

Apatinib Promotes Ferroptosis in Colorectal Cancer Cells by Targeting ELOVL6/ACSL4 Signaling

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Background: Colorectal cancer (CRC) is a common digestive system malignancy. Ferroptosis, a new form of regulated cell death, plays a vital role in the pathogenesis and therapy of cancers.

Objective: We aimed to study the role of apatinib in ferroptosis of CRC cells and its potential mechanisms.

Materials and Methods: Human CRC HCT116 cells were exposed to apatinib. Cell viability was examined using a CCK-8 kit. The concentrations of intracellular iron and reactive oxygen species (ROS) were detected using kits. Additionally, Western blot analysis was used to determine the expression of ferroptosis-related proteins. Elongation of very long-chain fatty acids family member 6 (ELOVL6) was one of the targets of apatinib predicted by SwissTargetPrediction. Therefore, ELOVL6 expression was evaluated after treatment with apatinib. Subsequently, the effects of ELOVL6 overexpression on ferroptosis of HCT116 cells were investigated. Finally, STRING database was applied to predict the potential proteins interacting with ELOVL6, and co-immunoprecipitation (co-IP) assay was applied for confirmation.

Results: Results indicated that apatinib decreased cell viability and increased the contents of intracellular iron ROS. Moreover, significantly upregulated ACSL4 expression was observed, accompanied by notable downregulation of GPx4 and FTH1 expression after apatinib exposure. Furthermore, ELOVL6 expression was remarkably enhanced in HCT116 cells, which was dramatically inhibited under apatinib intervention. ELOVL6 overexpression reversed the effects of apatinib on cell viability and ferroptosis of HCT116 cells. Moreover, ACSL4, a vital regulator of ferroptosis, could interact with ELOVL6 directly, which was confirmed by the result of co-IP.

Conclusion: These findings demonstrated that apatinib promoted ferroptosis in CRC cells by targeting ELOVL6/ACSL4, providing a new mechanism support for apatinib application in the clinical treatment of CRC.

Keywords: colorectal cancer, apatinib, ferroptosis, reactive oxygen species, ELOVL6

Introduction

Colorectal cancer (CRC) is a common and adverse malignancy in digestive system and a major cause for cancer-related mortality worldwide.^{1,2} Recent statistics have revealed that 1.8 million CRC cases are diagnosed annually and that each year, 881,000 CRC-related deaths occur worldwide.³ Although considerable progress has been made in the treatment of CRC recently, the prognosis of patients with CRC still remains dismal.⁴ Therefore, it is urgent to obtain a clearer understanding about the mechanisms underlying the anti-cancer effects of drugs, thereby developing more effective preventive strategies for CRC therapy.

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Ferroptosis was recently identified as a new form of regulated cell death by Brent R. Stockwell's laboratory in 2012.⁵ It is characterized by an overwhelming and iron-dependent accumulation of lipid peroxide and reactive oxygen species (ROS) upon a distinct set of genes, thus evoking a different method of death from apoptosis and necrosis.^{6,7} For excessive growth, tumor cells present a higher demand for iron than non-tumor cells, which enables tumor cells more susceptible to iron-catalyzed necrosis.⁸ Ferroptosis plays a vital role in the pathogenesis and cancer therapy.⁹ A growing body of literature has proved that ferroptosis, as a new form to regulate cell death, is a novel method for the destruction of cancer cells.¹⁰ Therefore, triggering ferroptosis might offer a novel therapeutic strategy for treating CRC.

Apatinib, also known as YN968D1, is a new oral small-molecule agent with antiangiogenic effect for the treatment of multiple solid tumors, including gastric cancer, colorectal cancer and hepatocellular carcinoma.^{11,12} It has been well reported that apatinib serves as an optional treatment in metastatic CRC.¹³ Apatinib, used as third-line treatment, effectively improves the prognosis of patients with heavily treated metastatic CRC.¹⁴ Besides, apatinib could induce apoptosis and protective autophagy in CRC cells by endoplasmic reticulum stress.¹⁵ Additionally, the SwissTargetPrediction website (<http://www.swisstargetprediction.ch/>) predicts that elongation of very long-chain fatty acids family member 6 (ELOVL6) ELOVL6 is one target of apatinib. Accumulating evidence has demonstrated that ELOVL6 is high-expressed and serves as a negative clinical predictor in a plenty of carcinomas.^{16,17} However, whether the anti-CRC activity of apatinib acts by regulating ferroptosis of CRC cells through ELOVL6 remains to be elucidated.

In the present study, human colorectal cancer HCT116 cells were exposed to apatinib for investigating its effects on viability and ferroptosis. Afterwards, ELOVL6 was overexpressed to explore the underlying regulatory mechanisms of apatinib in ferroptosis of CRC cells. Our study is of great significance since it provides a new mechanism support for apatinib application in the clinical treatment of colorectal cancer.

Materials and Methods

Cell Culture and Treatment

Human colorectal cancer HCT116 cells and human normal intestinal epithelial cell line (HIEC) were procured from the Cell Bank of Shanghai Institute of Biochemistry and

Cell Biology, Chinese Academy of Sciences. Cells were maintained in Dulbecco's modified Eagle medium (DMEM; Invitrogen, Carlsbad, USA) supplemented with 10% heat-inactivated FBS (Gibco; Thermo Fisher Scientific, Inc.) at 37°C in fully humidified air of 95% air and 5% CO₂. HCT116 cells were, respectively, exposed to apatinib (Hengrui Medicine Co. Ltd., Jiangsu, China) of different concentrations at 5 μM, 10 μM and 20 μM for 24 h. Untreated HCT116 cells were used as the control cells.

