

Iron metabolism and drug resistance in cancer

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Abstract Iron is an essential inorganic element for various cellular events. It is directly associated with cell proliferation and growth; therefore, it is expected that iron metabolism is altered in tumor cells which usually have rapid growth rates. The studies on iron metabolism of tumor cells have shown that tumor cells necessitated higher concentrations of iron and the genes of iron uptake proteins were highly overexpressed. However, there are limited number of studies on overall iron metabolism in drug-resistant tumor cells. In this article, we evaluated the studies reporting the relationship between drug resistance and iron metabolism and the utilization of this knowledge for the reversal of drug resistance. Also, the studies on iron-related cell death mechanism, ferroptosis, and its relation to drug resistance were reviewed. We focus on the importance of iron metabolism in drug-resistant cancer cells and how alterations in iron metabolism participate in drug-resistant phenotype.

Keywords Iron metabolism · Chemotherapy · Drug resistance · Ferroptosis

Introduction

Iron is an inorganic element which is critical for cell proliferation and growth by incorporating into iron- or heme-containing enzymes. Those enzymes are involved in respiratory complexes, DNA synthesis, cell cycle and detoxification processes. Hence, iron is essential in terms of cell replication, cellular metabolism and growth. Beside these effects, iron can create reactive oxygen species (ROS) by participating in Fenton reaction where hydroxyl radical is produced. ROS can damage DNA and be mutagenic. Thus, iron is not only fundamental for cell survival but it can be related to carcinogenesis (Torti and Torti 2013a, b).

Several studies in the literature relating iron metabolism and cancer revealed that iron is highly demanded by cancer cells. It can facilitate the formation and metastasis of tumor cells, and this detrimental effect can be overcome by iron chelators (Richardson et al. 2009; Torti and Torti 2011, 2013a, b). However, the role of iron metabolism in resistance to chemotherapeutic agents in cancer has not been fully understood.

Drug resistance describes different mechanisms of tumor cells to avoid the cytotoxic effects of anticancer agents with little to no common similarity in terms of structure, function and therapeutic target. Drug-resistant phenotype shows itself with increased insensitivity to chemotherapeutics, leading to decreased success of cancer treatment (Housman et al. 2014). Longley and Johnston stated more than 90% of cancer patients

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with metastatic tumors are believed to experience failure in chemotherapy due to multidrug resistance (Longley and Johnston 2005). The greater comprehension of cellular mechanisms that drive cancer cells to become drug-resistant will remarkably increase the clinical success of chemotherapy.

Drug resistance, either intrinsic or acquired, develops through alterations in various cellular mechanisms. The cumulative effect of these alterations provides drug-resistant cells with a survival advantage over drug-sensitive ones, leading to formation of drugresistant tumors. Several studies demonstrated that when over-expressed, ATP-binding cassette (ABC) transporter family proteins pump a wide range of anticancer drugs out of the cell, preventing intracellular drug concentration to reach cytotoxic levels. Apart from the ATP-driven efflux, structural changes in plasma membrane lower membrane permeability; thus, decrease drug influx. Increased activity of cellular detoxification, altered drug targets, enhanced DNA repair, alkaline shift of cellular pH, evasion of apoptosis and increased pro-survival signaling are of importance in development of drug-resistant phenotype (Krishna and Mayer 2000; Lage 2003; Leonessa and Clarke 2003; Longley and Johnston 2005; Simon and Schindler 1994).

In the present review, we evaluated the studies that combine iron metabolism with drug resistance in cancer. We also focused on the strategies to utilize iron metabolism in order to overcome drug resistance. Finally, we compiled the ferroptosis studies associating with the drug resistance in cancer.

Iron metabolism and drug resistance in cancer cells

Cellular iron metabolism comprises of three major processes: iron uptake, storage and export (Fig. 1). Firstly, to uptake iron inside the cell, transferrin (Tf) binds free iron (Fe(III)) in circulation and Tf-Fe conjugate then binds to transferrin receptor (TfR) on cell membrane. Tf-Fe-TfR conjugate is internalized via clathrin-dependent endocytosis. Cytoplasmic iron is stored by ferritin heteropolymers which consist of heavy (FtH) and light (FtL) chains. Finally, excess iron is exported by ferroportin (SLC40A1; Hentze et al. 2004).

