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Self-supplying Cu²⁺ and H₂O₂ synergistically enhancing disulfiram-mediated melanoma chemotherapy†

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Disulfiram (DSF) can target and kill cancer cells by disrupting cellular degradation of extruded proteins and has therefore received particular attention for its tumor chemotherapeutic potential. However, the uncontrollable Cu²⁺/DSF ratio reduces the efficacy of DSF-mediated chemotherapy. Herein, self-supplying Cu²⁺ and oxidative stress synergistically enhanced DSF-mediated chemotherapy is proposed for melanoma-based on PVP-coated CuO₂ nanodots (CPNDs). Once ingested, DSF is broken down to diethyldithiocarbamate (DTC), which is delivered into a tumor *via* the circulation. Under the acidic tumor microenvironment, CPNDs produce sufficient Cu²⁺ and H₂O₂. DTC readily chelates Cu²⁺ ions to generate CuET, which shows antitumor efficacy. CuET-mediated chemotherapy can be enhanced by H₂O₂. Sufficient Cu²⁺ generation can guarantee the maximum efficacy of DSF-mediated chemotherapy. Furthermore, released Cu²⁺ can be reduced to Cu⁺ by glutathione (GSH) and O²⁻ in tumor cells, and Cu⁺ can react with H₂O₂ to generate toxic hydroxyl radicals (·OH) *via* a Fenton-like reaction, promoting the efficacy of CuET. Therefore, this study hypothesizes that employing CPNDs instead of Cu²⁺ ions could enhance DSF-mediated melanoma chemotherapy, providing a simple but efficient strategy for achieving chemotherapeutic efficacy.

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Introduction

Melanoma, one of the skin cancers with a high mortality rate, is seriously threatening to human health.^{1–8} Surgery shows very limited efficacy for advanced melanoma, and classic treatments such as chemotherapy and targeted therapy are very costly and have side effects.^{9–11} Herein, a new treatment is proposed for advanced melanoma using clinically approved “old drugs”,^{12,13} with reduced clinical trial time and high biosafety.

Disulfiram (tetraethylthiuram disulfide, DSF), a drug approved by the Food and Drug Administration (FDA) for chronic alcoholism, has aroused great interest in the field of tumor chemotherapy.^{14–17} Once ingested, DSF can be metabolized to DTC, which chelates Cu²⁺ ions to form CuET, which shows high cytotoxicity. CuET induces p97 segregase adapter, NPL4, aggregation in the nucleus, which triggers a heat-shock

response, resulting in further death of cells.^{18–20} Thus, Cu²⁺ ions play an important role in DSF-based chemotherapy.

Previous reports showed that under oxidative conditions, CuET can release Cu²⁺ ions, which then bind to the zinc motif of NPL4, changing the conformational structure of NPL4.¹⁹ This change leads to inhibition of ubiquitin protein metabolism and promotes further cell apoptosis. The generation of Cu²⁺ ions suggests the existence of a cycle of DSF, DTC, and CuET and excessive DTC neutralizes CuET toxicity by chelating released Cu²⁺ ions.^{19,21} Therefore, it is necessary to provide sufficient Cu²⁺ ions and to create an oxidative microenvironment to improve the therapeutic efficacy of CuET-induced chemotherapy in tumor cells. In recent years, many studies focused on DSF-induced chemotherapy have paid more attention to copper-based nanoplatfoms to increase the efficacy of such chemotherapy.^{22–24} However, in many delivery strategies, the ratio between the released Cu²⁺ ions and DSF is uncontrollable, which affects the efficiency of DSF-based chemotherapy. The therapeutic efficacy of DSF-based chemotherapy is only enhanced when the concentration of Cu²⁺ ions exceeds that of DTC in tumor cells. Meanwhile, most nanoparticle platforms are transported intravenously and preferentially accumulate in a tumor depending on the EPR effect.^{25–29} This passive targeting mechanism has very low accumulation efficiency, resulting in about 0.7% ID (median) at the tumor.²⁸ Furthermore, most nanoparticles are distributed in normal tissues during