Cell Viability Assay

The cytotoxicity of apatinib on CRC cells was determined using a Cell Counting Kit-8 (CCK-8) assay kit (Shanghai Yi Sheng Biotechnology Co. Ltd., China). Briefly, HCT116 cells in the logarithmic growth period were collected and dispensed into 96-well tissue culture plates (4000 cells/well), followed by treatment with the indicated concentrations of apatinib. Following incubation for 24, 48 or 72 h, ten microliters of CCK-8 solution was added to each hole. After incubation at 37°C for further 2 h, the optical density (OD) at 450 nm of cells was detected utilizing a microplate reader (Bio-Rad Laboratories, Richmond, CA, USA).

Cell Transfection

Prior to cell transfection, HCT116 cells were plated into a 6-well plate and cultured until 80% confluence at 37°C in a humidified environment of 5% CO₂. ELOVL6-overexpressing plasmid (Oe-ELOVL6) or the empty vector plasmid (Oe-NC), provided by Shanghai GenePharma co., Ltd (Shanghai, China), was transfected into HCT116 cells. Transfection was performed by Lipofectamine 3000 (Invitrogen, Carlsbad, CA, USA) as per the manufacturer's protocol. At 24 h after transfection, cells were harvested and the transfection efficiency was evaluated by applying reverse transcription-quantitative polymerase chain reaction (RT-qPCR).

Measurement of Total Iron Content and Ferrous Iron (Fe²⁺) Concentration

An iron assay kit (Baiaolaibo, Beijing, China) was employed to measure total iron concentration in cell lysates in accordance with the manufacturer's guidelines. The detection of relative intracellular Fe²⁺ level was performed with the iron assay kit (Abcam) following

manufacturer's recommendations. The experiments were independently repeated at least three times.

Detection of Intracellular Reactive Oxygen Species (ROS)

The accumulation of intracellular ROS was examined by a ROS assay kit (Nanjing Jiancheng Bioengineering Institute, Nanjing, China) with usage of 2, 7-dichlorofluorescein diacetate (DCFH-DA, Invitrogen, USA) as a fluorescence probe. After appropriate treatment, the HCT116 cells were collected and washed with ice-cold phosphate buffered saline (PBS) thrice, followed by incubation in DMEM supplemented with 10 μ M DCFH-DA for 20 min at 37°C in the dark. After centrifugation at 800 g, 4°C for 5 min, the fluorescence was evaluated by a fluorimeter excited at 488 nm and emitted at 525 nm. The images were photographed by a confocal microscope.

Test for the Markers of Oxidative Stress

Cells were collected and centrifuged at 3000 x g for 10 min at 4°C, and the supernatant was collected for subsequent examinations. The concentrations of malondialdehyde (MDA) and glutathione (GSH) as well as the activity of glutathione peroxidase (GPx) in HCT116 cells were detected using commercial kits purchased from Nanjing Jiancheng Bioengineering Institute (Nanjing, China) according to the colorimetric methods.

RT-qPCR Analysis

Total RNA in cells was extracted using TRIzol reagent (Invitrogen). The complementary DNA (cDNA) was synthesized by a PrimeScript RT Reagent Kit (Takara, Japan). PCR then was performed with 2 μ g cDNA as the template using Power SYBR Master Mix (Applied Biosystems, Foster, USA) on the ABI 7500 PCR system (Applied Biosystems). All primers used in the present study were synthesized by Shanghai GenePharma co., Ltd (Shanghai, China). Glyceraldehyde 3-phosphate dehydrogenase (GAPDH) was the internal reference gene. Relative expression was calculated using the $2^{-\Delta\Delta Ct}$ method.

Western Blot Analysis

Total proteins were extracted from cells by using RIPA Lysis Buffer (Beyotime, Shanghai, China). A bicinchoninic acid protein assay kit (Beyotime, Shanghai, China) was applied to detect the protein concentration. A total of 40 μ g protein samples in cell lysates were loaded on SDS-PAGE, followed

by transferring onto PVDF membranes. Membranes were subsequently immersed into 5% nonfat milk, and then incubated with primary antibodies. All blots were incubated with HRP-conjugated secondary antibody (A0216, Beyotime, Shanghai, China) for 1 h and next visualized using Odyssey Infrared Imaging Scanner (LI-COR Biosciences). GAPDH was used as the loading control. The intensities of bands were calculated using Image-J software (National Institutes of Health).

Co-Immunoprecipitation (IP) Assay

For co-IP assay, HCT116 cells were washed in 2 mL Phosphate Buffer Saline (PBS; Beyotime, Shanghai, China) and centrifuged at 850 x g at room temperature for 5 min to collect the cells. Then, cells were lysed in lysis Buffer (Beyotime, Shanghai, China). Lysates were incubated with specific antibody and control IgG plus Protein A/G beads (Santa Cruz Biotechnology, USA) at 4°C for 2 h. After washing the beads with PBS three times, immunoprecipitates were analyzed by Western blot analysis.

Statistical Analysis

All data were represented as mean values \pm standard deviation. Statistical analysis was performed with GraphPad Prism 6 (GraphPad Software, Inc., La Jolla, CA, USA). Differences between groups were evaluated by Student's *t*-test. Comparisons involving multiple samples were analyzed by one-way analysis of variance (ANOVA) followed by Turkey's post hoc test. P values less than 0.05 were considered statistically significant.