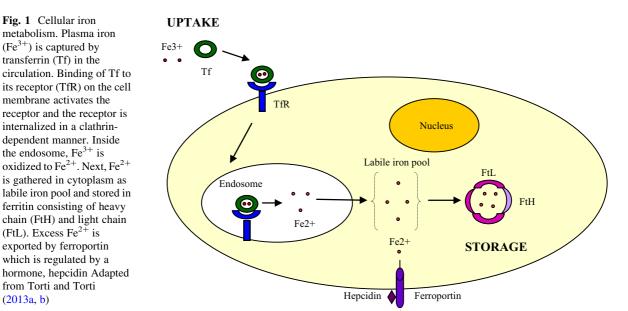
In this section, we will basically focus on how expression patterns of these proteins participating in iron uptake, storage and export as well as intracellular free iron levels change in drug-resistant cancer cells compared to their drug-sensitive counterparts. We summarized the altered expression levels of iron metabolism proteins in drug-resistant cell lines in Table 1 and the differences between expression levels of these proteins are given in Fig. 2.

Transferrin receptor and drug resistance

Transferrin receptor (TfR) is a membrane glycoprotein which has a role in importing iron via conjugation to a plasma glycoprotein, transferrin. TfR is a homodimer with 180 kDa molecular weight. Each monomer contains a C-terminal domain that binds transferrin, a single-pass transmembrane domain and a short N-terminal domain. There are four glycosylation sites in C-terminal and this modification is crucial for activity of the receptor. The transcription of TfR gene is regulated by intracellular iron concentration via binding of iron regulatory proteins (IRPs) to the iron response elements (IREs) on the 5' untranslated region of TfR transcript (Ponka and Lok 1999; Daniels et al. 2006).

Barabas and Faulk have shown that TfR expression was up-regulated in doxorubicin-resistant human chronic myelogenous leukemia cells (K562) and promyclocytic leukemia cells (HL60). Verapamil, a well-known P-glycoprotein inhibitor and a calcium channel blocker, decreased IC50 value of doxorubicin in drug-resistant HL60 cells, which did not overexpress P-glycoprotein, in comparison to drug-sensitive cells, by down regulating TfR. TfR can be upregulated by growth hormones, such as insulin-like growth factor which decreases the rate of receptor endocytosis and epidermal growth factor that increases rate of receptor exocytosis (Davies et al. 1987). This induction was shown to be triggered by calcium channels (Neckers 1991) and the blockade of such mechanism could re-sensitize drug-resistant cells (Cornwell et al. 1987). This may explain the association of TfR with drug resistance through the function of calcium-sensitive receptors (Barabas and Faulk 1993).

A proteomic study to identify membrane protein differences between drug-sensitive and mitoxantrone (MXR)-resistant human breast adenocarcinoma MCF7 cell lines revealed that TfR was highly expressed in MXR-resistant cells while not detected



in drug-sensitive parents (Rahbar and Fenselau 2005). Similarly, acquired tamoxifen- and Faslodex-resistant MCF7 cells displayed elevated TfR expression at both mRNA and protein levels (Habashy et al. 2010).

Gallium is a chemical element used as a chemotherapeutic agent, particularly in bladder cancer and lymphoma treatment (Chitambar 2004). Gallium resistance occurs through a unique pathway involving iron metabolism. Because of the fact that gallium has a similar chemistry in terms of ionic radii and bonding to iron, gallium remarkably interferes with iron metabolism (Davies et al. 2006); it can easily conjugate to transferrin and be internalized by TfR-mediated endocytosis. Through this competitive binding to transferrin, gallium inhibits cellular iron uptake and incorporation, and blocks DNA synthesis (Chitambar and Seligman 1986; Chitambar and Zivkovic 1987; Harris and Pecoraro 1983; Larson et al. 1980). In gallium-resistant HL60 cells, TfR was found to be up-regulated, although surprisingly gallium uptake was not increased by high TfR expression compared to drug-sensitive cells. Gallium treatment caused arrest in cellular proliferation in drug-susceptible parents by inhibiting iron uptake whereas drug-resistant HL60 cells continued normal proliferation. Gallium-resistant cells had rapidly uptaken exogenous iron as a result of high levels of TfR on plasma membrane; thus, drug-resistant HL60 cells overcame the effect of gallium by enhanced incorporation of iron (Chitambar et al. 1990). In contrast, Davies et al. reported TfR expression was downregulated in gallium-resistant HL60 cells and as a result, iron uptake was reduced (Davies et al. 2006). Similarly, in gallium-resistant human leukemic CCRF-CEM cells, TfR expression, and subsequently the iron uptake, was found to be decreased (Chitambar and Wereley 1997).

