

# New perspectives on ferroptosis and its role in hepatocellular carcinoma

Tianhao Cong, Yingen Luo, Yan Fu, Yu Liu, Yujie Li, Xiao Li

Department of Interventional Therapy, National Cancer Center/National Clinical Research Center for Cancer/Cancer Hospital, Chinese Academy of Medical Sciences and Peking Union Medical College, Beijing 100021, China.

## Abstract

For a long time, the morbidity and mortality rates of hepatocellular carcinoma (HCC) have remained high. Since the concept of ferroptosis was introduced in 2012, researchers' perspectives have shifted toward finding novel ferroptosis-related treatment strategies, especially for tumors that are resistant to apoptosis. In recent years, there have been an increasing number of studies on ferroptosis, and these studies have found that ferroptosis has great potential and promise for cancer treatment. Ferroptosis is a kind of regulated cell death (RCD); unlike apoptosis, ferroptosis is an iron-dependent type of RCD driven by lipid peroxidation. The whole process of ferroptosis mainly revolves around three pathways (system  $x_c^-$ /glutathione peroxidase 4 [GPX4]), lipid peroxidation, and iron metabolism), which are also regulated by various metabolic factors. This review will attempt to analyze the relationship between the system  $x_c^-$ /GPX4 pathway, lipid peroxidation, iron metabolism, and ferroptosis from three aspects (triggering, execution, and regulation), and the regulatory factors for ferroptosis will be summarized. In this review, we will also illustrate the relationship between ferroptosis and tumors as well as its application in tumors from the perspective of HCC. Finally, we will summarize the current limitations and needs and provide perspectives related to the focus of development in the future.

**Keywords:** Ferroptosis; Hepatocellular carcinoma; Molecular targets

## Introduction

Primary liver cancer is the sixth most commonly diagnosed cancer and ranks third in terms of mortality rate; hepatocellular carcinoma (HCC) accounts for 75% of primary liver cancers.<sup>[1,2]</sup> In the long-term fight against cancer, we are committed to exploring strategies for the specific elimination of cancer cells while retaining normal cells. To date, most clinical anticancer drugs cause cancer cell apoptosis. Apoptosis is a type of programmed cell death, which is a constitutive suicide process cells undergo that can be regulated by various internal or external signals.<sup>[3]</sup> However, apoptosis resistance is a general feature of cancer that forces us to look for new opportunities to exploit regulated cell death (RCD), which is a broader concept.<sup>[4]</sup> RCD refers to cell death that is regulated by molecules or signals but can develop without going through a specific procedure.<sup>[5]</sup> Ferroptosis is an iron-dependent type of RCD driven by lipid peroxidation. Ferroptosis is distinct from apoptosis, necrosis, and autophagy in terms of morphological changes, biochemical characteristics and genetic standards.<sup>[6]</sup> With the increase in research on ferroptosis, there is

an increasing evidence that ferroptosis may play a physiological role in the occurrence and development of tumors, which is particularly important, especially in the era of drug resistance. In this paper, we first introduced the factors mediating the triggering, execution and regulation of ferroptosis at the molecular level and then introduced the relationship of ferroptosis with HCC, including its application in HCC. Finally, we discussed the prospects regarding the development of ferroptosis.

## Molecular Mechanism of Ferroptosis

The concept of ferroptosis was described by Dixon *et al*<sup>[7]</sup> in 2012. However, Yang and Stockwell<sup>[8]</sup> described the phenomenon of ferroptotic cell death for the first time in a high-throughput screening study to identify compounds that can be selectively lethal to cells carrying ras sarcoma (RAS) mutant subtypes as early as 2008. Dixon *et al*<sup>[7]</sup> found that erastin can induce a regulated but nonapoptotic form of cell death, which is accomplished by phospholipid peroxidation, and this process depends on the reactive oxygen

### Access this article online

Quick Response Code:



Website:  
www.cmj.org

DOI:  
10.1097/CM9.0000000000002327

**Correspondence to:** Professor Xiao Li, Department of Interventional Therapy, National Cancer Center/National Clinical Research Center for Cancer/Cancer Hospital, Chinese Academy of Medical Sciences and Peking Union Medical College, Beijing 100021, China  
E-Mail: simonlixiao@gmail.com

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Chinese Medical Journal 2022;135(18)

**Received:** 21-02-2022; **Online:** 09-11-2022 **Edited by:** Xiuyuan Hao and Rongman Jia

species (ROS), polyunsaturated fatty acids (PUFAs), and iron metabolic products. Glutathione peroxidase 4 (GPX4) and system  $x_c^-$  are the two most important pathways and are involved in nearly the entire mechanism of triggering, executing, and regulating ferroptosis; thus, they have a direct impact on this process. Iron and lipid peroxidation also play vital roles in ferroptosis. The regulatory factors of ferroptosis, such as ferroptosis suppressor protein 1 (FSP1) and nuclear factor erythroid 2-related factor 2 (NRF2), were also briefly summarized in this article.