Results

Apatinib Inhibits Cell Viability and Promotes Ferroptosis in HCT116 Cells

The chemical structural formula of apatinib was presented in [Figure 1A](#). The viability of HCT116 cells after treatment with apatinib at different concentrations was determined using a CCK-8 assay kit. As exhibited in [Figure 1B](#), apatinib dose-dependently decreased cell viability compared with the untreated control group. Apatinib at concentration of 5, 10 or 20 μ M to treat HCT116 cells was selected for the following experiments, which was consistent with the previous study.¹⁵ Iron accumulation and lipid ROS production are two critical signaling events in ferroptosis.¹⁸ Result of [Figure 1C](#) and [D](#) indicated that the levels of total iron and Fe²⁺ were dramatically reduced after apatinib intervention relative to

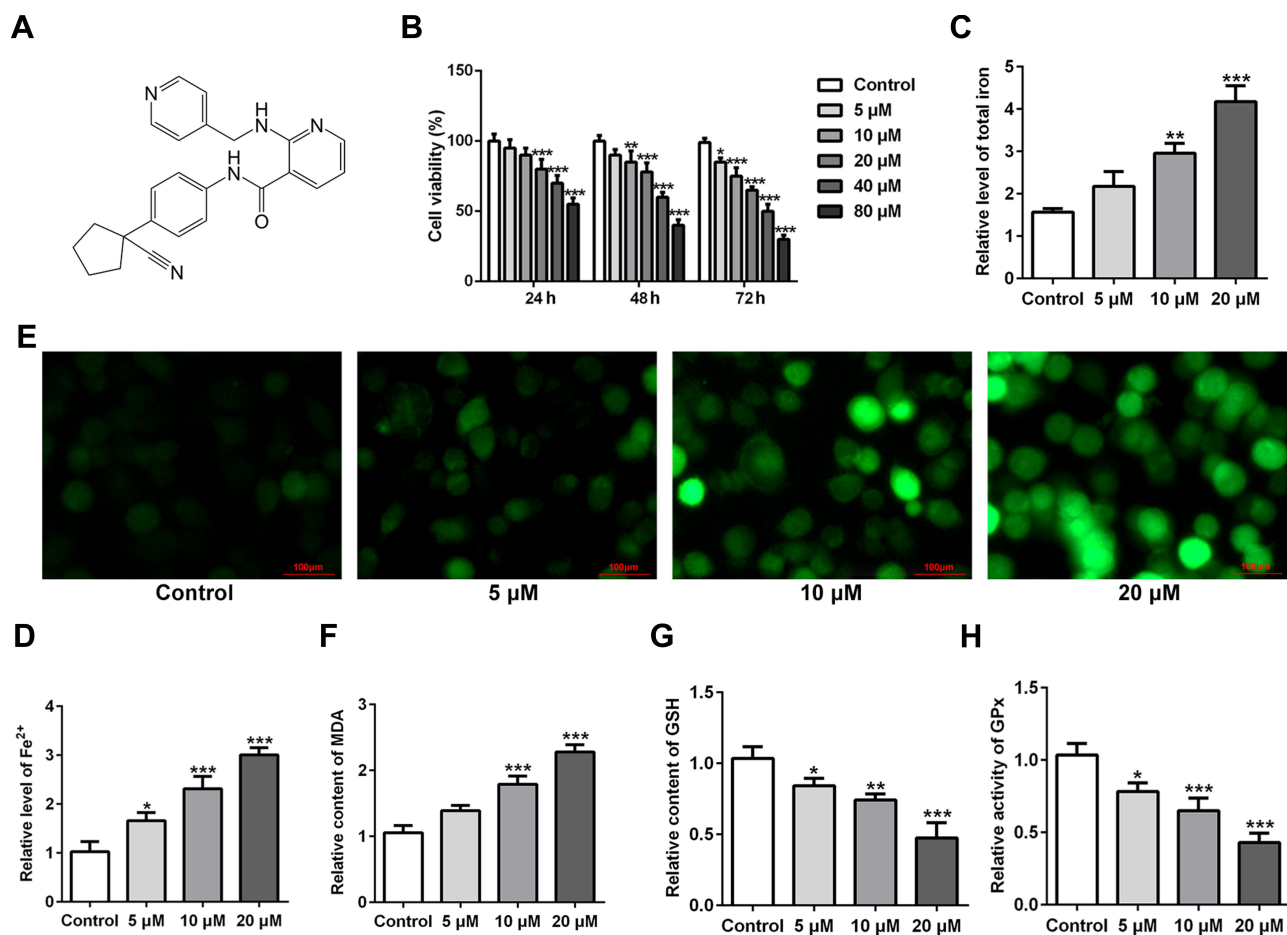


Figure 1 Apatinib inhibits cell viability and promotes ferroptosis in HCT116 cells. **(A)** The chemical structural formula of Apatinib. **(B)** Cell viability of HCT116 cells after treatment with apatinib (5, 10 and 20 μM) was determined using a cell counting kit-8 assay kit. The relative concentrations of **(C)** total iron and **(D)** Fe²⁺ were measured using commercially available kits, respectively. **(E)** The level of ROS was detected by a ROS assay kit using 2, 7-dichlorofluorescein diacetate (DCFH-DA, Invitrogen, USA). The levels of **(F)** MDA, **(G)** GSH and **(H)** GPx were evaluated using commercial kits. *P<0.05, **P<0.01 and ***P<0.001 vs control.