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Chekhun et al. used doxorubicin- and cisplatinresistant MCF7 cells to investigate differences in expressions of iron metabolism-related proteins. Drug-resistant cells exhibited higher TfR1 expression and iron uptake than drug-sensitive parents. However, intracellular unincorporated Fe(III) levels did not follow the same pattern; in cisplatin-resistant MCF7 cells, free iron levels were 2 times higher than those of drug-sensitive cells whereas there was twofold decrease in doxorubicin-resistant cells (Chekhun et al. 2013, 2017).

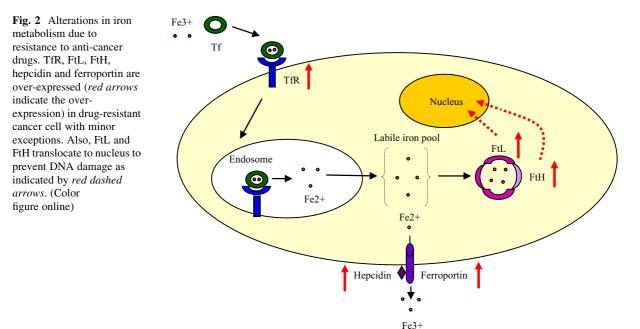
Protein	Cell line	Drug	Expression	Reference
Transferrin receptor	MCF7	Doxorubicin	Up-regulated	Chekhun et al. (2013)
		Cisplatin		
		Mitoxanrone		Rahbar and Fenselau (2005)
		Tamoxifen		Habashy et al. (2010)
		Faslodex		
	K562	Doxorubicin		Barabas and Faulk (1993)
	HL60			
	U266	Bortezomib		Campanella et al. (et al. 2013)
	RPMI-8226			
	H69VP	Artemisinin		Sadava et al. (2002)
	HL60	Gallium		Chitambar et al. (1990)
			Down-regulated	Davies et al. (2006)
Ferritin Light Chain	MCF7	Doxorubicin	Up-regulated	Chekhun et al. (2013)
		Cisplatin		
Ferritin Heavy Chain	MCF7	Doxorubicin	Down-regulated	Chekhun et al. (2013)
		Cisplatin	Up-regulated	
	MEL	Colchicine		Epsztejn et al. (1999)
		Vinblastine		
	SNU	Cisplatin		Kang et al. (2004)
	CCRF-CEM	Gallium		Chitambar and Werely (1997)
	U266	Bortezomib		Campanella et al. (et al. 2013)
	RPMI-8226			
Total Ferritin Content	HL60	Gallium	Down-regulated	Chitambar et al. (1990)
	CCRF-CEM			Chitambar and Werely (1997)
Ferroportin	MCF7	Doxorubicin	Up-regulated	Chekhun et al. (2013)
		Cisplatin		
Hepcidin	MCF7	Doxorubicin	Up-regulated	Yalovenko et al. (2016)
		Cisplatin		

Table 1 Expression levels of iron metabolism-related proteins in drug-resistant cell lines compared to their drug-sensitive counterparts

Ferritin and drug resistance

Ferritin is an intracellular protein which is responsible for iron sequestration, storage and release. It is composed of two subunits; heavy chain and light chain. These two chains self-assemble and form hetero-24mers in which the ratio of chains is tissuedependent. Ferritin light chain (FtL) is responsible for mineral nucleation due to its greater negative charge. Ferritin heavy chain (FtH) is involved in nucleation process and also displays ferroxidase activity (Broxmeyer et al. 1991; Koorts and Viljoen 2007; Uchida et al. 2010). The transfection of *FtH* has been shown to trigger acquired multidrug resistance (MDR) in murine erythroid leukemia (MEL) cells. Epsztejn et al. revealed expression levels of MDR1 and FtH at both mRNA and protein levels were directly proportional. FtH-overexpressing cells were three-to-seven fold resistant to hydrophobic cytotoxic drugs, colchicine and vinblastine, and remarkably affected by P-glycoprotein blockers and MDR reversal agents, cyclosporine and verapamil. The fact that MDR blockers significantly reduced the cellular growth in FtH-overexpressing cells indicates the direct correlation between FtH and multidrug resistance in those cells (Epsztejn et al. 1999).

In gallium-resistant CCRF-CEM cells, heavy chain ferritin mRNA levels were shown to be increased; nevertheless, total ferritin content was lower. The



excess transferrin treatment sensitized resistant cells to gallium, implying that gallium resistance is closely linked to iron metabolism (Chitambar and Wereley 1997).