### System $x_c^-$ and GPX4

System  $x_c^-$  is an amino acid antiporter that belongs to the heteromeric amino acid transporter (HAT) family, whose function is to mediate the exchange of extracellular L-cystine and intracellular L-glutamate across the cellular plasma membrane.<sup>[9]</sup> It is a heterodimer formed by the 12-pass transmembrane transporter protein solute carrier family 7 member 11 (SLC7A11) and the single-pass transmembrane regulatory protein solute carrier family 3 member 2 (SLC3A2) with a disulfide bond as a link.<sup>[10]</sup> It is necessary to discuss GPX4 before introducing the role of system  $x_c^-$ . GPX4 is a restricted selenoprotein that is important for the survival of mammalian cells and can catalyze the reduction of lipid peroxides (LPOs) in a complex cell membrane environment.<sup>[11,12]</sup> It can restrain the activity of lipoxygenase (LOX) and resist lipid peroxidation by converting phospholipid hydroperoxide (PLOOH) to the corresponding alcohol phospholipid hydroxide.<sup>[13]</sup> A number of studies have shown that the inhibition of GPX4 can increase lipid peroxidation and eventually lead to ferroptosis, which also illustrates the source of LPOs in normal cells.<sup>[14-16]</sup> Research has also shown that GPX4 is the central regulatory factor of ferroptosis and one of the most fatal signals of ferroptosis. Notably, to play a corresponding role in the physiological state, glutathione (GSH) acts as the basic auxiliary factor for GPX4, and cysteine is a precursor of GSH synthesis.<sup>[12,17]</sup> Simultaneously, system  $x_c^-$  can regulate the level of intracellular cysteine directly, thus controlling the occurrence of ferroptosis by affecting the activity of GPX4. In addition, research has shown that SLC7A11 is regulated by multiple genes at the same time; for example, *NRF2* can promote its expression and activation,<sup>[18]</sup> while some tumor suppressor genes, such as *p53*, breast cancer susceptibility gene 1 associated protein 1 (*BAP1*), and beclin 1 (*BECN1*), play a negative regulatory role.<sup>[19-21]</sup> System  $x_c^-$  and GPX4 expression can be restrained by erastin and RSL3, respectively.<sup>[14,22]</sup>

### Lipid peroxidation

Previous studies have revealed that the association between cellular metabolism and ferroptosis is particularly complex.<sup>[23-26]</sup> It has been confirmed that ferroptosis is driven by lipid peroxidation.<sup>[7]</sup> Lipid peroxidation occurs via a sequential process (initiation, propagation, and termination), and these three stages are closely linked.<sup>[27,28]</sup> However, when GPX4 cannot eliminate redundant LPOs for some reasons, such as the inhibition of RSL3, LPOs will be converted to an alkoxyl radical (LO·) under the premise of the action of ferrous iron, and

then LO· reacts with PUFAs, ultimately entering into another lipid radical chain reaction.<sup>[27]</sup> Consequently, the lipid peroxidation process cannot enter the final termination phase, which promotes the accumulation of LPOs, such as malondialdehyde (MDA), eventually leading to toxicity or the initiation of cell death cascades.<sup>[27,29]</sup> LOXs are nonheme, iron-containing enzymatic protein effectors, whose primary substrates are PUFAs. They play an important role in lipid peroxidation in ferroptosis, and LOX knockout can resist ferroptosis mediated by erastin.<sup>[30,31]</sup> In addition to the reported long-chain fatty acid-coenzyme A (CoA) ligase 4 (ACSL4)/lysophospholipid-acyltransferase 3 (LPCAT3)/15-LOX pathway involved in the production of LPO during ferroptosis,<sup>[15,32]</sup> a recent study found that the *p53*/SLC7A11/12-LOX pathway was also related to the production of LPO.<sup>[29]</sup> Although we understand the metabolic process of lipid peroxidation in ferroptosis, it is not clear how to induce cytotoxicity via lipid peroxidation or which mechanism regulates cell death in this context. Lipid peroxidation in the course of ferroptosis is a multicascade process that is organized by the balance between the oxidation system and the antioxidant system, lipid scavenging mechanisms, autophagy, and membrane repair mechanisms.<sup>[33]</sup> In addition, since GPX4 also plays a crucial role in other RCD processes,<sup>[34-36]</sup> it suggests that the “intersection” position of lipid peroxidation in RCD is reasonable.<sup>[33]</sup>

### Iron metabolism

The accumulation of intracellular iron due to various conditions, such as increased iron absorption, enhanced release or limited intracellular iron outflow, is capable of promoting the occurrence of ferroptosis through comprehensive pathways.<sup>[37]</sup> Why is this process iron-dependent? First, the Fenton reaction, which is a non-enzymatic reaction chain, is an indispensable step for the occurrence of ferroptosis: the Fenton reaction can enhance PLOOH production, which is a landmark event of ferroptosis.<sup>[38]</sup> In addition, iron is an essential component of some enzymes involved in ferroptosis; for example, phospholipid peroxidation metabolism-related LOX and Cytochrome P450 Reductase (POR) need iron to support catalysis, and iron is also vital for a large number of metabolic enzymes involved in the production of ROS in cells.<sup>[16]</sup> Therefore, metabolic regulators related to iron metabolism, including iron uptake, storage, transport, or degradation, may affect ferroptosis. Among them, the binding of iron regulatory proteins (IRPs) to the iron response element (IRE) in mRNA is the primary pathway regulating the transport, storage, and release of iron,<sup>[39,40]</sup> and IRE-binding protein 2 (IRE-BP2) is considered to be a very important factor in inducing ferroptosis.<sup>[41]</sup> Transferrin (TF) can bind to iron in the blood and transport it to various tissues and organs. Research by Gao *et al*<sup>[23]</sup> has shown that TF is necessary for inducing ferroptosis, which is also a crucial factor for ferroptosis induced by amino acid starvation. Transferrin receptor (TFR) is a membrane protein that is responsible for the cellular uptake of iron-loaded TF and plays a vital role in mediating ferroptosis.<sup>[41]</sup> Oncogenic RAS can enrich the iron pool of cells by promoting the function of TFR, thus increasing the sensitivity of cells to ferroptosis induced by erastin.<sup>[8]</sup> In addition, a study has shown that the