Abbreviations: ROS, reactive oxygen species; MDA, malondialdehyde; GSH, glutathione; GPx, glutathione peroxidase.

the control group. Additionally, the ROS accumulation was significantly elevated in a dose-dependent manner of apatinib (Figure 1E). Moreover, notably increased concentration of MDA while decreased contents of GSH and GPx were observed under the influence of apatinib (Figure 1F–H). Summing up, these data suggest that apatinib suppresses cell viability and promotes ferroptosis in HCT116 cells.

Apatinib Regulates the Expression of Ferroptosis-Associated Proteins in HCT116 Cells

Subsequently, the expression of ferroptosis-associated proteins was separately examined via applying Western blot analysis. It is observable from Figure 2 that apatinib remarkably upregulated the expression of long-chain acyl-CoA synthetases 4 (ACSL4) whilst downregulated the

levels of glutathione peroxidase 4 (GPx4) and ferritin heavy chain (FTH1) compared with the control group. These data show evidence that apatinib can regulate the expression of ferroptosis-associated proteins in HCT116 cells.

Apatinib Inhibits the ELOVL6 Expression in HCT116 Cells

To study the underlying mechanisms of apatinib in modulating ferroptosis of HCT116 cells, the SwissTargetPrediction website (<http://www.swisstargetprediction.ch/>) was employed. ELOVL6 was noticed as a target of apatinib. The GEPIA database (<http://gepia.cancer-pku.cn/>) displayed the high expression of ELOVL6 in CRC tissues compared with the normal tissues (Figure 3A). Then, the expression of ELOVL6 in HCT116 cells was determined using RT-qPCR and Western blot

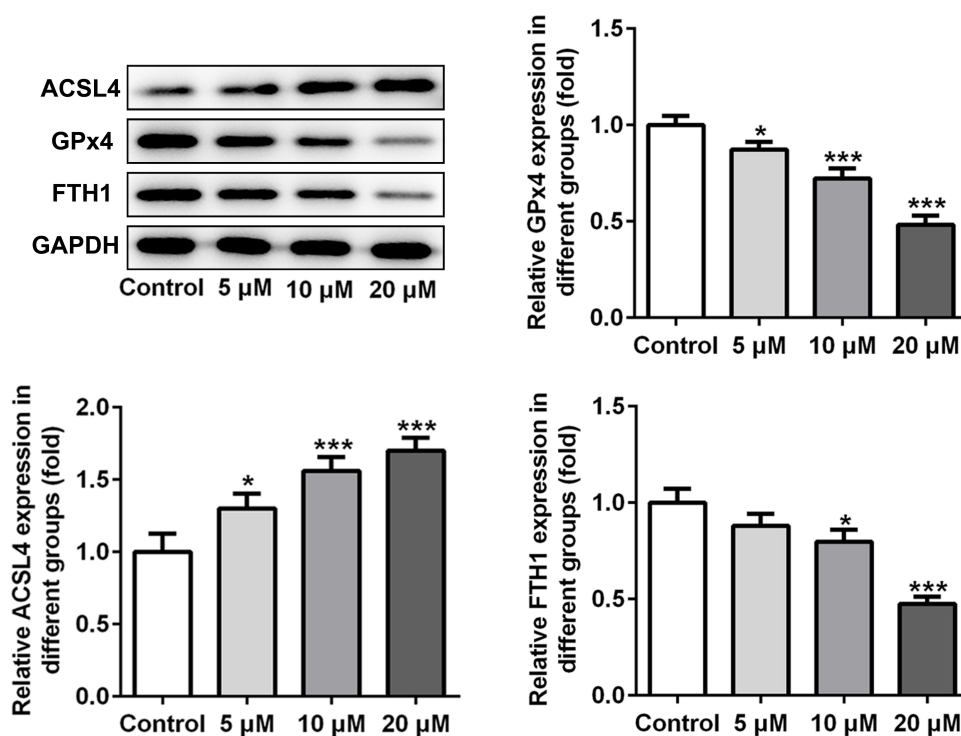


Figure 2 Apatinib regulates the expression of ferroptosis-associated proteins in HCT116 cells. The expression of ACSL4, GPx4 and FTH1 was determined using Western blot analysis. * $P < 0.05$ *** $P < 0.001$ vs control.

Abbreviations: ACSL4, long-chain acyl-CoA synthetases 4; GPx4, glutathione peroxidase 4; FTH1, ferritin heavy chain.

analyses. As shown in [Figure 3B](#) and [C](#), ELOVL6 was heightened in HCT116 cells in comparison with the HIEC cells. Moreover, apatinib exposure remarkably downregulated ELOVL6 at the transcriptional and protein levels in a dose-dependent manner ([Figure 3D](#) and [E](#)). These observations reveal that apatinib possesses an inhibitory effect on the ELOVL6 expression in HCT116 cells.