Checkhun et al. proposed the expression of FtH as cell-specific; up-regulated in cisplatin-resistant MCF7 cells but down-regulated in doxorubicin-resistant ones. On the contrary to FtH, FtL was found to be up-regulated in both drug-resistant MCF7 sub-lines; however, sub-cellular localization of FtL differed between drug-sensitive and -resistant cells. As reviewed by Alkhateeb and Connor, nuclear ferritin protects DNA from DNA damage-inducing agents (Alkhateeb and Connor 2010). FtL was cytoplasmic in drug-sensitive parents whilst it was mostly nuclear in drug-resistant counterparts (Chekhun et al. 2013).

FtH expression was also shown to be over-expressed in cisplatin-resistant gastric cancer cells, SNU-620R-CIS/2000, SNU-638R-CIS/400 and SNU-668RCIS/ 400 in a microarray-based study (Kang et al. 2004). Liu et al. showed that 1,3-bis(2-chloroethyl)-1-nitrosurea (BCNU), which is a DNA-alkaylating agent used in brain tumor therapy (Madajewicz et al. 1981), caused supercoiling of plasmid DNA (pUC19); however, exogenous addition of FtH reversed supercoiling and re-linearized DNA. The change in DNA structure suggests that FtH protected DNA from any damage caused by chemotherapeutics and the presence of FtH in nucleus resulted in decreased sensitivity against anti-cancer agents (Liu et al. 2011).

Bortezomib is a 26S proteasome inhibitor that has an effect on the regulation of various signaling pathways (Chauhan et al. 2005). Campanella et al. reported that bortezomib resistance in multiple myeloma cell lines was correlated with FtH levels. Even though iron supplementation triggered generation of ROS in drug-sensitive cells (MM.1S and KMS-18), such effect was not observed in resistant ones (U266 and RPMI-8226). Down-regulation of FtH in U266 and RPMI-8226 cell lines successfully sensitized the cells to bortezomib (Campanella et al. 2013).

FtH and *FtL* mRNAs contain iron response elements in their 5' UTR. These regions were shown to form hairpin loops and impact the regulation of protein synthesis by interacting with other proteins and small molecules (Hentze et al. 2004; Oliveira et al. 1993). The anthracyclines, doxorubicin and daunorubicin, were found to interact with 5' UTRs of *FtH* and *FtL* mRNA (Canzoneri and Oyelere 2008).

Hereditary hemochromatosis protein and drug resistance

Hereditary hemochromatosis protein (HFE) is an atypical major histocompatibility class I molecule that acts as an iron sensor by interacting with TfR1 and

TfR2 and regulates cellular iron uptake and whole body iron homeostasis (Goswami and Andrews 2006). Davies et al. revealed a novel mechanism in which HFE formed a secondary complex with transferrin and TfR in gallium-resistant HL60 cells; however, such a complex was absent in drug-sensitive parents. Since HFE competes with transferrin to bind to TfR, it limits the uptake of both iron and gallium in resistant cells (Davies et al. 2006). Moreover, C282Y mutation in *HFE* gene was shown to be related with resistance to Temodar, geldanamycin, doxorubicin and gammaradiation in human neuroblastoma (SH-SH5Y) and glioma cells (Lee et al. 2011a, b).

Hepcidin and drug resistance

Hepcidin is a hormone that regulates iron metabolism. It has three isoforms containing 20, 22 and 25 amino acids. The connection between hepcidin and iron metabolism was revealed by iron-overloading studies where hepcidin was over-expressed in hepatocytes. Hepcidin-defficient mice were also shown to have iron over-load in pancreas and liver and hepcidin-overexpressing mice displayed iron deficiency and microcytic hypochromic anemia. By these studies, hepcidin was proposed as a controller of iron transport (Reviewed by Ganz 2003; Nicolas et al. 2002). Yalovenko et al. determined that hepcidin levels were 1.5-2 times higher in doxorubicin- and cisplatinresistant MCF7 cells. The exogenous administration of hepcidin as well as free iron caused the development of resistance to doxorubicin in rats bearing Walker-256 carcinosarcoma cells (Yalovenko et al. 2016).

Iron efflux, iron accumulation and iron-related proteins, and drug resistance

Watts et al. showed that nitric oxide (NO)-mediated iron efflux together with glutathione was greater in MCF-VP cells which are resistant to VP-16 etoposide and over-express multidrug resistance-associated protein 1 (MRP1) compared to wild-type cells (Watts et al. 2006). Similarly, a multidrug resistance-related protein, breast cancer resistance protein (BCRP; ABCG2) was also found to be responsible for the efflux of heme, a complex of iron and protoporphyrin IX (PPIX) and a regulatory molecule affecting gene expression and translation (Krishnamurthy et al. 2007).

Checkhun et al. showed that free iron complexes were elevated in cisplatin-resistant Guerin carcinomatransplanted rats (Chekhun et al. 2014).