accumulation of TFR exists only in the process of ferroptosis, not apoptosis.<sup>[42]</sup> The degradation of ferritin can be accomplished by selective autophagy mediated by nuclear receptor coactivator 4 (NCOA4), and the knockout of *NCOA4* can inhibit ferroptosis induced by erastin in fibroblasts and pancreatic cancer cells by blocking ferritin degradation.<sup>[43]</sup> In addition, other proteins or regulators related to iron metabolism, such as heme oxygenase 1 (HO1) and prominin-2 (PROM2), may affect the process of ferroptosis.<sup>[44-46]</sup>

### Regulatory Factors of Ferroptosis

At present, the known regulatory signaling pathways related to ferroptosis are very complex, and there are also many regulatory factors. In recent years, FSP1 has become a research hotspot. FSP1, formerly known as apoptosis inducing factor mitochondria associated 2 (AIFM2), was once thought to be a proapoptotic protein<sup>[47]</sup> Doll *et al*<sup>[48]</sup> found for the first time that the inhibitory effect of FSP1 on ferroptosis was not affected by intracellular glutathione, the level or activity of GPX4, long-chain fatty acyl-CoA ligase 4 (ACSL4) expression, or other factors, which shows the particularity of the inhibition mechanism of FSP1 on ferroptosis. Their research showed that this kind of mechanism was mediated by coenzyme Q10 (CoQ10), and FSP1-CoQ10-nicotinamide adenine dinucleotide (NAD[P]H) existed as an independent parallel pathway, which cooperated with the GPX4 pathway and glutathione to restrain phospholipid peroxidation and ferroptosis.<sup>[48]</sup> Moreover, Bersuker *et al*<sup>[49]</sup> found that the expression of FSP1 was positively correlated with ferroptosis resistance in hundreds of cancer cell lines, and they also confirmed that FSP1 is a key component of a non-mitochondrial CoQ10 antioxidant system and works in parallel with the GPX4 pathway. Therefore, FSP1 is a promising candidate for targeting ferroptosis.<sup>[48-50]</sup> Regulatory factors of ferroptosis are listed in [Table 1].

In general, ferroptosis mainly involves three types of pathways: the system  $x_c^-$ -GPX4 pathway, the lipid peroxidation pathway and the iron metabolism pathway. Most other regulatory signals can affect ferroptosis by targeting these three pathways. Strikingly, although we have made great progress in the study of the triggering mechanism of ferroptosis, it is still unclear what kind of mechanism mediates cell death after ferroptosis. Is the cytotoxicity of lipids or the accumulation of ROS dominant? Does it lead to the loss of the integrity of

the cell membrane or mitochondrial membrane? Or does it lead to the destruction of some cellular key macromolecules? In our opinion, exploration of the execution mechanism and process of ferroptosis is needed and will provide great contributions to the clinical translation and application of ferroptosis. Next, we discussed the link between ferroptosis and HCC and its application.

### Molecular Targets of Ferroptosis in HCC

Currently, research on ferroptosis in the field of cancer has attracted an increasing attention, and one of the main directions is exploring the intricate and close relationship between cancer and ferroptosis. Many studies have shown that various tumor suppressors can regulate the sensitivity of cancer cells to ferroptosis. For example, *p53* can enhance the sensitivity of cells to ferroptosis by inhibiting the SLC7A11 subunit and controlling cell metabolism and the redox state;<sup>[19,51-53]</sup> in contrast, the inhibitory effect of *p53* on ferroptosis has been reported.<sup>[54,55]</sup> Tumor suppressors can also downregulate the expression of SLC7A1.<sup>[20]</sup> On the other hand, a study has shown that ectopic expression of oncogenic *RAS* mutants can reduce the sensitivity of rhabdomyosarcoma (RMS)13 cells to ferroptosis.<sup>[56]</sup> The occurrence of HCC involves alterations in specific gene expression patterns caused by related mutant genes, including mutant *RAS*, *p53*, retinoblastoma (*Rb*), *NRF2*/Kelch-like ECH-associated protein 1 (*Keap1*), and other genes or common changes in signaling pathways.<sup>[57]</sup>

The p62-Keap1-NRF2 pathway has been the focus of research in recent years. It has been reported that an imbalance in the p62-Keap1-NRF2 pathway can be observed in both HCC patients and mouse liver cancer models.<sup>[58,59]</sup> p62 is the key component of Mallory-Denk bodies and hyaline granules and is found in most precancerous lesions of the liver and in HCC.<sup>[60]</sup> *NRF2* influences the cellular stress program related to the occurrence and progression of cancer,<sup>[61]</sup> and participates in the regulation of proliferation of HCC and liver cancer stem cells.<sup>[62]</sup> Mutations in the *NRF2* gene were found in approximately 15% of HCC cases;<sup>[63]</sup> *NRF2* can act as both a tumor suppressor and an oncogene,<sup>[64]</sup> and the *NRF2*-mediated antioxidant response pathway is considered to be a “double-edged sword” with respect to the occurrence and development of cancer. Second, in the regulatory network of ferroptosis, the p62-Keap1-NRF2 pathway regulates lipid peroxidation, oxidative stress, and iron metabolism in HCC cells by transcribing downstream