ELOVL6 Overexpression Reverses the Effects of Apatinib on Cell Viability and Ferroptosis

To further investigate the specific role of ELOVL6 in CRC cells, ELOVL6 expression was overexpressed by transfecting overexpressing plasmid. After transfection, significant upregulation of ELOVL6 mRNA and protein expression are observed in [Figure 4A](#) and [B](#). Afterwards, cell viability was tested using CCK-8 assay. In [Figure 4C](#), ELOVL6 upregulation notably elevated the viability of HCT116 cells as compared to the 20 μM apatinib+Oe-NC group. As displayed in [Figure 4D](#), the expression of ELOVL6 protein was conspicuously upregulated after ELOVL6 overexpression in apatinib-treated HCT116 cells. Additionally, the levels of intracellular total iron and Fe^{2+} were examined to analyze

the function of ELOVL6 overexpression in ferroptosis. Results of [Figure 4E](#) and [F](#) elucidated that the suppressive effects of apatinib on both total iron and Fe^{2+} in HCT116 cells were reversed with ELOVL6 overexpression. Moreover, the production of ROS was significantly suppressed in apatinib-treated and ELOVL6-upregulated HCT116 cells relative to the apatinib-challenged HCT116 cells ([Figure 4G](#)). Consistently, in comparison with the 20 μM apatinib+Oe-NC group, ELOVL6 overexpression notably reduced the content of MDA while enhanced the levels of GSH and GPx ([Figure 4H–J](#)). Collectively, these results implicate that ELOVL6 overexpression neutralizes the effects of apatinib on cell viability and ferroptosis.

ELOVL6 Overexpression Restores the Regulatory Effects of Apatinib on the Expression of Ferroptosis-Associated Proteins via Regulating ACSL4 Expression

Then, the role of ELOVL6 overexpression in modulating the levels of ferroptosis-associated proteins was determined through Western blot analysis. As displayed in [Figure 5A–D](#), the increase in ACSL4 expression as well as the decrease in GPx4 and FTH1 expression resulted

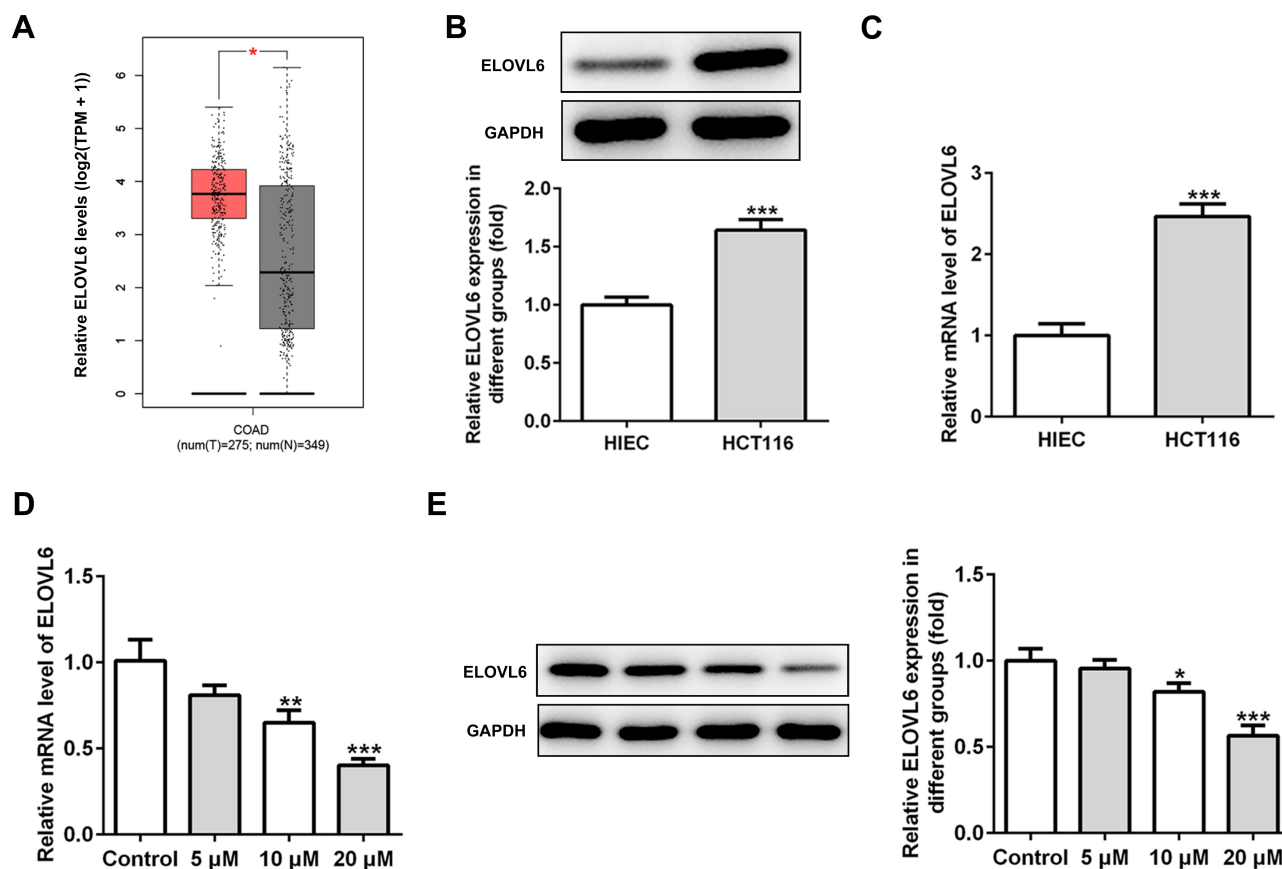


Figure 3 Apatinib inhibits the ELOVL6 expression in HCT116 cells. **(A)** The GEPIA database (<http://gepia.cancer-pku.cn/>) displayed that ELOVL6 is highly expressed in CRC tissues compared with the normal tissues. The expression of ELOVL6 in HCT116 and HIEC cells was detected using **(B)** Western blot analysis and **(C)** RT-qPCR, respectively. *** $P < 0.001$ vs HIEC. **(D)** RT-qPCR and **(E)** Western blot analyses (40 μg protein sample per lane) were employed to evaluate the level of ELOVL6 after treatment of apatinib. * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$ vs control.