Tumor associated macrophages (TAMs) was shown to trigger IL-6 secretion in breast cancer cell lines, 4T1 and MCF7, in a paracrine loop manner, resulting in an IL-6 rich niche which generates *de novo* drug resistance. Iron was proved to enhance this positive feedback loop between TAMs and IL-6, which further strengthens resistance. The fact that an iron chelator, desferrioxamine (DFO), could prevent resistance status by interfering with the function of iron (Li et al. 2016) strongly indicates the importance of iron metabolism in drug resistance. Contrary to that study, Yang et al. determined that DFO promoted resistance to BCNU by activating hypoxia-inducible factor 1 (HIF-1) in rat C6 glioma cells (Yang et al. 2004).

Utilization of iron metabolism to overcome the drug-resistance

From now on, we will focus on the studies that utilize iron metabolism to overcome the drug resistance in cancer cell lines. We deliberately eliminated the resistance status of the drug-resistant cancer cells to iron-related drugs, mainly iron chelators, and were only interested in the restoring sensitivity to anticancer drugs via manipulation of cellular iron metabolism. Table 2 summarizes fold decreases in IC_{50} (inhibitory concentration 50) values of anticancer agents upon alterations in iron metabolism. As seen in Table 2, we classified these strategies into three groups which are explained below.

Transferrin receptor-targeted strategies

Transferrin-conjugated drug studies to overcome the drug resistance were partly reviewed by different research groups (Daniels et al. 2012; Head et al. 1997; Li and Qian 2002; Li et al. 2002; Qian et al. 2002).

Doxorubicin-transferrin conjugate is internalized through a transmembrane mechanism contrary to the function of P-glycoprotein which is active when doxorubicin is administered alone (Li and Qian 2002). Hatano et al. used transferrin-doxorubicin

Table 2 Efficiencies of iron metabolism-related manipulations to overcome drug resistance in cancer

Cell Line	Drug	Method	IC ₅₀ fold decrease	Reference
K562	Doxorubicin	TfR-targeted	5.6	Fritzer et al. (1992)
			5.5	Hatano et al. (1993)
			3.5	Suzuki et al. (1997)
			5.2	Wu et al. (2007)
L929			130	Lai et al. (1998)
SBC-3			3.3	Kobayashi et al. (2007)
HL60			8.7	Fritzer et al. (1992)
			200	Lubgan et al. (2009)
KB-8-5			4	Fritzer et al. (1996)
H69VP	Artemisinin	Tf-treated	4.4	Sadava et al. (2002)
CCRF-CEM	Galium		3-4*	Chitambar and Wereley (1997)
MCF-7/VP	Etoposide	Iron chelating	16.7	Whitnall et al. (2006)
KB-V1	Vinblastine		8.3	
RPMI-8226	Bortezomib	FtH downregulation	6.7	Campanella et al. (2013)
U251	BCNU		2.5**	Liu et al. (2011)
MDA-MB-231	Doxorubicin		5	Shpyleva et al. (2011)
MCF-7			7.9	Chekhun et al. (2013)
	Cisplatin		2	

* Cell death ratio

** LD value

conjugate against doxorubicin-resistant K562 cells and showed that this conjugate reduced the IC_{50} value from 20 to 3.6 µM (Hatano et al. 1993). Likewise, Fritzer et al. conjugated doxorubicin to transferrin to overcome the multidrug resistance by targeting TfR in multidrug-resistant HeLa contaminant KB cell lines, KB-8-5, KB-C1 and KB-V1. It has been shown that the conjugate completely re-sensitized drug-resistant cells, lowering IC₅₀ values of doxorubicin to that of sensitive ones (Fritzer et al. 1996). In another study by Lai et al., doxorubicin-resistant murine L929 cell line was used to assess the effect of transferrin-doxorubicin conjugate. The transferrin-doxorubicin conjugate mainly localized in cytoplasm while free doxorubicin dispersed through cell membrane, cytoplasm and, nucleus. The conjugation to transferrin did not only provide higher uptake of doxorubicin through TfR but also helps to achieve higher cytoplasmic concentrations of the drug (Lai et al. 1998). The transferrindoxorubicin conjugate was also shown to lower IC_{50} of doxorubicin from 7 to 0.035 µM in doxorubicinresistant human promyelocytic cell line, HL60 (Lubgan et al. 2009). In another study, Fritzer et al. utilized transferrin-doxorubicin conjugate against doxorubicin-resistant HL60 and K562 cell lines and showed that Tf-Dox conjugate significantly decreased the IC_{50} values in both cell lines (Fritzer et al. 1992).