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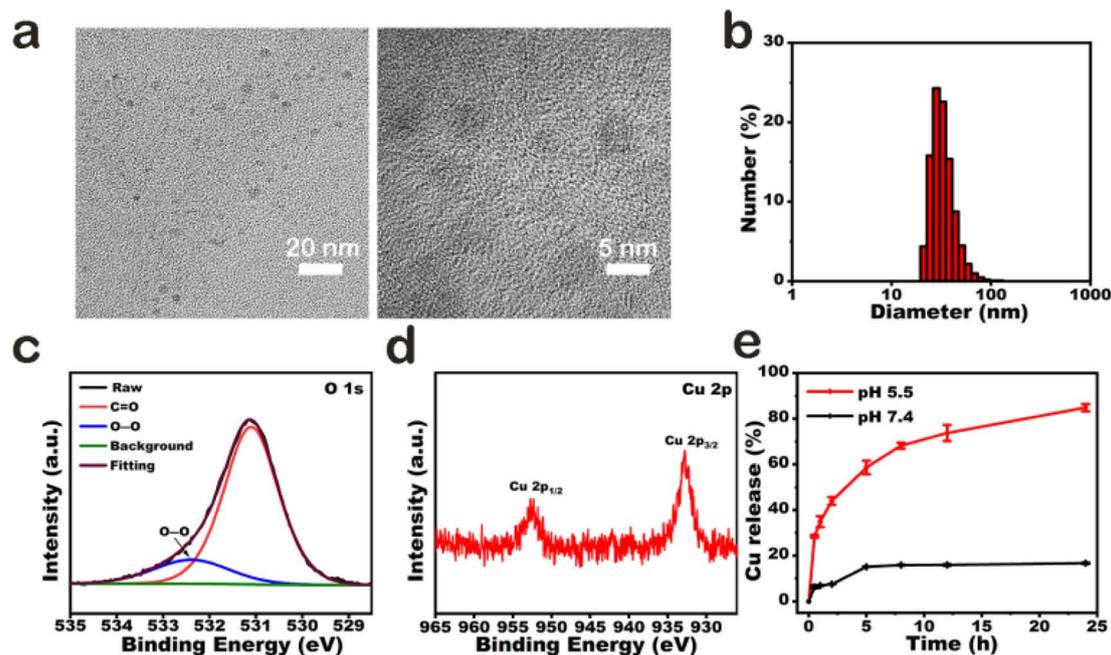



Fig. 1 (a) TEM images of CPNDs. (b) Dynamic light scattering (DLS) measurement of CPNDs. (c) High-resolution O 1s and (d) Cu 2p XPS spectra of CPNDs. (e) Cumulative Cu release from CPNDs under different pH conditions.

(TME),³⁷ CPNDs are decomposed into Cu^{2+} ions and H_2O_2 . The release of Cu^{2+} ions by CPNDs was enhanced under acidic conditions (Fig. 1e). An assay using 3,3',5,5'-tetramethylbenzidine (TMB), which can be oxidized by $\cdot\text{OH}$ with maximum

absorbance at 650 nm, was used to detect the generation of $\cdot\text{OH}$. The absorption peak at 650 nm increasing under the acidic condition indicated that the $\cdot\text{OH}$ generation of CPNDs could be pH dependent (Fig. S2†). DSF can be metabolized into DTC,

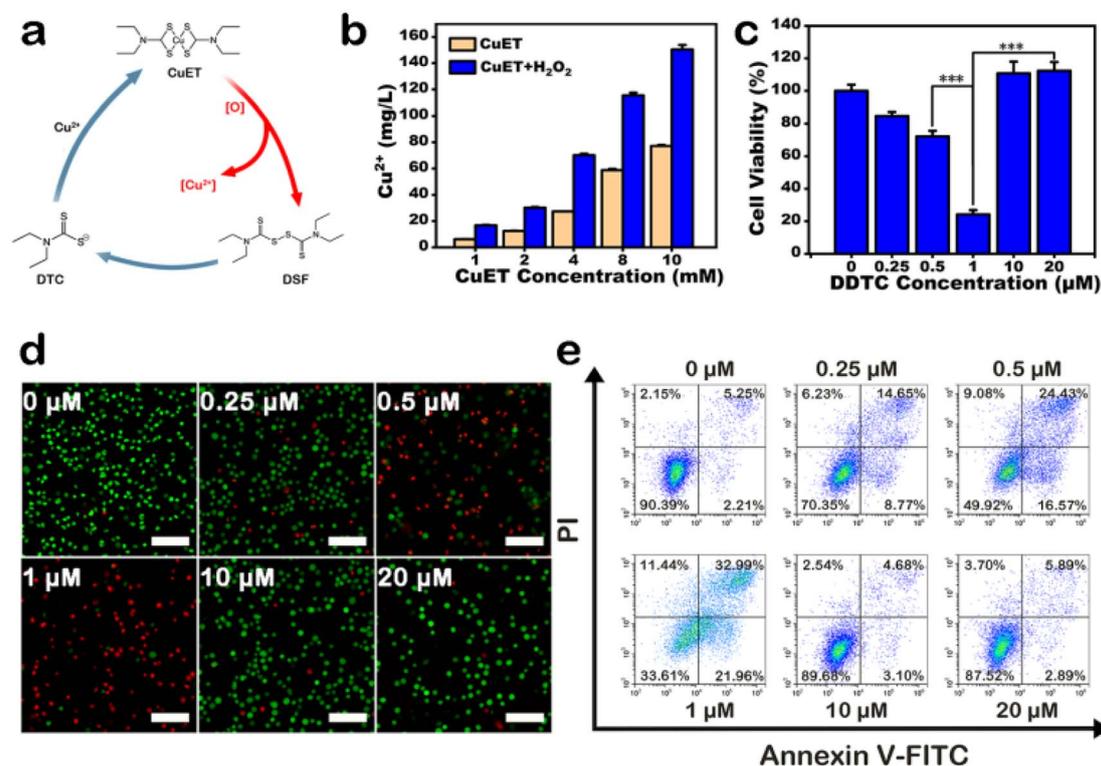


Fig. 2 (a) Scheme of CuET cycle. (b) Cu^{2+} release of CuET under oxidative conditions. Performance of CuET cycle *in vitro* via (c) CCK-8 assay, (d) live-dead cell staining assay (scale bar is 100 μm) and (e) cell apoptosis analysis.



which can chelate Cu^{2+} to form CuET,^{38–43} a complex with anti-tumor activity (Fig. S3†). To synthesize CuET *in vitro*, we used sodium diethyldithiocarbamate (DDTC) to chelate Cu^{2+} to afford CuET. The DDTC and CuCl_2 reaction mixture showed an absorption peak at ~ 444 nm,⁴⁴ which indicated the generation of CuET (Fig. S4†). To investigate the reaction between CPNDs and DDTC, DDTC was added into an aqueous solution of CPNDs at different pH. The results showed that the absorbance at ~ 444 nm increased at acidic condition, indicating that lower pH promoted CuET production and Cu^{2+} ion release (Fig. S5 and S6†).

