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Correction: Copper-Based Targeted Nanocatalytic Therapeutics for Non-Small Cell Lung Cancer

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Yongfei Fan and Jiao Chang have contributed equally to this work.

The original article can be found online at <https://doi.org/10.1007/s40820-025-01998-5>.

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Correction to:

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Following publication of the original article [1], the authors reported the article needed to be updated. The required corrections are detailed below.

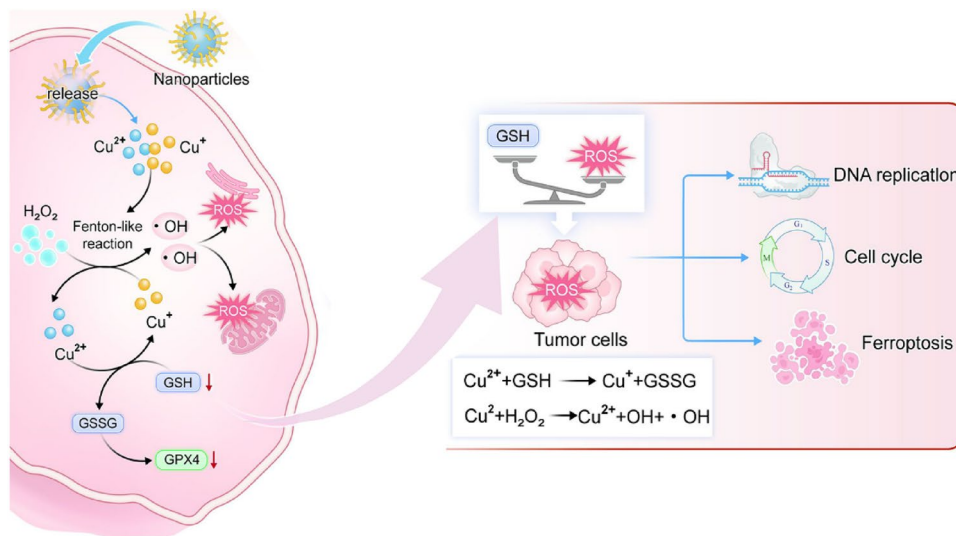
- 1) The **full name of DMSA** was incorrectly written at its first occurrence in the **Highlights, Abstract, Introduction, Fig. 1 legend, and Supplementary Information (Section S1, Experimental Section)**. The correct expression is “**copper (Cu) and dimercaptosuccinic acid (DMSA).**”

- 2) The **chemical equations in the Table of Contents graphic (TOC) and Fig. 1** were incorrectly written. The correct expressions are “ **$\text{Cu}^+ + \text{H}_2\text{O}_2 \rightarrow \text{Cu}^{2+} + \text{OH}^- + \bullet\text{OH}$** .”
- 3) Due to an error during image assembly involving copy-and-paste operations, **the statistical plot for PC-9 in Fig. 6i was incorrectly duplicated from that of NCI-H1975.**

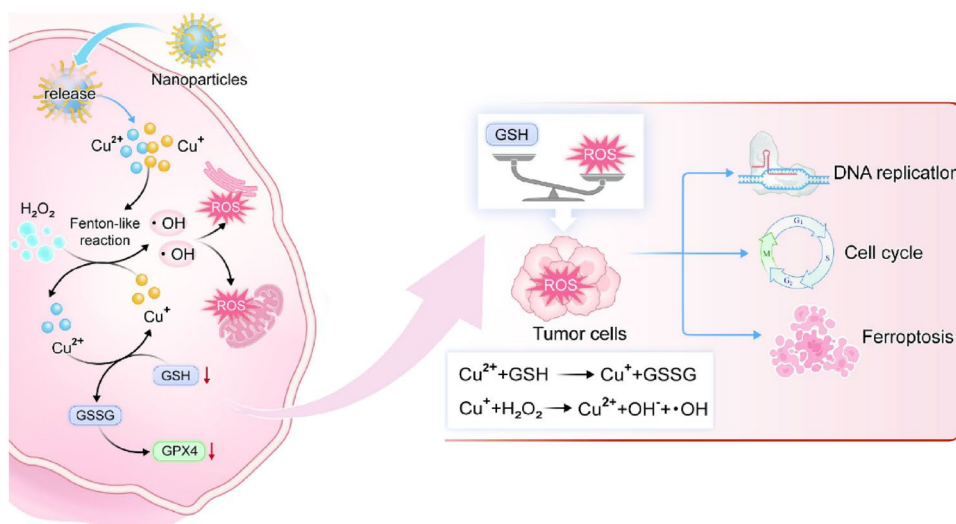
All the above corrections do not affect the final conclusions of the article. The correct Table of Contents graphic, Figs. 1 and 6 have been provided in this Correction. The Supplementary file has been updated as well.