TfR was targeted by egg-phosphatidylcholine (PC)/cholesterol liposomes encapsulating doxorubicin to overcome MDR in human small cell lung cancer cell line, SBC-3. Transferrin was covalently conjugated to liposomes so as to target TfR-mediated endocytic pathway and overcome P-glycoproteinmediated drug export. Short-term exposure to TfRtargeted liposome-doxorubicin conjugate increased the toxicity compared to free doxorubicin. Targeted liposomes resulted in high doxorubicin accumulation compared to non-targeted liposome and free doxorubicin in both drug-sensitive and -resistant cells (Kobayashi et al. 2007). In a similar study, Wu et al. designed transferrin-conjugated liposomes co-encapsulating doxorubicin and verapamil (Tf-L-Dox/Ver) to minimize systemic cardiotoxicity of drugs while maximizing cytotoxicity in K562 cell lines. Tf-L-Dox/Ver displayed 5.2 times greater cytotoxicity than non-targeted conjugate and 2.8 times greater cytotoxicity than Tf-L-Dox alone in doxorubicin-resistant K562 cells (Wu et al. 2007). Additionally,

doxorubicin-encapsulated anti-TfR antibody-conjugated immunoliposomes were shown to increase intracellular doxorubicin and the duration of intracellular drug accumulation as well as to re-sensitize doxorubicin-resistant cells against the drug (Suzuki et al. 1997). In addition to liposomes, nanoparticles were also used in TfR-targeting strategies. A biodegradable polymer, poly(lactic-co-glycolide) (PLGA) was loaded with paclitaxel and conjugated to transferrin, and assessed in both drug-sensitive and multidrug-resistant MCF7 (MCF7/ADR) cells. Intracellular drug levels, and consequently cell death, were enhanced when treated with Tf-drug conjugate compared to nanoparticles without transferrin conjugation. Such approaches were proposed as new strategies to overcome the multidrug resistance in cancer cells (Sahoo and Labhasetwar 2005).

Artemisinin (ART) is a drug used predominantly in treatment of drug-resistant malaria and also found to be effective in several tumor cells (Klayman 1985; Olliaro et al. 2001; Li et al. 2001). ART only becomes toxic in the presence of Fe^{2+} . Sadava et al. reported human multidrug-resistant small cell-lung cancer (SCLC) cell line H69VP was 10-fold resistant to ART compared to parental H69. Transferrin treatment increased cytotoxicity of ART, achieving complete resensitization in H69VP cells. Drug-resistant H69 cell line was shown to have high levels of TfR on cell membrane; therefore, iron uptake was greater than drug-sensitive H69 which, as a result, was not affected by transferrin treatment. They proved the effect of iron by using iron chelator, desferrioxamine mesylate, which reversed the IC50 values of ART with transferrin to ART alone in H69VP while did not affect parental H69 cells. Thus, ART-transferrin co-therapy was offered for the treatment of drug-resistant SCLC (Sadava et al. 2002).

Iron chelators

Iron chelators are complexation agents used to bind free iron and are widely used in treatments of different diseases (Pierre et al. 2002; Sheth 2014; Torti and Torti 2013a). Here, we evaluated the studies that used iron chelators to overcome the drug resistance in cancer.

Whitnall et al. assessed the effect of an iron chelator, di-2-pyridylketone-4,4,-dimethyl3-thiosemicarbazone (Dp44mT), which significantly reversed drug-resistance in etoposide-resistant MCF7 and vinblastine-resistant KB-V1 epidermal carcinoma cells (Whitnall et al. 2006).

In a clinical study, a patient with gemtuzumab ozogamicin-resistant acute monocytic leukemia with a high serum ferritin level was treated with iron chelator, deferasirox. Deferasirox reduced the serum ferritin, eliminating continuous need of blood transfusions and helping complete remission in the patient (Fukushima et al. 2011). Deferasirox was also shown to be effective to overcome the imatinib-resistance in chronic myeloid leukemia cell lines by inducing apoptosis in vitro (Kim et al. 2016).

Iron deprivation by DFO was shown to overcome the multidrug resistance in K562 cell lines by decreasing intracellular iron level and MDR1 expression (Fang et al. 2010).

Ganguly et al. synthesized a novel iron chelator, iron N-(2-hydroxy acetophenone) glycinate (FeNG) that was found to be highly effective in doxorubicin– resistant T lymphoblastic leukemia cell lines, CCRF-CEM. In doxorubicin-resistant cells, FeNG triggered the production of reactive oxygen species (ROS), leading cells to apoptosis (Ganguly et al. 2010). Similarly, FeNG treatment resulted in depletion of intracellular glutathione and increase in effectiveness of doxorubicin in drug-resistant Ehrlich ascites carcinoma cells ex vivo and in vivo (Ganguly et al. 2012).