**Table 1: Regulatory factors of ferroptosis.**

Regulatory factors	Possible pathways	References
NRF2, <i>p53</i> , BAP1, BECN1, erastin, RSL3	System $x_c^-$ -GPX4	[14,18,20-22]
LOXs, ACSL4, NRF2	Lipid peroxidation	[30-32,65]
IRE-BP2, TF, TFR, NCOA4, HO1, PROM2, NRF2	Iron metabolism	[8,23,41,43-46,65]
FSP1	FSP1-CoQ10-NAD(P)H	[48,49]

ACSL4: Long-chain fatty acid-coenzyme A (CoA) ligase 4; BAP1: Breast cancer susceptibility gene 1 (BRCA1) associated protein 1; BECN1: Beclin 1; CoQ10: Coenzyme Q10; FSP1: Ferroptosis suppressor protein 1; GPX4: glutathione peroxidase 4; HO1: Heme oxygenase 1; IRE-BP2: Iron responsive element binding protein 2; LOXs: Lipoxygenases; NAD(P)H: Nicotinamide adenine dinucleotide; NCOA4: Nuclear receptor coactivator 4; NQO1: Quinone oxidoreductase 1; NRF2: Nuclear factor erythroid 2-related factor; PROM2: Prominin-2; RSL3: *RAS*-selective lethal 3; TF: Transferrin; TFR: Transferrin receptor.

target genes such as quinone oxidoreductase 1 (*NQO1*), *HO1*, and ferritin heavy chain 1 (*FTH1*), thus inhibiting ferroptosis.<sup>[65]</sup> In addition, studies have shown that many steps of the ferroptosis cascade are centered on NRF2.<sup>[66–68]</sup> Sun *et al*<sup>[65]</sup> showed that the inhibition of the p62–Keap1–NRF2 pathway could significantly enhance the anticancer activity of erastin and sorafenib in HCC cells both *in vivo* and *in vitro* and finally proved that the p62–Keap1–NRF2 pathway was a key negative regulator of ferroptosis in HCC cells: p62 can competitively bind to Keap1, thus increasing the transcriptional activity and stability of NRF2.<sup>[69]</sup> Umemura *et al*<sup>[60]</sup> showed that the accumulation of p62 is an important prerequisite for the malignant development of precancerous liver lesions, which may occur via the prevention of HCC-initiating cell death induced by oxidative stress. This way of preventing HCC-initiating cell death may be related to ferroptosis inhibition because the accumulation of p62 eventually leads to the non-mutagenic activation of three key cancer drivers including NRF2; however, there is no direct evidence yet. On the other hand, the interaction between autophagy and ferroptosis is still unclear.<sup>[70]</sup> It has been reported that when autophagy is defective, NRF2 can be activated through the atypical Keap1–NRF2 pathway because p62 autophagy degradation is inhibited.<sup>[71]</sup> In the occurrence of this series of events, autophagy and ferroptosis seem to be located in upstream and downstream of the p62–Keap1–NRF2 pathway, respectively, and interact with each other on the axis of this pathway. Interestingly, a study reported that NRF2 was the common intermediary of hepatomegaly.<sup>[72]</sup> We hold the opinion that it is essential for the study of ferroptosis to establish biomarkers that can characterize the occurrence of ferroptosis, which will be of great contribution to our exploration of the relationship between ferroptosis and cancer.

Glutathione S-transferase zeta 1 (*GSTZ1*) belongs to the glutathione S-transferase (*GST*) superfamily, which is involved in the catabolism of phenylalanine/tyrosine and catalyzes the isomerization of maleylacetoacetate to fumarylacetoacetate.<sup>[73]</sup> It has been reported that *GSTZ1* shows low expression in HCC, which leads to the accumulation of the metabolite succinylacetone in cells, and activation of the NRF2 pathway, finally promoting the proliferation of HCC cells.<sup>[74]</sup> Wang *et al*<sup>[73]</sup> showed that *GSTZ1* was strikingly downregulated in sorafenib-resistant HCC, and the deletion of *GSTZ1* promoted the activation of the NRF2 pathway, increased the level of *GPX4*, and decreased iron levels and lipid peroxidation, which finally inhibited ferroptosis induced by sorafenib.

Metallothionein-1G (*MT-1G*) belongs to the metallothionein (*MT*) family, which includes low molecular weight and cysteine-rich proteins, and it exists in mammals as an antioxidant and heavy metal antidote.<sup>[75,76]</sup> It has been reported that *MT-1G* can inhibit carcinogenesis,<sup>[75]</sup> inhibit metastasis and promote differentiation.<sup>[77,78]</sup> Sun *et al*<sup>[79]</sup> found that sorafenib could induce the expression of *MT-1G* in HCC cells through the activation of NRF2, but not *p53* or hypoxia-inducible factor (*HIF*)-1 $\alpha$ , while the inhibition of *MT-1G* could increase the sensitivity to sorafenib therapy, which finally proved that *MT-1G* was a negative regulator of ferroptosis in HCC. Their research

also found that inhibiting *MT-1G* could enhance the anticancer activity of sorafenib *in vitro* and in tumor xenograft models.