Under oxidative conditions, CuET has been shown to decompose into Cu^{2+} ions, which interact with zinc finger motifs in NPL4,¹⁹ which is involved in cellular protein metabolism. First, we synthesized a solution of CuET by reacting DDTC and CuCl_2 , with H_2O_2 as an oxidative reagent. After reacting for 24 h, the solution was centrifuged, and the supernatant was collected to detect Cu^{2+} ion content using ICP-MS (Fig. 2b and S7†). The results showed that the concentration

of Cu^{2+} ions was higher in the CuET solution with H_2O_2 , which indicated that oxidative conditions accelerated CuET decomposition into Cu^{2+} ions. In addition, a cycle of DSF, DTC, and Cu^{2+} was observed in cells containing CuET, which could increase the cellular Cu^{2+} ion level and bypass the copper transporter system (Fig. 2a). To confirm the existence of this cycle, the cytotoxicity of DDTC at different concentrations (0, 0.25, 0.5, 1, 10, and 20 μM) with 1 μM CuCl_2 was investigated. Cell viability decreased with increasing DDTC concentration until 1 μM (Fig. 2c). However, at 10 μM and 20 μM DDTC, cell viability was 110.8% and 112.3%, respectively, compared with 24.2% at 1 μM DDTC. This result indicated that excessive DDTC rescued the damage to the structure of NPL4 by chelating Cu^{2+} ions released from CuET. To directly visualize live or dead cells after treatment with different concentrations of DDTC with CuCl_2 , calcein-AM/propidium iodide (PI) fluorescent dyes were used to label A375 cells. The result showed that dead cells were fewer when concentrations of DDTC were much higher than that of Cu^{2+} ions, which was consistent with the CCK-8 assay