Beside the iron chelators, several complexes containing iron have been also used to overcome drug resistance in cancer. Lee et al. synthesized a novel iron complex (FeIII(salophene)Cl) which proved remarkably effective in vincristine- and daunorubicin-resistant leukemic Nalm-6 cells (Lee et al. 2011a, b). Another Fe(III)-salophene conjugate was shown to decrease cell viability in a dose-dependent manner by triggering S phase arrest and apoptosis in platinumresistant ovarian epithelial adenocarcinoma cell lines, SKOV-3 and OVCAR-3 (Lange et al. 2008).

Genetic manipulations

In this strategy, iron metabolism-related proteins are over-expressed and/or silenced to examine the changes in drug resistance upon these alterations.

Liu et al. hypothesized that FtH could protect DNA from damage and the silencing of *FtH* could increase the sensitivity of glioma cells against the chemotherapeutic agents. It was reported that silencing of *FtH*

decreased lethal dosage (LD) value of BCNU with high active caspase 3 levels in U251 glioma, MCF7 breast adenocarcinoma and malignant peripheral nerve sheath tumor cell lines. Downregulation of *FtH* also reduced tumor size in drug-treated animal models (Liu et al. 2011). Similarly, in MDA-MB-231 cells, down-regulation of *FTH1* by miR-200b brought about increased sensitivity to doxorubicin (Shpyleva et al. 2011).

Chekhun et al. silenced *FTL* in doxorubicin- and cisplatin-resistant MCF-7 cells by targeting with miR-133a, and showed that downregulation of *FTL* increased the effectiveness of both drugs on drug-resistant cells (Chekhun et al. 2013).

Ferroptosis and drug resistance

Ferroptosis is a form of non-apoptotic cell death which is dependent on intracellular iron and reactive oxygen species. Intracellular iron triggers reactive oxygen species via the oncogenic Ras-selective lethal small molecule erastin. A small molecule, called ferrostatin-1 and iron chelators were shown to inhibit the ferroptosis (Dixon et al. 2012).

At morphological, biochemical and genetic levels, ferroptosis is distinguishable from other regulated and non-regulated cell death mechanisms. Cells that undergo ferroptosis are rounded up and have dysmorphic mitochondrial phenotype with condensed mitochondrial membrane density (Cao and Dixon 2016; Xie et al. 2016). The induction of cell death in apoptosis-deficient Bax/Bak double knockout mouse embryonic fibroblasts after erastin treatment indicates that ferroptosis in dependent of apoptotic machinery since erastin do not cause apoptotic changes such as chromatin margination, activation of caspases or cleavage of PARP (Dixon et al. 2012). Ferroptosis cannot be attenuated by deletion of Bcl2 family proteins, necroptosis and autophagy inhibitors suggesting that ferroptotic cell death requires different cellular stimulants than other cell death mechanisms (Cao and Dixon 2016). Ferroptosis is triggered by accumulated toxic lipid peroxidation products and iron metabolism-derived reactive oxygen species. This occurs by activation of mitochondrial voltagedependent anion channels and mitogen-activated protein kinases, enhanced endoplasmic reticulum stress, and inhibition of cystine/glutamate antiporter, unbalancing antioxidant defenses of the cell and resulting in iron-dependent oxidative cell death (Dixon et al. 2012). Misregulated ferroptosis has been shown to be implicated in pathological and physiological conditions (Xie et al. 2016).

Yu et al. used acute myeloid leukemia cell lines to demonstrate the role of ferroptosis inducer, erastin towards activity of the chemotherapeutic drugs, cytarabine and doxorubicin. Treatment with low-dose erastin significantly increased the anticancer activity of cytarabine and doxorubicin in HL60 cells. The enhanced sensitivity to cytarabine and doxorubicin was in a RAS-independent manner given that inhibition of c-Jun and p38, but not ERK, induced resistance to erastin and other anticancer agents (Yu et al. 2015).

