. After washing with water, cells were stained with 0.1% crystal violet (61135; Sigma-Aldrich) in 10% ethanol for 20 min, then washed again with water until clear and allowed to air dry. After drying, crystal violet stain was dissolved in 10% acetic acid by shaking the plate at room temperature for 30 min. This solution was diluted 1:4 with water, and absorbance was measured at 590 nm on a BioTek Synergy H1 Hybrid Reader. Background absorbance from wells containing only tissue culture medium was subtracted from all readings, and values were averaged across technical replicates and normalized to the untreated controls to obtain percent viability.

Statistics and reproducibility. Data were analyzed in Microsoft Excel (Office 2011) and GraphPad Prism 7 and are represented as mean with individual data points. P values were obtained using two-sided Dunnett’s multiple comparisons test unless otherwise indicated. *P < 0.05. **P < 0.01. ***P < 0.001. ****P < 0.0001. The following analyses were done using data from one experiment: spatiotemporal
analyses in Figs. 1d–h, 4b, d, e, and Extended Data Fig. 1c,e, except data for MCF7+H2O2 in Fig. 1g, h which was from two independent experiments. The images in Fig. 1a are representative of one experiment, and images in Fig. 2f are representative of two independent experiments. All other data were derived from three or more biologically independent experiments; exact n are reported in the figure legends. For LDH measurements in Figs. 3a–d and Fig. 5b, and crystal violet assay in Extended Data Fig. 1a, each experiment consisted of three technical replicates that were averaged for each condition.

**Reporting Summary.** Further information on research design is available in the Nature Research Reporting Summary linked to this article.

**Data availability**
The statistical source data that support the findings of this study have been provided as part of this publication. All other data are available from the corresponding authors upon request. Source data are provided with this paper.

**Code availability**
Our source code is available via GitHub at https://github.com/AssafZaritskyLab/PropagationOfCellDeath. This repository includes all code used to measure the mean time difference between neighboring deaths and to run the random simulations, as well as a demo dataset.

**References**
31. Ma, K. et al. Control of ultrasmall sub-10 nm ligand-functionalized fluorescent core–shell silica nanoparticle growth in water. *Chem. Mater.* **27**, 4119–4133 (2015).
32. Du, Q. & Gunzburger, M. Grid generation and optimization based on centroidal Voronoi tessellations. *Appl. Math. Comput.* **133**, 591–607 (2002).

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**Author contributions**
M.R. and M.O. designed the study. M.R. designed, performed and analysed experiments. L.S., C.G., T.L., N.S. and A.Z. wrote the analysis code and performed computational analyses. M.R., A.Z. and M.O. wrote the paper. S.J.D., U.W., M.S.B. and P.N. provided key reagents and edited the manuscript.

**Competing interests**
Memorial Sloan-Kettering Cancer Center and three investigators involved in this study (M.S.B., U.W. and M.O.) have financial interests in Elucida Oncology. Research involving C′ dots may involve one or more US or international patent applications.

**Additional information**
Extended data is available for this paper at https://doi.org/10.1038/s41556-020-0565-1.

**Supplementary information** is available for this paper at https://doi.org/10.1038/s41556-020-0565-1.

**Correspondence and requests for materials** should be addressed to A.Z. or M.O.

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Extended Data Fig. 1 | Treatment of cells with FAC and BSO induces ferroptosis. a, Viability of HAP1 cells after treatment with FAC and BSO and either DMSO or ferroptosis inhibitors as measured by crystal violet staining. N=three independent experiments. Dunnett’s test; *p = 0.0024 for Lip-1; "p = 0.0045 for Fer-1; ‘p = 0.0107 for Trolox. b, Confocal images of HAP1 cells treated with FAC and BSO and stained with C11-BODIPY581/591. Non-oxidized probe is shown in red, oxidized probe is shown in green (arrow). Scale bar = 10 μm. Images are representative of three independent experiments. c, Values from the analysis of the experiment shown in panels 1c and d. Note that the experimental mean time difference between neighbors (μ_{exp} Δt) is much smaller than the mean (μ_{perm} Δt) and 95th percentile (μ_{95perm} Δt) obtained from the randomly permuted data. (d) Spatiotemporal distribution of cell death in HAP1 cells treated with ML162 to induce ferroptosis. Each dot represents a cell from a single movie representative of five fields of view from one experiment. Colors indicate relative times of cell death as determined by SYTOX Green staining. e, Distribution of experimental time differences between neighboring deaths in blue and averaged distribution of the corresponding permuted data in orange. Data belong to the experiment shown in panel d and are representative of five fields of view from one experiment. Statistical source data can be found at Source data Extended Data Fig. 1.
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Sample size  Sample size was determined based on published literature and previous experience. All cell biological experiments were performed as at least three replicates, or more when greater variability in outcomes was observed to ensure reproducibility. No statistical methods were used to predetermine sample sizes.

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Cell line source(s)  MCF7 cells were obtained from Lombardi Cancer Center, Georgetown University, which is the original source of this cell line. H11080 cells, Hela cells, 316L/10 cells, U937 cells, and MCF10A cells were obtained from ATCC. HAP2 cells were a kind gift from Dr. Jan Carette [Stanford School of Medicine], who originally developed this cell line.

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