The *Rb* tumor suppressor gene is a key regulatory factor in the cell cycle that is activated in response to numerous anti-mitotic signals and plays a vital role in the G1/S checkpoint.<sup>[80]</sup> The loss of the *Rb* function is a key event in the process of HCC.<sup>[81–83]</sup> A study showed that the cell death level of HCC cells with low *Rb* expression induced by sorafenib was 2–3 times higher than that of the control group and resulted in complete tumor regression in 50% of sorafenib-treated mice that received HCC cell xenografts with low *Rb* expression, but the tumors in the control group were stable, and the difference was statistically significant.<sup>[84]</sup> This study finally proved that the low expression of *Rb* can effectively promote the induction of HCC cell ferroptosis by sorafenib.

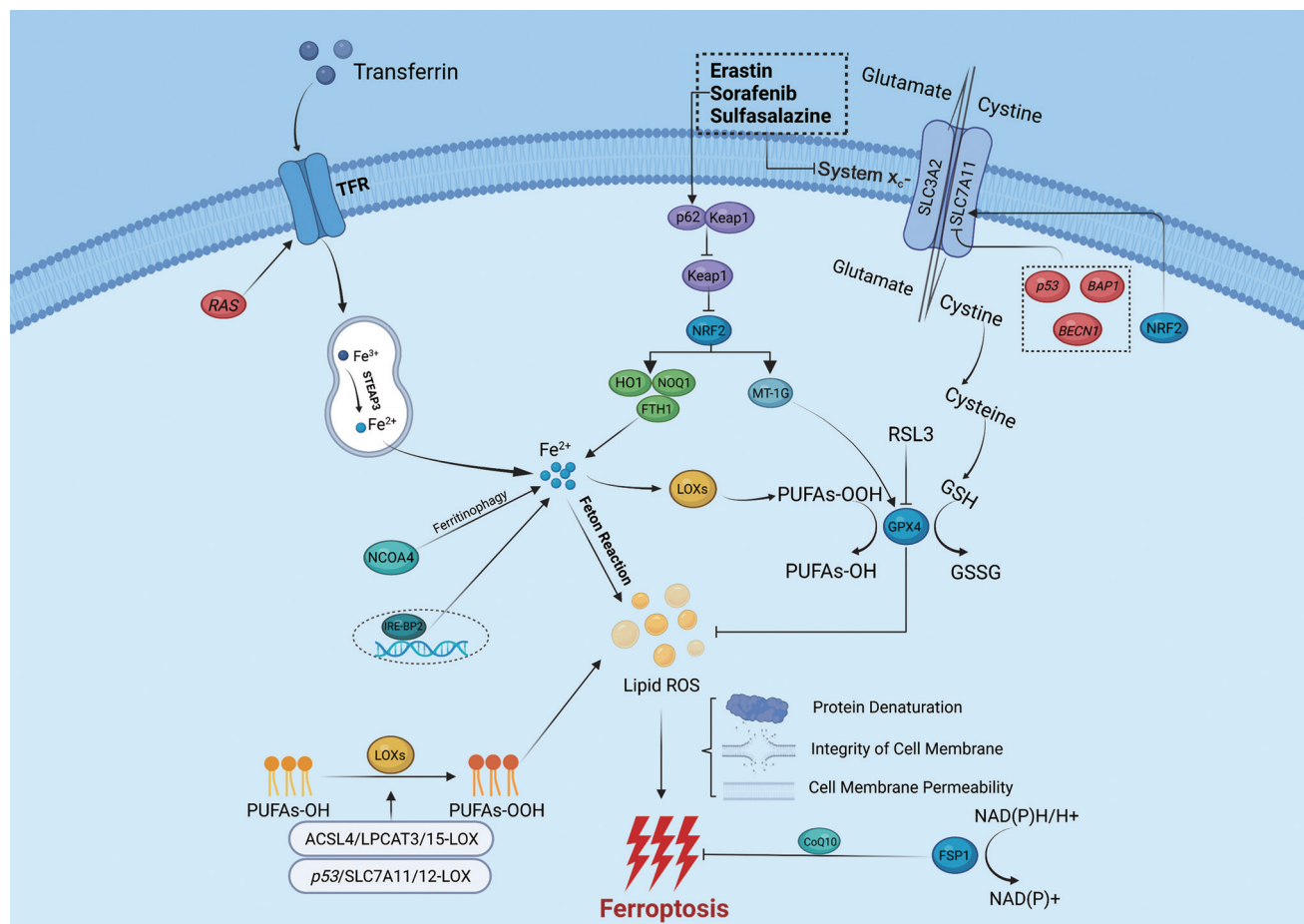
In addition to the regulatory factors mentioned above that affect ferroptosis through participation in iron metabolism,<sup>[41–46]</sup> recent progress has been made in the identification of regulatory factors that affect ferroptosis by regulating the iron homeostasis of HCC. O-linked N-acetylglucosaminylation (O-GlcNAcylation) is a reversible post-translational modification catalyzed by O-linked N-acetylglucosamine (O-GlcNAc).<sup>[85]</sup> There is evidence that yes-associated protein (*YAP*) O-GlcNAcylation can enhance the sensitivity of HCC to ferroptosis by regulating the expression of *TFR*, which may be a promising treatment for HCC.<sup>[86]</sup> Ceruloplasmin (*CP*), a copper-containing protein, belongs to the multicopper oxidase family and can prevent the production of ROS mediated by ferrous ions and is widely regarded as an effective protective agent against antioxidant stress.<sup>[87,88]</sup> A study has shown that deletion of *CP* can promote the ferroptosis of HCC cells caused by erastin and *RSL3* and can lead to the accumulation of intracellular ferrous ions and lipid ROS, which ultimately proves that *CP* can suppress ferroptosis by affecting iron homeostasis in HCC.<sup>[89]</sup>