The incorrect Table of Contents graphic is:



The correct Table of Contents graphic is:



The incorrect Fig. 1 is:

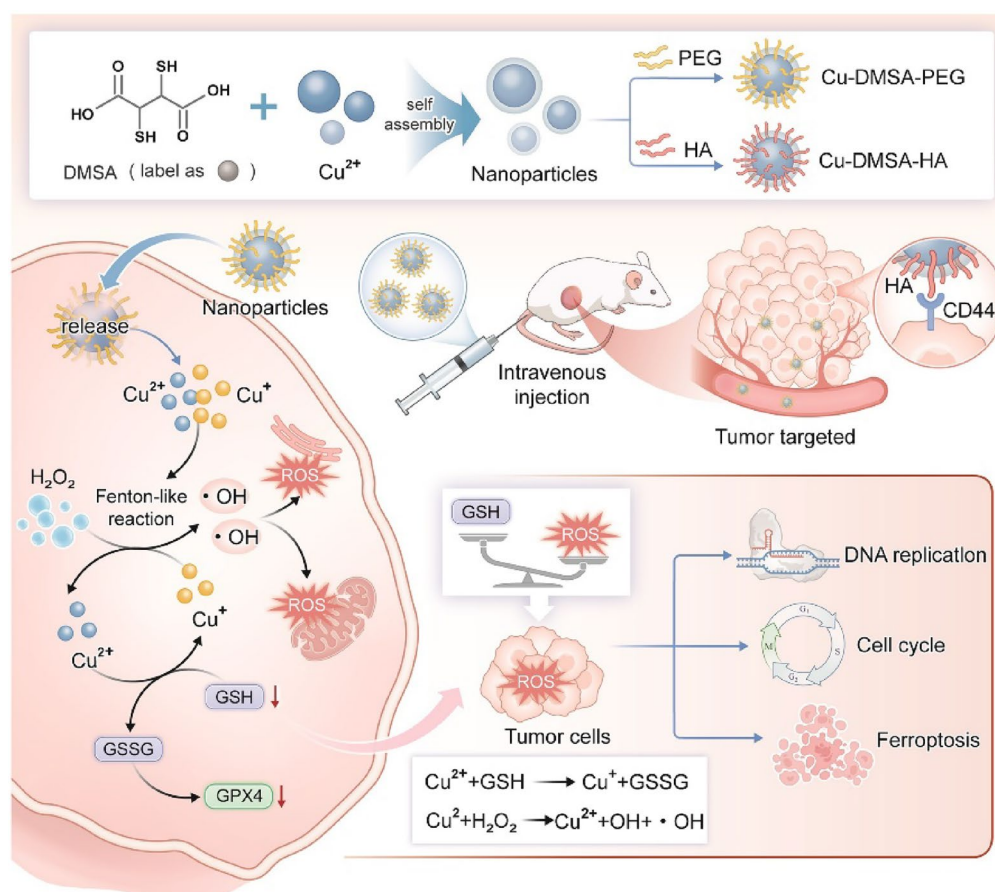


Fig. 1 Preparation schematics of nanoparticles (NPs) and proposed therapeutic mechanism. **a** Schematic illustration of the synthesis of copper-N,N-dimethyl-N-phenylsulfonylbisamine (DMSA) NPs modified with polyethylene glycol (Cu-DMSA-PEG NPs) and hyaluronic acid (HA) (Cu-DMSA-HA NPs), respectively. **b** Proposed biological mechanism of Cu-DMSA-HA-induced ferroptosis in tumor cells. Following cellular internalization, Cu-DMSA-HA NPs, which are assembled via coordination chemistry, catalyze intracellular reactive oxygen species (ROS) generation through glutathione (GSH)-mediated reduction and subsequent Fenton-like reactions. The resulting ROS accumulation triggers oxidative damage to mitochondria and the endoplasmic reticulum of tumor cells, leading to downregulation of glutathione peroxidase 4 (GPX4), a key regulator protecting against ferroptosis. **c** Mechanism of in vivo tumor targeting by Cu-DMSA-HA. Cu-DMSA-HA facilitates selective binding to cluster of differentiation 44 (CD44) receptors overexpressed on tumor cells, enhancing tumor-specific accumulation. **d** Antitumor mechanism of Cu-DMSA-HA in vivo. Cu-DMSA-HA inhibits tumor progression by suppressing DNA replication and cell cycle progression while inducing ferroptosis.