Abbreviations: ELOVL6, elongation of very long-chain fatty acids family member 6; CRC, colorectal cancer.

from apatinib treatment was reversed by ELOVL6 upregulation. Subsequently, STRING database was applied to predict the potential proteins interacting with ELOVL6, which illustrated that ACSL4, a vital regulator of ferroptosis, could interact directly with ELOVL6 (Figure 5E). Further, Co-IP assay explored their binding affinity, proving the strong interaction between ELOVL6 and ACSL4 (Figure 5F). Through the above findings, we conclude that ELOVL6 elevation abolishes the regulatory impacts of apatinib on the expression of ferroptosis-associated proteins via ACSL4.

Discussion

Existing studies have indicated that apatinib inhibits the progression of CRC via promoting apoptosis. However, the detailed mechanisms underlying apatinib-induced cell death require further exploration. Among the mechanisms accounting for cell death, ferroptosis is a newly identified and unique category of regulated cell death.¹⁹ Ferroptosis

is featured with iron dependence and lipid peroxidation production, which makes it distinguished from other kinds of cell death such as apoptosis and necrosis.²⁰ Numerous studies have confirmed that ferroptosis plays a crucial role in killing tumor cells and inhibiting tumor growth in multiple cancers. For instance, ferroptosis activation suppresses CRC cell growth and tumorigenesis.²¹ In the present study, we demonstrated that apatinib treatment led to a decrease in cell viability and an increase in ferroptosis at least partly via targeting ELOVL6/ACSL4 signaling in HCT116 cells.

Iron accumulation and lipid ROS production are two critical events in ferroptosis, which is characterized by high intracellular Fe^{2+} concentration and ROS level.¹⁸ The decrease in total iron, Fe^{2+} and ROS levels has been widely reported in diverse cancers.^{22,23} Significant increases in intracellular iron content, ferrous iron concentration and ROS level were revealed in HCT116 cells after apatinib intervention. As is known to us all, ferroptosis is linked to a fatal accumulation of lipid peroxidation, evoking