Similarly, the effect of lipid metabolism on ferroptosis is mainly regulated by *ACSL4*.<sup>[32,90]</sup> Doll *et al*<sup>[32]</sup> showed through oxi-lipidomics analysis that *ACSL4* can influence the composition of lipids required for ferroptosis. The main role of *ACSL4* in ferroptosis is to ligate coenzyme A to long-chain PUFAs; once inside the membranes, these PUFAs may undergo peroxidation, which is the essential element for ferroptosis.<sup>[15,38]</sup> In HCC, *ACSL4* can restore the lipid metabolism of HCC cells through the c-Myc/sterol regulatory element-binding protein 1 (*SREBP1*) pathway and then participate in the proliferation and metastasis of HCC.<sup>[91]</sup> In addition, in recent years, a number of studies have shown that *ACSL4* can adequately predict the prognosis of sorafenib-treated HCC patients, especially those who exhibit sorafenib-mediated ferroptosis.<sup>[92,93]</sup> The molecular mechanisms of ferroptosis and key regulatory pathways are shown in [Figure 1] (created with BioRender.com).

### Application of Ferroptosis in HCC

Sorafenib is the first multikinase inhibitor approved for unresectable HCC. It is worth noting that several studies





**Figure 1:** Molecular mechanisms of ferroptosis and key regulatory pathways. Ferroptosis is directly triggered by iron-dependent lipid peroxidation, through three possible mechanisms: (1) denaturation of related proteins on the cellular membrane; (2) disruption of the integrity of the cellular membrane; and (3) increased permeability of the cellular membrane. Lipid alcohols are oxidized to lipid hydroperoxides in a reaction mediated by LOXs, and eventually participate in lipid peroxidation. Lipid peroxidation requires the Fenton reaction and the overreaction of LOXs; simultaneously, GPX4 can prevent lipid peroxidation by consuming two molecules of GSH. GSH enters the cell through a process mediated by system  $x_c^-$ . Iron is transported to TFR by transferrin, then imported into the cell by TFR, and eventually converted to  $Fe^{2+}$  in endosomes. Subsequently,  $Fe^{2+}$  participates in the Fenton reaction and the synthesis of lipid peroxidation-related enzymes such as LOXs. ACSL4: Long-chain fatty acid-coenzyme A (CoA) ligase 4; BAP1: Breast cancer susceptibility gene 1 (BRCA1) associated protein 1; BECN1: Beclin 1; CoQ10: coenzyme Q10; FSP1: Ferroptosis suppressor protein 1; FTH1: Ferritin heavy chain 1; GPX4: Glutathione peroxidase 4; GSH: Glutathione; GSSG: Oxidized glutathione; HO1: Heme oxygenase-1; IRE-BP2: Iron responsive element binding protein 2; Keap1: Kelch-like ECH-associated protein 1; LOXs: Lipoxygenases; LPCAT3: Lysophospholipid-acyltransferase 3; MT-1G: Metallothionein-1G; NAD(P)H: Nicotinamide adenine dinucleotide; NCOA4: Nuclear receptor coactivator 4; NQO1: Quinone oxidoreductase 1; NRF2: Nuclear factor erythroid 2-related factor; PUFAs: Polyunsaturated fatty acids; ROS: Reactive oxygen species; SLC3A2: Solute carrier family 3 member 2; SLC7A11: Solute carrier family 7 member 11; STEAP3: Six transmembrane epithelial antigen of prostate 3; TFR: Transferrin receptor.

have shown that the mechanism of ferroptosis induced by sorafenib is mediated by non-kinase inhibition.<sup>[65,94]</sup> On the one hand, sorafenib can inhibit the activity of the system  $x_c^-$  receptor and cause GSH consumption, which eventually leads to ferroptosis.<sup>[84]</sup> On the other hand, a study showed that the responsive expression of NRF2 increased in HCC cells after treatment with sorafenib, which may be the mechanism of sorafenib resistance in some HCC patients,<sup>[65]</sup> and the latest research shows that sorafenib may be involved in the regulation of protein phosphorylation signals in the key regulatory pathways as a means of regulating ferroptosis.<sup>[95]</sup> Recently, research has focused on enhancing the effect of sorafenib on ferroptosis in various ways. Artesunate (ART) itself can induce ROS-mediated cell death by inducing ferritinophagy and disrupting the unstable iron pool in HCC,<sup>[96,97]</sup> while its combination with sorafenib can produce a synergistic effect to induce ferroptosis.<sup>[98]</sup> Quiescin sulphydryl oxidase 1 (QSOX1) can promote ferroptosis

induced by sorafenib in HCC by driving the nuclear endosome transport of the epidermal growth factor receptor (EGFR) and restraining the activity of the NRF2.<sup>[99]</sup> In addition, changes in GSTZ1 and MT-1G expression,<sup>[73,79]</sup> as mentioned above, can synergize with sorafenib to improve the induction of ferroptosis in tumor cells. However, although a study has shown that ACSL4 can be used as a biomarker for predicting the sensitivity of sorafenib in HCC,<sup>[92]</sup> we still do not understand the relationship between the kinase inhibition and ferroptosis effects of sorafenib and their dominance in antitumor activity.