The correct Fig. 1 is:

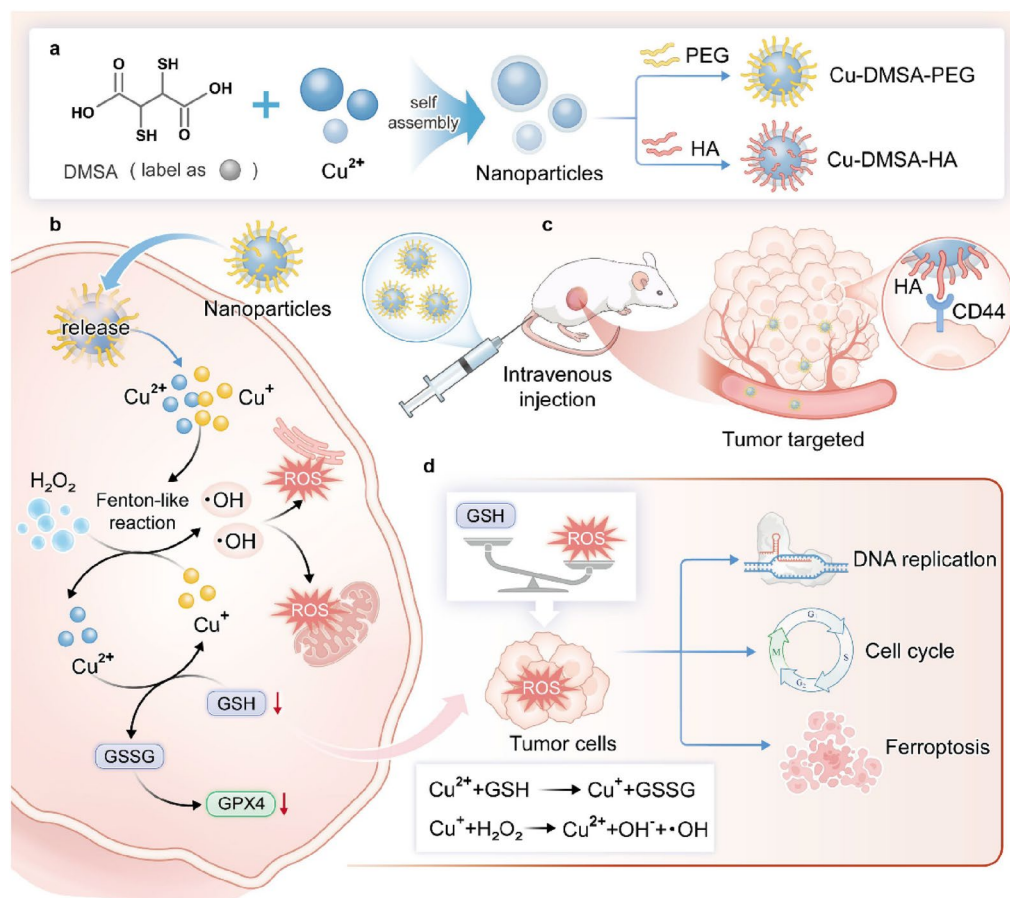
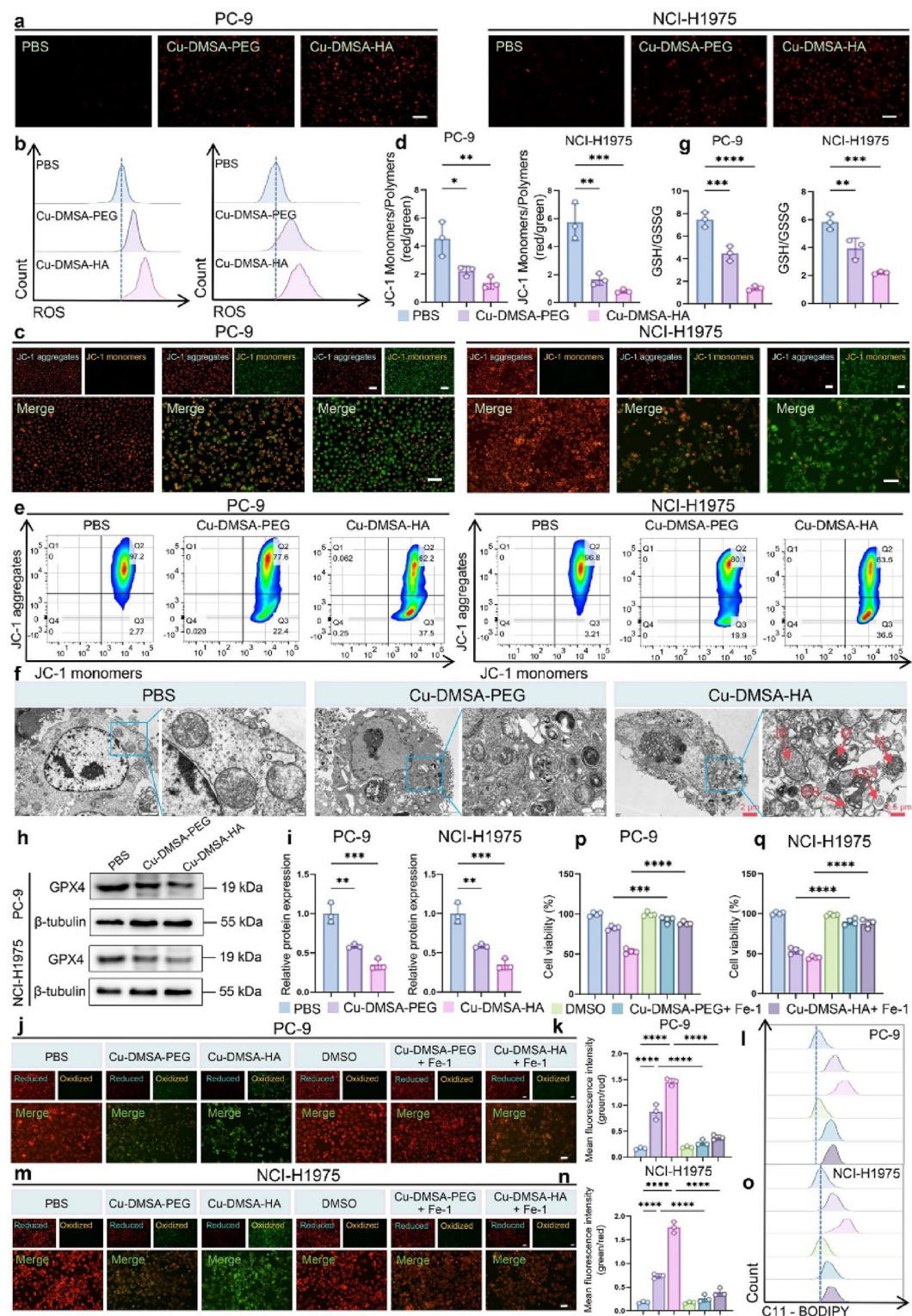