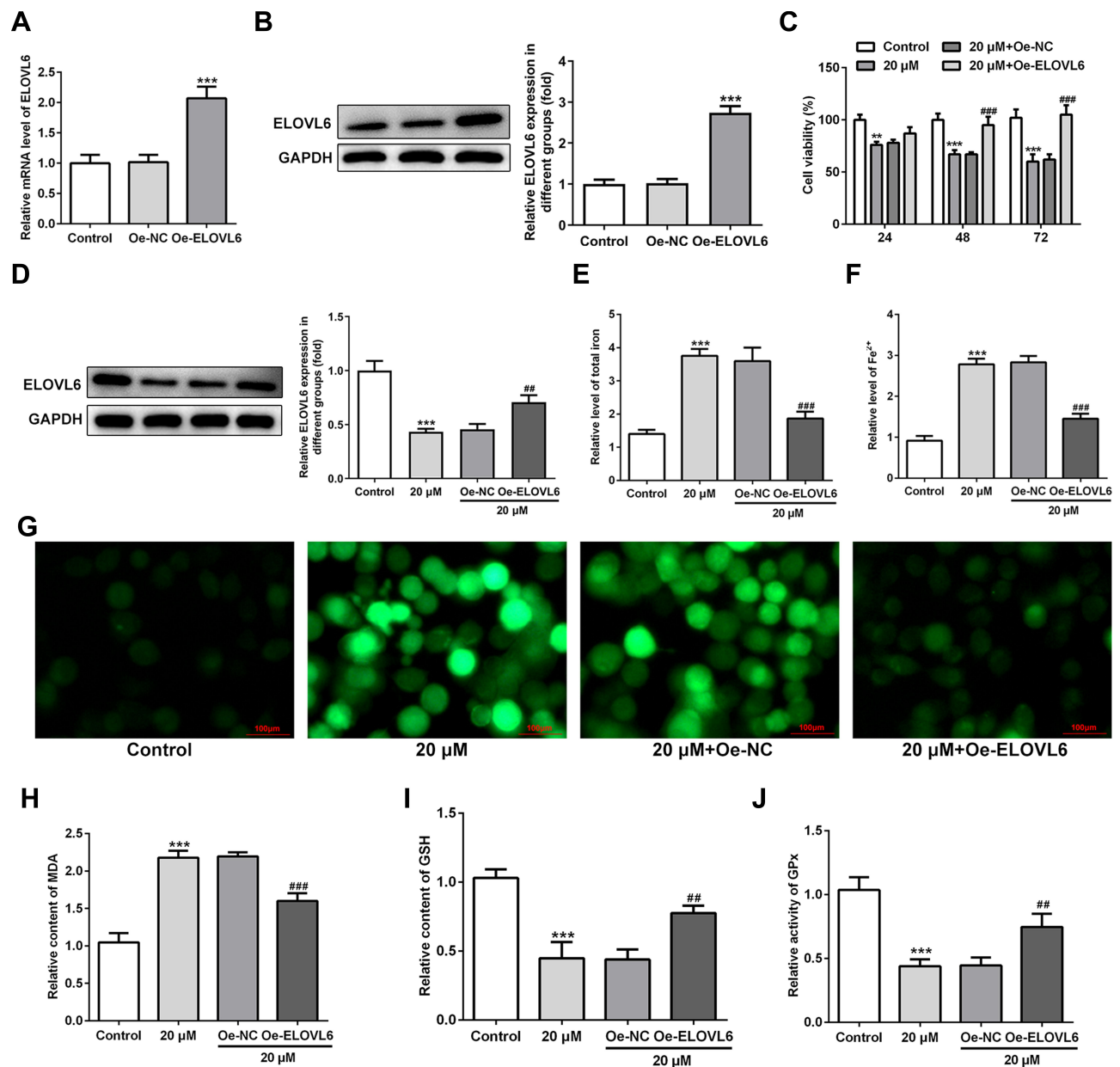


Figure 4 ELOVL6 upregulation reverses the effects of apatinib on cell viability and ferroptosis. (A and B) ELOVL6 overexpression was examined, respectively, using RT-qPCR and Western blot analysis after transfection. *** $P < 0.001$ vs Oe-NC. (C) Cell viability of HCT116 cells was determined using a cell counting kit-8 assay kit. (D) The protein level of ELOVL6 after apatinib treatment upon transfection was examined with Western blotting. The relative concentrations of (E) total iron and (F) Fe^{2+} were detected using respective kits. (G) ROS level was detected by a ROS assay kit with 2, 7-dichlorofluorescein diacetate (DCFH-DA, Invitrogen, USA). The levels of (H) MDA, (I) GSH and (J) GPx were evaluated using commercial kits. ** $P < 0.01$ and *** $P < 0.001$ vs control; ### $P < 0.01$, #### $P < 0.001$ vs Oe-NC.

Abbreviations: ELOVL6, elongation of very long-chain fatty acids family member 6; ROS, reactive oxygen species; MDA, malondialdehyde; GSH, glutathione; GPx, glutathione peroxidase.

membrane rupture and cell content release.²⁴ MDA, a representative end-product of lipid peroxidation, contributes to cell membrane injury and then ferroptosis.⁶ Moreover, the GSH antioxidant system is recognized as a major cell-protection system, preventing ferroptotic cell death.²⁵ Furthermore, GPx is a unique member of glutathione peroxidases that regulates the ferroptosis to protect cells against detrimental lipid peroxidation.²⁶ The present

study unveiled that apatinib notably enhanced the levels of ROS and MDA, accompanied by decreases in GSH content and GPx activity, suggesting the activation of ferroptosis after apatinib exposure in HCT116 cells. It has been documented that excessive iron involves in lipid peroxidation through multiple mechanisms.²⁷ ACSL4, a limiting enzyme for catalyzing the conversion of long-chain fatty acids to acyl-coenzyme, plays a crucial role in the process of iron-