However, the bioavailability of sorafenib is limited to a certain extent because of its water solubility and targeting properties. In one study, a novel cascaded copper-based metal-organic framework (MOF) nanocatalyst was constructed using the Hong Kong University of Science and Technology (HKUST) -1 framework by incorporating the cyclooxygenase-2 (COX-2) inhibitor meloxicam (Mel)

and sorafenib to improve the treatment effect in HCC.<sup>[100]</sup> This method can substantially improve the water solubility of sorafenib and effectively regulate the levels of GSH and COX-2 in HCC cells, which provides a new therapeutic strategy. A study constructed sorafenib-loaded MIL-101 (Fe) nanoparticles and combined them with an iRGD peptide containing a tumor-homing motif (RGD) and a tissue-penetrating motif (CendR), which may represent a promising treatment for HCC by inducing ferroptosis.<sup>[101]</sup> Tang *et al*<sup>[102]</sup> introduced manganese-doped mesoporous silica nanoparticles (MMSNs), which could induce the ferroptosis of tumor cells through the consumption of intracellular GSH caused by the degradation of MMSNs. They loaded MMSNs with sorafenib at a drug loading rate of  $2.68 \pm 0.32\%$  and released the drug into the tumor microenvironment (TME) to achieve GSH inhibition. On the other hand, MMSNs themselves can also consume intracellular GSH, thus achieving synergism. In recent years, the therapeutic strategy of ferroptosis caused by the strong permeability, retention, and targeting of nanomaterials in cancer has been widely studied. Nanotechnology not only enhances the targeting characteristics of tumors but also prolongs the life cycle of ferroptosis inducers,<sup>[103]</sup> and is thus a very promising field.

In addition to studies on erastin, sorafenib, and RSL3, there have been an increasing number of studies on drugs that induce ferroptosis. It has been reported that disulfiram (DSF) can react with Cu to generate the anticancer metabolite DSF/Cu and inhibit the activity of the p97-dependent proteasome.<sup>[104]</sup> Ren *et al*<sup>[105]</sup> showed that DSF/Cu can seriously disrupt mitochondrial homeostasis, increase the free iron pool, and enhance lipid peroxidation, eventually leading to ferroptosis, and can cooperate with sorafenib to inhibit the NRF2 and mitogen-activated protein kinase (MAPK) pathways. DSF/Cu can also significantly increase the cytotoxicity of sorafenib while inducing ferroptosis in HCC. The nuclear protein 1 (NUPR1) inhibitor ZZW-115 can induce mitochondrial dysfunction in pancreatic cancer and HCC cells, which leads to ferroptosis after the accumulation of ROS. This phenomenon can be inhibited by the Transcription Factor A, Mitochondrial (TFAM).<sup>[106]</sup> Furthermore, solasonine has been reported to promote ferroptosis in HCC cells by suppressing GPX4 and glutathione synthetase (GSS).<sup>[107]</sup> Haloperidol can adequately increase GSH levels, ferrous ion levels and lipid peroxidation in cells, thus promoting ferroptosis induced by erastin and sorafenib.<sup>[108]</sup> Loss of leukemia inhibitory factor (LIF) receptor subunit alpha (LIFR), the gene that encodes the LIF receptor, can induce liver tumorigenesis and resistance to drug-related ferroptosis through upregulation of the iron-sequestering cytokine lipocalin 2 (LCN2) mediated by nuclear factor kappa-B (NF- $\kappa$ B), and this situation can be altered by an LCN2-neutralizing antibody.<sup>[109]</sup> Yao *et al*<sup>[109]</sup> found that this kind of antibody was able to elevate the ferroptosis-inducing and anticancer effects mediated by sorafenib, and they finally proved that co-administration with sorafenib and LCN2-neutralizing antibody had a better therapeutic response in mice bearing HCC patient-derived xenograft tumors.

Immunotherapy has been a great breakthrough in the field of cancer in recent years and has a promising therapeutic

effect on malignant tumors, including HCC, and recent studies have begun to explore the role of ferroptosis in cancer immunotherapy and TME. Xu *et al*<sup>[110]</sup> have indicated that ferroptosis plays a two-way role in the TME. Specifically, ferroptosis may restrain the activity of antitumor immune cells, including cluster of differentiation 8 (CD8)<sup>+</sup> T cells, natural killer (NK) cells, and dendritic cells (DCs),<sup>[111-113]</sup> leading to a decrease in the antitumor immunity. For example, Ma *et al*<sup>[114]</sup> found that CD36 can mediate the uptake of fatty acids in the TME by tumor-infiltrating CD8<sup>+</sup> T cells, induce ferroptosis, reduce the production of cytotoxic cytokines, and weaken the antitumor ability, while blocking CD36 can effectively restore the antitumor activity of CD8<sup>+</sup> T cells, and the combination with anti-programmed death 1 (PD1) antibodies possessed stronger antitumor efficacy. At the same time, some immunosuppressive immune cells, including M2 tumor-associated macrophages (TAMs), and T regulatory cells (Tregs), are also affected by ferroptosis, which leads to the reversal of immunosuppression.<sup>[115,116]</sup> A recent study showed that the higher resistance of M1 TAMs to ferroptosis mediated by GPX4 deletion can be used to inhibit the survival of M2 TAMs, and eventually enhance antitumor immunity.<sup>[115,117]</sup> On the other hand, the effect of ferroptosis on cancer cells also changes the TME, and ferroptotic cancer cells can release immunostimulant signals such as damage-associated molecular pattern (DAMP) signals, so that immune cells can locate cancer cells accurately.<sup>[118,119]</sup> In addition, ferroptotic cancer cells can induce greater tumor-specific immune responses to enhance the efficacy of immunotherapy.<sup>[120]</sup> However, ferroptotic cancer cells can also suppress anti-tumor immunity through various mechanisms. Studies have shown that ferroptotic cancer cells are associated with the release of prostaglandin E2 (PGE2), which is a key immunosuppressive factor and capable of inhibiting the activity of antitumor immune cells.<sup>[14,121]</sup> In short, in view of the complex crosstalk between the TME and ferroptosis, the relationship between ferroptosis and immunotherapy is more than just a synergistic one; it should also be determined according to different immunophenotypes, different induction mechanisms of ferroptosis, and different types of immune cells. At present, the combination of ferroptosis and immunotherapy is relatively rare in HCC. Clinical or preclinical application of ferroptosis in hepatocellular carcinoma are summarized in Table 2.