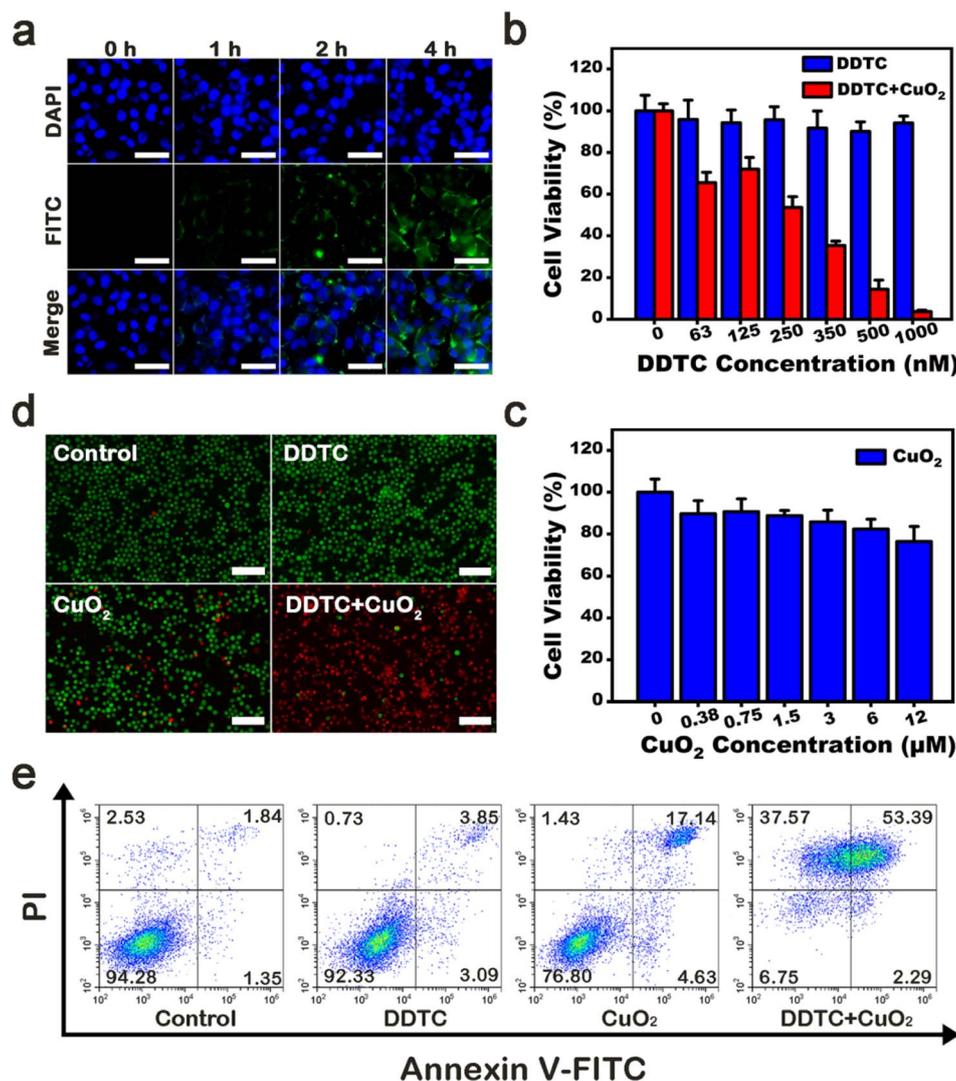


Fig. 3 (a) Cellular uptake assay of CPNDs; scale bar is 50 μm . (b) CCK-8 assay of DDTC, (c) CPNDs and combined (d) live-dead cell staining images; scale bar is 100 μm . (e) Cell apoptosis analysis images of A375 cells after different treatments.

(Fig. 2d). This conclusion was further confirmed by apoptosis assay using flow cytometry (Fig. 2e). In addition, the assay indicated that CuET induced apoptotic A375 cell death. In conclusion, to improve the efficiency of CuET-induced chemotherapy, in addition to a suitable amount of DSF, excess Cu^{2+} ions and oxidative conditions are very important.

CPNDs were endocytosed into tumor cells and later rapidly decomposed in the acidic environment of endo/lysosomes.⁴⁵ First, we used fluorescence microscopy to observe the efficiency of intracellular uptake of FITC-labeled CPNDs. After co-incubation for 4 h, enhanced green fluorescence signals were seen around the nucleus, which indicated that most CPNDs were localized in the cytoplasm (Fig. 3a). This result was confirmed by quantitative flow cytometry evaluation (Fig. S8†). Then, the synergistic effect of DDTC combined with CPNDs and cytotoxicity of DDTC at different concentrations (0, 63, 125, 250, 350, 500, and 1000 nM) with or without CPNDs were investigated. To ensure sufficient Cu^{2+} ions, the concentration of the Cu^{2+} donor, CPNDs, was 8 μM .⁴⁶ Experimental results depicted that the viability of A375 cells remained high after incubation with free DDTC. In contrast, the

viability of cells treated with DDTC combined with CPNDs decreased, which could be ascribed to CuET-induced chemotherapy and CPND-induced CDT (Fig. 3b). In addition, the viability of cells treated with CPNDs at action concentrations remained high, which indicated that the cytotoxicity of CuET-induced chemotherapy was much stronger than that of CPND-induced CDT (Fig. 3c). To directly observe the efficiency of these agents, calcein-AM/propidium iodide (PI) fluorescent dyes were used to label A375 cells (Fig. 3d). The results showed that cells in the control group and DDTC-treated cells displayed strong green fluorescence, indicating that they were living cells. In contrast, cells in the CPND group showed little red fluorescence, and cells in the DDTC group combined with the CPND group showed strong red fluorescence, indicating that they were dead cells. These observations indicated that treatment with DDTC combined with CPNDs had a significant therapeutic effect on A375 cells. Furthermore, an apoptosis assay was performed using Annexin V-FITC/PI staining and flow cytometry. The results showed that the combined treatment of DDTC and CPNDs could increase the apoptosis of A375 cells (Fig. 3e).