Induction of ferroptosis by inhibition of cystine/ glutamate antiporter and glutathione peroxidase by cystine/glutamate antiporter inhibitors, erastin and sulfasalazine was shown to overcome cisplatin resistance in head and neck cancer cell lines (AMC-HN3R. -HN4R and -HN9R). Also, high glutamine without cysteine decreased the cell viability even in resistant cell lines, resulting in elevated levels of reactive oxygen species and leading to ferroptosis (Roh et al. 2016). Erastin and sulfasalizine were also assessed to increase the efficiency of Temozolomide (TMZ), an alkylating agent, towards glioblastoma multiforme (GBM) which were proved to be resistant to radio- and chemotherapy. TMZ was shown to increase the expression of xCT, the subunit of glutamate/cysteine transporter system x_c⁻ and associating with glutathione (GSH) synthesis and ROS accumulation, via upregulating NF-E2 related factor 2 (Nrf2) and activating transcription factor 4 (ATF4) transcription factors. Inhibition of xCT by erastin and sulfasalazine decreased the level of GSH and increased the amount of ROS in GBM cell lines, which sensitized the cells towards TMZ via enhanced ferroptosis (Chen et al. 2015). In another study, Nrf2 has been shown to inhibit ferroptosis in cisplatin-resistant head and neck cancer cell lines. Artesunate treatment decreased the cellular GSH levels and increased the ROS formation, which resulted in ferroptosis. Genetic silencing of Nrf2 reversed the inhibition of ferroptosis in these cell lines (Roh et al. 2017).

As an inhibitor of xCT, sulfasalazine also promotes ferroptosis in a similar manner to erastin (Sehm et al. 2016). Therefore, it is not surprising that sulfasalazine has been shown to sensitize colorectal cancer cell lines to cisplatin (Ma et al. 2015); lung adenocarcinoma cell lines and breast cancer cell lines to doxorubicin (Lay et al. 2007; Narang et al. 2007) by triggering ferroptosis.

Sun et al. studied the mechanisms of sorafenib, an inhibitor of multiple oncogenic kinases, in primary tumor hepatocytes, hepatocellular carcinoma lines Huh7, HepaG2 and Hep3b, and in animal models. Sorafenib induced the expression of metallothionein (MT)-1G, which functions in heavy metal detoxification process. Inhibition of MT-1G increased the sensitivity to sorafenib in the cell lines and tumor xenografts models. Furthermore, silencing *MT-1G* was shown to promote glutathione depletion and lipid peroxidation, triggering ferroptosis in aforementioned cells (Sun et al. 2016).

Conclusions and future perspectives

In this review, we focused on iron metabolism and drug resistance in cancer cells. Resistance to anticancer drugs remarkably alters the iron metabolism depending on type of cancer and drugs used. Although drug-resistant phenotype exhibits itself as insensitivity to chemotherapeutics, the genetic alterations leading to drug resistance differ from cell to cell, even in the same cell line that are resistant to different concentrations of the same anticancer agent (Iseri et al. 2012). Since iron is an essential element for cellular proliferation, various drug-resistant cells with rapid proliferation rates, in general, enhance iron uptake and increase intracellular iron level. Iron chelators are proved to be effective to overcome drug resistance in these cells by binding free iron and interfering with overall iron metabolism (Whitnall et al. 2006; Fang et al. 2010; Fukushima et al. 2011; Kim et al. 2016). In other cases, drug resistance seemed to be significantly reversed by supplementation of iron-containing compounds through arrest of cellular growth and, subsequently, enhanced apoptosis (Lee et al. 2011a, b; Lange et al. 2008) although such studies are lack of definition of mechanisms by which these compounds exert their effects. Still, strategies that target iron metabolism definitely increase the efficacy of anticancer drugs regardless of the approach. However, more comprehensive studies are still required in this area. As a ROS-dependent cell death mechanism, ferroptosis is greatly linked to drug resistance in cancer. It is known that chemotherapeutic agents trigger elevated ROS formation (Conklin 2004; Arun et al. 2016), however, drug resistant cancer cells develop several mechanisms, particularly increasing GSH levels, to avoid the detrimental effects of ROS (Chen et al. 2015). Any attempt disrupting these mechanisms would inhibit the detoxification processes and could promote ferroptosis. Moreover, accumulation of intracellular free iron would stimulate ROS formation and, thus, ferroptosis (Roh et al. 2017), which would further explain the logic beyond the success of iron treatment in reversal of drug resistance. Furthermore, sulfasalazine, as a ferroptosis-related inhibitor, was shown to be a substrate of breast cancer resistance protein (BCRP/ABCG2; van der Heijden et al. 2004; Tomaru et al. 2013). Elevated levels of BCRP substrates may decrease the rate of drug efflux, which, in turn, triggers ferroptosis and increases the efficiency of drugs. Still, to explore the role of iron metabolism in drug resistance and to overcome the resistance through utilization of iron metabolism seems to be a great challenge.

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