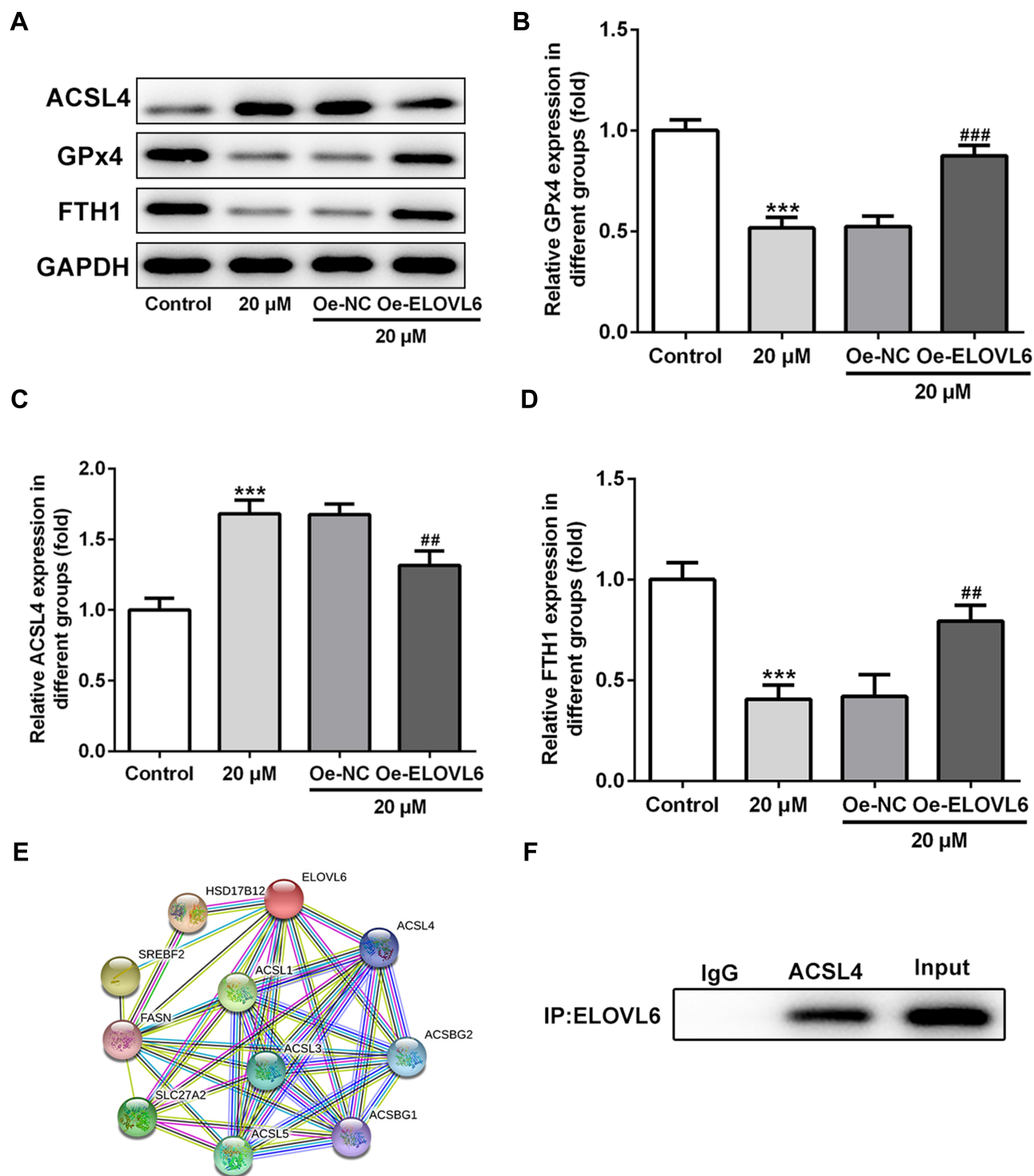


Figure 5 ELOVL6 overexpression restores the regulatory effects of apatinib on the expression of ferroptosis-associated proteins via ACSL4. (A–D) The expression of ACSL4, GPx4 and FTH1 was determined using Western blot analysis. *** P <0.001 vs control; ### P <0.01, #### P <0.001 vs Oe-NC. (E) The protein–protein interaction network for ELOVL6. (F) Immunoprecipitation was performed with indicated antibodies in HCT116 cells. The immunocomplexes were subjected to Western blot assay.

Abbreviations: ELOVL6, elongation of very long-chain fatty acids family member 6; ACSL4, long-chain acyl-CoA synthetases 4; GPx4, glutathione peroxidase 4; FTH1, ferritin heavy chain.

dependent oxidative stress.^{28,29} GPx4, a significant antioxidant enzyme in mammals, has been proved to perform a suppressive role in lipid ROS production during

ferroptotic cell death.³⁰ Besides, the inhibitory effect of GSH on ferroptosis is mainly attributed to the activation of GPx4.²⁶ Moreover, FTH1 is an antioxidant that can cause

the resistance to ferroptosis.³¹ In the current study, significant upregulation of ACSL4 and downregulation of GPx4 and FTH1 were viewed in apatinib-treated HCT116 cells, affirming the regulatory effects of apatinib on ferroptosis-related proteins.

To fully investigate the underlying mechanisms of apatinib in ferroptosis of HCT116 cells, the SwissTargetPrediction website (<http://www.swisstargetprediction.ch/>) was employed, the result of which delineated that ELOVL6 was targeted by apatinib. ELOVL6 was reported to be a tumor suppressor gene or an oncogene in various tumors. For instance, in hepatocellular carcinoma, low level of ELOVL6 indicates poor prognosis.³² Inhibition of ELOVL6 potentiates upregulation of genes related to breast cancer development and metastasis in vivo.³³ Additionally, in a study investigating pediatric primary central nervous system germ cell tumors, the expression of ELOVL6 was observed to be abundant in germinoma.³⁴ However, the GEPIA database (<http://gepia.cancer-pku.cn/>) displayed that ELOVL6 is expressed at high level in CRC tissues. Meanwhile, in the current study, the expression of ELOVL6 in mRNA and protein levels in HCT116 cells presented the same result. Therefore, whether apatinib targeted ELOVL6 to promote ferroptosis of HCT116 cells was explored. Results demonstrated that apatinib inhibited the expression of ELOVL6, and ELOVL6 overexpression reversed the effects of apatinib on ferroptosis of HCT116 cells. Additionally, to further excavate the molecular mechanisms underlying ELOVL6-mediated ferroptosis, STRING database was applied to search for the potential proteins interplaying with ELOVL6. Among the gained proteins, ACSL4, a vital regulator of ferroptosis, was identified as a biomarker and contributor of ferroptosis.²⁸ It has been reported that ACSL4 was downregulated in human glioma tissues and cells, and it could suppress glioma cells proliferation via activating ferroptosis.³⁵ Further, the interplay between ELOVL6 and ACSL4 was confirmed by co-IP assay. Thus, these findings suggest that apatinib inhibits cell viability and promotes ferroptosis in HCT116 cells at least partly via targeting ELOVL6/ACSL4 signaling.

Conclusion

In summary, the present research first discovered the effect of apatinib on the ferroptosis of CRC HCT116 cells. Mechanically, we demonstrated that apatinib boosted ferroptosis in CRC cells by targeting ELOVL6 and subsequently regulating ACSL4 expression, which provides

a new mechanism support for apatinib application in the clinical treatment of patients suffering from colorectal cancer. However, the usage of single CRC cell line is a limitation of this study. Therefore, a comprehensive analysis is required in the future.

Data Sharing Statement

The datasets used and/or analyzed during the present study are available from the corresponding author on reasonable request.

Disclosure

The authors declare that they have no competing interests.

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