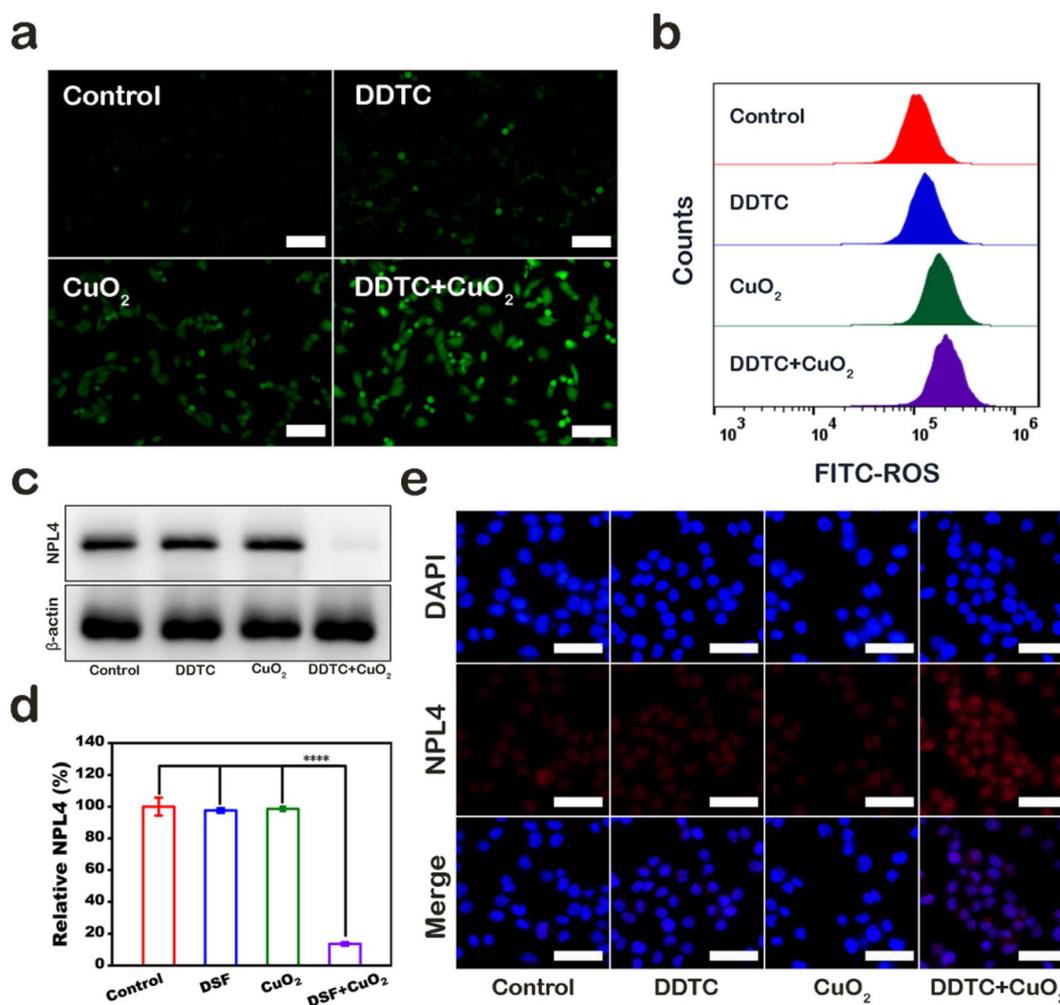


Fig. 4 (a) DCFH-DA fluorescence images of A375 cells; scale bar is 100 μm . (b) Flow cytometry demonstrating cell phagocytosis of CPNDs. (c) Western blot assay of NPL4 expression in A375 cells after different treatments. (d) Quantitative results of the expression of NPL4. (e) NPL4 aggregation was visualized by immunofluorescence staining after different treatments; scale bar is 50 μm .



The mechanism of A375 cell damage was also explored. First, we used 2',7'-dichlorofluorescein diacetate (DCFH-DA), which can be oxidized by ROS into green fluorescent 2',7'-dichlorofluorescein (DCF), to detect the level of ROS in A375 cells after treatment. Cells in control, DDTC, CPND group, and combined treatment groups showed no fluorescence, little green fluorescence, and strong green fluorescence, respectively (Fig. 4a). Furthermore, the result of the quantitative flow cytometry evaluation was consistent with fluorescence patterns (Fig. 4b). Melanoma cells were shown to be highly sensitive to DSF, which could increase the level of intracellular ROS. In addition, CPNDs could increase the ROS level in cells *via* CDT. Our results demonstrated that the combined treatment caused cell damage by inducing high ROS levels, which could be attributed to the

synergistic effect of CPNDs and DDTC. DTC, a metabolite of DSF, has been shown to chelate Cu^{2+} to form CuET. Under an oxidative environment, CuET could be decomposed into Cu^{2+} , which binds and immobilizes NPL4, an adaptor of p97 segregase, inhibiting protein metabolism. First, we extracted the cytoplasmic protein of A375 cells and measured NPL4 in the cytoplasm. The results showed that the expression of NPL4 decreased after the combined treatment (Fig. 4c and d). Then, we used immunofluorescence staining to evaluate the NPL4 protein status. The result showed that the NPL4 protein aggregated in the nucleus after treatment with DDTC combined with CPNDs (Fig. 4e).

Encouraged by the therapeutic efficacy *in vitro*, we further evaluated the therapeutic effect of DSF combined with CPNDs on A375 tumor-bearing mice (Fig. 5a). The mice were divided into