## Conclusion and Perspective

Ferroptosis is a kind of RCD that involves triggering, execution, and regulation processes; the mechanisms underlying these processes are complex but are mainly mediated by three central axes: system  $x_c^-$ /GPX4, iron metabolism and lipid peroxidation. Nevertheless, the specific mechanism underlying the induction of ferroptosis, such as the mechanism of cell death after triggering ferroptosis and the contributions of these processes, is unclear; these areas need to be further explored to facilitate the clinical translation and application of ferroptosis-related strategies. Moreover, we know little about the relationship between ferroptosis and other types of RCD. Regarding the relationship between ferroptosis and cancer,

**Table 2: Clinical or preclinical application of ferroptosis in hepatocellular carcinoma (HCC).**

Treatment	Target	Possible mechanism	Reference
DHA	LOXs	By promoting the binding of PEBP1 and 15-LO, and activating LOXs allosterically to induce lipid peroxidation	[30]
An LCN2-neutralizing antibody	LCN2	By targeting the upregulation of LCN2 mediated by loss of LIFR and upregulating Fe <sup>2+</sup> and lipid peroxidation levels	[109]
Sorafenib	System x <sub>c</sub> <sup>-</sup>	By inhibiting system x <sub>c</sub> <sup>-</sup> for absorption of cystine, and promoting the depletion of GSH	[123]
ART	Ferritin	By inducing ferritinophagy and releasing iron into the cytosol to produce ROS, which can also synergize with sorafenib	[97,98]
QSOX1	NRF2	By promoting the nuclear endosome transport of EGFR and restraining the activity of NRF2	[99]
A cascaded copper-based nanocatalyst	GSH	By improving the water solubility of sorafenib and increasing the levels of GSH and COX-2	[100]
MMSNs loaded with sorafenib	GSH	By promoting the depletion of GSH and increasing the targeting characteristics of sorafenib, achieving synergism	[102]
Co-administration of MIL-101(Fe)@sor NPs with iRGD	GSH; GPX4	By reducing the levels of GSH and activity of GPX4	[101]
RSL3@O2-ICG NBs	GSH	By promoting the depletion of GSH and damaging mitochondria	[122]
DSF/Cu	Iron; NRF2	By disrupting mitochondrial homeostasis, increasing the iron pool and inhibiting NRF2 pathway for synergizing with sorafenib	[105]
Solasonine	GPX4	By restraining the activity of GPX4 and GSS	[107]

ART: Artesunate; COX-2: Cyclooxygenase-2; DHA: Dihydroartemisinin; DSF: Disulfiram; EGFR: Epidermal growth factor receptor; GPX4: Glutathione peroxidase 4; GSH: Glutathione; GSS: Glutathione synthetase; ICG: Indocyanine green; LCN2: Lipocalin 2; LIFR: Leukemia inhibitory factor (LIF) receptor subunit alpha; LOXs: Lipoxygenases; MMSNs: Manganese-doped mesoporous silica nanoparticles; NBs: Nanobubbles; NPs: Nanoparticles; NRF2: Nuclear factor erythroid 2-related factor; PEBP1: Phosphatidylethanolamine binding protein 1; QSOX1: Quiescin sulfhydryl oxidase 1; RGD: Tumor-homing motif; ROS: Reactive oxygen species.

as far as the current situation is concerned, its role in the occurrence and development of tumors seems to be two-sided, and we need to explore and explain the effects of the related genes and factors on ferroptosis as much as possible, which is a crucial but difficult area. Finally, we described the influence, targets, and clinical applications of ferroptosis in HCC. We have taken a small step toward the application of ferroptosis in HCC. As mentioned previously, ferroptosis is a novel opportunity for treatment in the era of apoptosis resistance, and we should pay more attention to the identification of biomarkers of ferroptosis, which is one of the most important prerequisites for improving ferroptosis assessment and clinical application.

### Conflicts of interest

None.

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**How to cite this article:** Cong T, Luo Y, Fu Y, Liu Y, Li Y, Li X. New perspectives on ferroptosis and its role in hepatocellular carcinoma. *Chin Med J* 2022;135:2157–2166. doi: 10.1097/CM9.00000000000002327