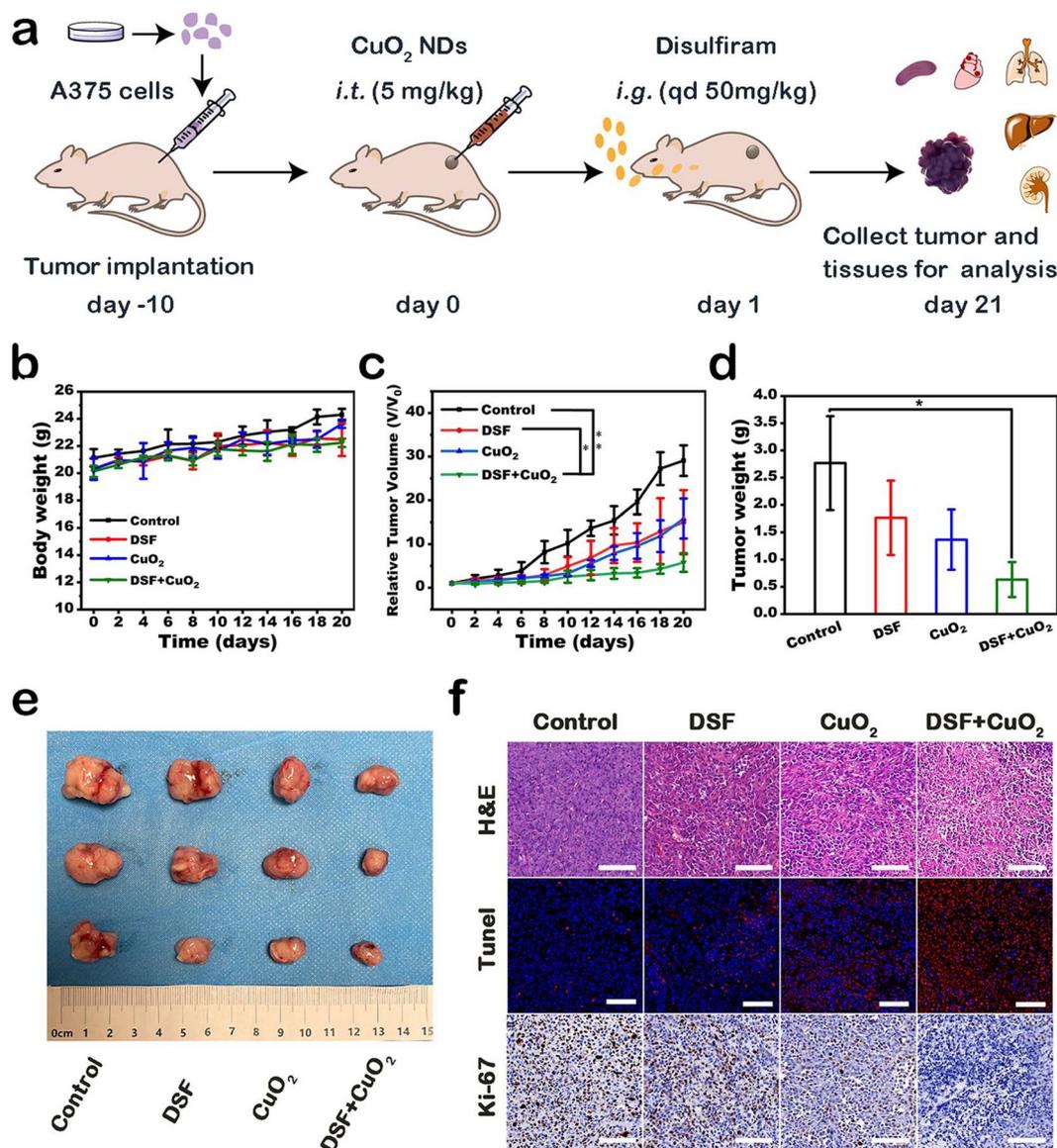


Fig. 5 (a) Schematic illustration of A375 subcutaneous tumor-bearing mice model development and treatment process. (b) Body weight, (c) relative tumor volume and (d) tumor weight change curves of A375 tumor-bearing female Balb/c nude mice after different treatments. (e) Digital photographs of excised tumors after different treatments. (f) H&E, TUNEL, and Ki-67 antigen staining of tumor tissues after different treatments; scale bar is 100 μm .



four groups: control, DSF, CuO₂, and DSF + CuO₂. After treatment, the inhibition rate in the DSF group and CuO₂ group was 48.5% and 45.6%, respectively. Comparatively, the inhibition rate of the combined treatment group was 80.1% (Fig. 5c). These results indicated that DSF combined with CuO₂ had better synergetic efficacy than the other groups. Tumor weights and corresponding digital pictures were directly consistent with these results (Fig. 5d and e). In addition, body weights (Fig. 5b) and major organs (heart, liver, spleen, lung, and kidney) of the mice showed no obvious changes and damage, which indicated that the treatments induced no significant side effects (Fig. S9†).

Hematoxylin–eosin (H&E) staining, Ki-67 immunohistochemistry staining, and TUNEL fluorescence staining were performed to show histological changes and cell inhibition in tumor sections (Fig. 5f). Remarkably, based on H&E staining results, the largest area of tissue necrosis was in the DSF + CuO₂ group. TUNEL staining and Ki-67 staining indicated significant tumor cell apoptosis and suppression of tumor cell proliferation, respectively. In general, these results provided show that the therapeutic strategy combining DSF with CPNDs exhibited good tumor inhibition efficacy.

Conclusion

Previous studies report that DSF-based chemotherapy still suffers from low efficacy and biosafety. In this work, exploiting the drug delivery mechanisms of nanomedicine, we constructed a novel treatment scheme for improving the efficacy of DSF-based chemotherapy and demonstrated its biosecurity *via* oral uptake of DSF orally and intratumoral injection of CPNDs to treat melanoma. In the acidic TME, CPNDs dissolved in saline can be broken up into Cu²⁺ ions and H₂O₂, and the released Cu²⁺ ions are reduced to Cu⁺ by GSH and O₂^{•−} in tumor cells, and Cu⁺ readily reacts with H₂O₂ to generate ·OH *via* a Fenton-like reaction. This smart DSF/Cu²⁺ delivery strategy effectively inhibited tumor growth by 80.1% *in vivo* through highly effective DSF-based chemotherapy and highly toxic ·OH *via* CPND-induced Fenton-like reaction. This scheme provides sufficient Cu²⁺ ions and oxidative condition for improving the efficacy of DSF-based chemotherapy. Meanwhile, the therapeutic schedule conforms to the principles of clinical medication, providing a new idea for the clinical treatment of melanoma. One limitation of this study is that the permeability and metabolism of nanoparticles *via* intratumoral injection were not studied. Therefore, it is necessary to explore the mechanism of intratumoral injection of nanoparticles to perfect this treatment scheme.

Experimental section

Synthesis and characterization of CPNDs

CPNDs were synthesized on the basis of a reported method.⁴² First, 0.0085 g of CuCl₂·2H₂O and 0.5 g of PVP were added to deionized water (5 mL) to form a homogeneous solution by magnetic stirring. Next, 5 mL of deionized water containing NaOH (0.004 g) was added to the above solution and continuously stirred to form a green solution. Subsequently, H₂O₂ (100

μL) was added to the mixed solution and stirred for 30 min. The resultant CPNDs were obtained by ultrafiltration and washed three times.

TEM imaging was carried out with a JEOL JEM-2100F transmission electron microscope. Hydrodynamic particle size distribution was determined using dynamic light scattering (DLS) (Zetasizer Nano ZS90; Malvern Instruments, Malvern, UK). XPS analysis was performed with a K-Alpha instrument (Thermo Fisher Scientific, Waltham, MA, USA). The content of copper was determined using inductively coupled plasma mass spectrometry (ICP-MS) (Agilent 5110, USA). UV/visible absorption spectra were measured with a Lambda-750 spectrometer (PerkinElmer, Shanghai, China).

pH-responsive decomposition of CPNDs

The acidic-induced dissociation of CPNDs was evaluated by ICP-MS. Briefly, CPNDs were dissolved in buffer solutions with different pH values (7.4 or 5.5). Then, dialysates were withdrawn from each sample at given times (0.5, 1, 2, 5, 8, 12, and 24 h), and released Cu²⁺ was determined by ICP-MS.

Synthesis of CuET

To prepare CuET, DDTC aqueous solution (1.5 mL and 1 mM) was slowly added into 1.5 mL of CuCl₂ at different concentrations (20, 10, 1, 0.1, and 0.05 mM). After that, to prove the generation of CuET that has a characteristic absorption peak at ~444 nm, the UV/visible absorption spectra of mixed solutions were measured.

pH-dependent reaction between DDTC and CPNDs

To verify that the amount of CuET, produced by the reaction between DDTC and CPNDs, is pH-dependent, 1.5 mL of CPNDs (1 mM) was mixed for 2 h at different pH values (7.4 or 5.0). Next, 1.5 mL of DDTC aqueous solution (1 mM) was slowly added to the above solution. Then, after mixing for 30 min, the generation of CuET was determined by the absorption increase at ~444 nm.

Cu²⁺ release from CuET under oxidative condition

First, 0.5 mL of CuCl₂ aqueous solution was mixed with 1 mL of DDTC aqueous solution to prepare CuET. The concentrations of the above two solutions were 1 mM, 2 mM, 4 mM, 8 mM, and 10 mM. Each concentration of the mixed solution was prepared in duplicate. Then, 200 μL of H₂O₂ was added to one group of the solutions. After 24 h, two groups of the solutions were centrifuged, and 1 mL of supernatant of each sample was collected. Finally, the supernatant was diluted with 4 mL of concentrated nitric acid and later analyzed for copper content using ICP-MS.

Cell culture

A human melanoma cell line (A375 cells) was purchased from the Cell Bank of the Chinese Academy of Sciences (Shanghai, China). A375 cells were seeded in DMEM supplemented with 10% FBS, penicillin (100 μg mL^{−1}), and streptomycin (100 μg mL^{−1}) at 37 °C and in 5% CO₂ atmosphere.



Performance of CuET cycle

The performance of the CuET cycle was verified by CCK-8 assay, live-dead cell staining assay, and cell apoptosis analysis. First, A375 cells were seeded in 96-well plates (1×10^4 per well). Then, the cells were incubated with varying doses of DDTC (0, 0.25, 0.5, 1, 10, and 20 μM) and 1 μM of $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$. After co-culturing for 24 h, the cell viability was determined by the CCK-8 assay.

To observe the live and dead cells, the cells were seeded in 6-well plates (1×10^5 per well) and treated as described above for 24 h. After that, the cells were stained by Calcein-AM (2 μM) and PI (4.5 μM) for 15 min and visualized by a fluorescence microscope (Nikon Corporation, Japan). For cell apoptosis analysis, after being treated as described above, the cells were incubated with FITC (5 μL) and PI (10 μL) in Annexin-binding buffer (195 μL) for 15 min and further analyzed using a flow cytometer (CytoFLEX, Beckman, USA).

Cellular uptake of CPNDs

Briefly, 1 mg of FITC was dissolved in 10 mL of DMSO, and then, 1 mL of the above solution was mixed with CPND aqueous solution and stirred for 12 h. FITC- CuO_2 nanodots were obtained by ultrafiltration and washed three times.

A375 cells were seeded in confocal dishes (1×10^5 per well) and treated with the FITC- CuO_2 nanodots (1 mL, 8 μM) at various times (0, 1, 2, and 4 h). For fluorescence microscopy observation, nuclei were stained with DAPI for 5 min, followed by washing three times and visualization with fluorescence microscopy. In addition, flow cytometry analysis was used to obtain the quantitative data of FITC reflecting cell phagocytosis.

In vitro anti-cancer efficacy of DDTC and CPNDs

A375 cells were seeded in 96-well plates (1×10^4 per well). Further, the cells were treated with 100 μL of various doses of CPNDs (0, 0.38, 0.75, 1.5, 3, 6, and 12 μM) and DDTC (0, 63, 125, 250, 350, 500, and 1000 nM) for 24 h. To investigate the cytotoxicity of the combination of CPNDs and DDTC, the cells were co-cultured with 50 μL of CPNDs (8 μM) and 50 μL of DDTC (0, 63, 125, 250, 350, 500, and 1000 nM) for 24 h. The CCK-8 assay was carried out to determine the cell viability. Subsequently, live-dead cell staining assay and cell apoptosis analysis were used to verify the therapeutic effects of the combination of CPNDs and DDTC on A375 cells.

In vitro ROS generation

A375 cells were treated with DDTC (1 μM), CPNDs (8 μM), and a combination of the two for 2 h. Then, the cells were stained with DCFH-DA (10 μM) for 20 min, and fluorescence images were recorded. Further, flow cytometry analysis was used to obtain the quantitative data of ROS generation.

Measurement of NPL4 by western blot assay

Western blot assay was used to determine the expression of NPL4 in the cytoplasm of A375 cells after different treatments. First, A375 cells were seeded in a 6-well plate (1×10^5 per well)

for 24 h. DDTC (1 μM), CPNDs (8 μM), and a combination of the two were then added into cells and incubated for 4 h, and proteins were lysed by RIPA lysis buffer. A BCA Protein Quantification Kit was used to determine the total protein. Next, the proteins were separated by sodium dodecylsulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and followed by transferring onto a polyvinylidene fluoride (PVDF) membrane. Further, the membrane was incubated with the primary antibody NPL4 (1 : 1000) at 4 $^\circ\text{C}$ overnight. After that, the membrane was incubated with a relevant secondary antibody (1 : 1000) for 1 h. Finally, the membrane was visualized on an enhanced chemiluminescence imaging system (ChemiScope 6100, Shanghai, China), and band intensities were quantified using ImageJ software.

Immunofluorescence staining

A375 cells were seeded in confocal dishes (1×10^5 per well) and treated with compounds at above-mentioned concentrations for 2 h. Subsequently, the cells were fixed with 4% methanol for 10 min at room temperature, washed with PBS, and permeabilized with 0.5% Triton X-100 for 5 min. Then, the cells were incubated in 1% BSA in 0.1% PBS-Tween for 1 h to block non-specific protein-protein interactions. Next, the cells were incubated with the primary antibody NPL4 (1 : 200) for 2 h at room temperature, followed by PBS washes and staining with fluorescently conjugated secondary antibody for 60 min at room temperature. The nuclei were stained with DAPI for 5 min, followed by washing three times and visualized with fluorescence microscopy.

Animal and tumor model

BALB/c nude mice (4–6 weeks) were purchased from Changzhou Cavens Experimental Animal Co. Ltd (Changzhou, China). All animal experiments were conducted according to protocols approved by the Institutional Animal Care and Use Committee (IACUC) of Nanjing Medical University (Animal Welfare Ethics acceptance number no. IACUC-2110015). To build the melanoma tumor model, 100 μL of A375 cells (1×10^7 cells) in PBS was injected into the subcutaneous space of the hind leg. When the tumor volume reached about 60–80 mm^3 , the mice were used for further examination.

In vivo anti-cancer efficacy of DSF and CPNDs

The A375 tumor-bearing mice were randomly assigned to four groups ($n = 3$ per group), and the treatment was set as follows. Group 1: saline; group 2: DSF (50 mg kg^{-1}); group 3: CPNDs (5 mg kg^{-1}); and group 4: DSF + CPNDs (50 mg kg^{-1} + 5 mg kg^{-1}). First, group 3 and group 4 were intratumorally injected with 50 μL of CPNDs. At 24 h postinjection, group 1 was given 50 μL of saline every day. Simultaneously, group 2 and group 4 were given 50 μL of saline containing DSF every day. During the treatment, the tumor size of each group was measured with a digital vernier caliper every other day for 21 days. The tumor volume was calculated using the following equation: volume = (horizontal \times vertical²)/2. Meanwhile, the body weight was measured. On day 21, the mice of all groups were sacrificed for further histological examination.



Statistical analysis

Statistical analysis was performed using SPSS 25 software. All data are presented as mean \pm standard deviation (SD). The statistical significance of differences was conducted with Student's two-tailed *t*-test (* $P < 0.05$; ** $P < 0.01$; and *** $P < 0.001$).

Author contributions

Yingqian Gao: conceptualization, methodology, data curation, investigation, writing – original draft, writing – review & editing. Xiaojun Cai: conceptualization, methodology, writing – review & editing. Weijuan Zou: conceptualization, methodology, data curation. Xiuzhen Tang: conceptualization, methodology, data curation. Lixian Jiang: conceptualization, methodology, data curation. Junnian Hao: conceptualization, methodology, data curation. Yuanyi Zheng: conceptualization, supervision, project administration, writing – review & editing. Xinhua Ye: writing – review & editing. Tao Ying: writing – review & editing. Ao Li: writing – review & editing.

Conflicts of interest

There are no conflicts to declare.

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