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The incorrect Fig. 6 is:

Fig. 6 Evaluation of oxidative stress and ferroptosis induction by Cu-DMSA-HA in NSCLC cells. **a** ROS staining of PC-9 and NCI-H1975 cells following treatment with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA. Scale bar: 100 μm . **b** Flow cytometry analysis of ROS levels in PC-9 and NCI-H1975 cells treated with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA. **c** Mitochondrial membrane potential ($\Delta\psi\text{m}$) staining of PC-9 and NCI-H1975 cells following treatment with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA, and **d** corresponding quantitative fluorescence intensity analyses ($n=3$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. Red fluorescence: JC-1 aggregates (healthy mitochondria, high membrane potential); green fluorescence: JC-1 monomers (damaged mitochondria, low membrane potential). Scale bar: 100 μm . **e** Flow cytometry analysis of $\Delta\psi\text{m}$ in PC-9 and NCI-H1975 cells treated with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA. **f** TEM images depicting ultrastructural changes in PC-9 and NCI-H1975 cells after treatment with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA. N: nucleus; M: mitochondria; ER: endoplasmic reticulum; ASS: autolysosome. **g** Quantification of reduced GSH and GSSG levels in PC-9 and NCI-H1975 cells following treatments with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA ($n=3$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. **h** Western blot analysis and **i** quantification of GPX4 protein expression in PC-9 and NCI-H1975 cells after treatments with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA ($n=3$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. C11-BODIPY staining and fluorescence microscopy analysis of lipid ROS accumulation in **j-k** PC-9 and **m-n** NCI H1975 cells after treatments with PBS, Cu-DMSA-PEG, Cu-DMSA-HA, DMSO, Cu-DMSA-PEG + ferrostatin-1 (Fer-1), or Cu-DMSA-HA + Fer-1 ($n=3$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. C11-BODIPY staining flow cytometry quantification of lipid ROS levels in **l** PC-9 and **o** NCI H1975 cells after treatments with PBS, Cu-DMSA-PEG, Cu-DMSA-HA, DMSO, Cu-DMSA-PEG + Fer-1, or Cu-DMSA-HA + Fer-1. CCK-8 assay of cell viability in **p** PC-9 and **q** NCIH1975 cells after treatments with DMSO, Cu-DMSA-PEG, Cu-DMSA-HA, or in combination with Fer-1 ($n=4$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. * $P<0.05$; ** $P<0.01$; *** $P<0.001$; **** $P<0.0001$.



The correct Fig. 6 is:

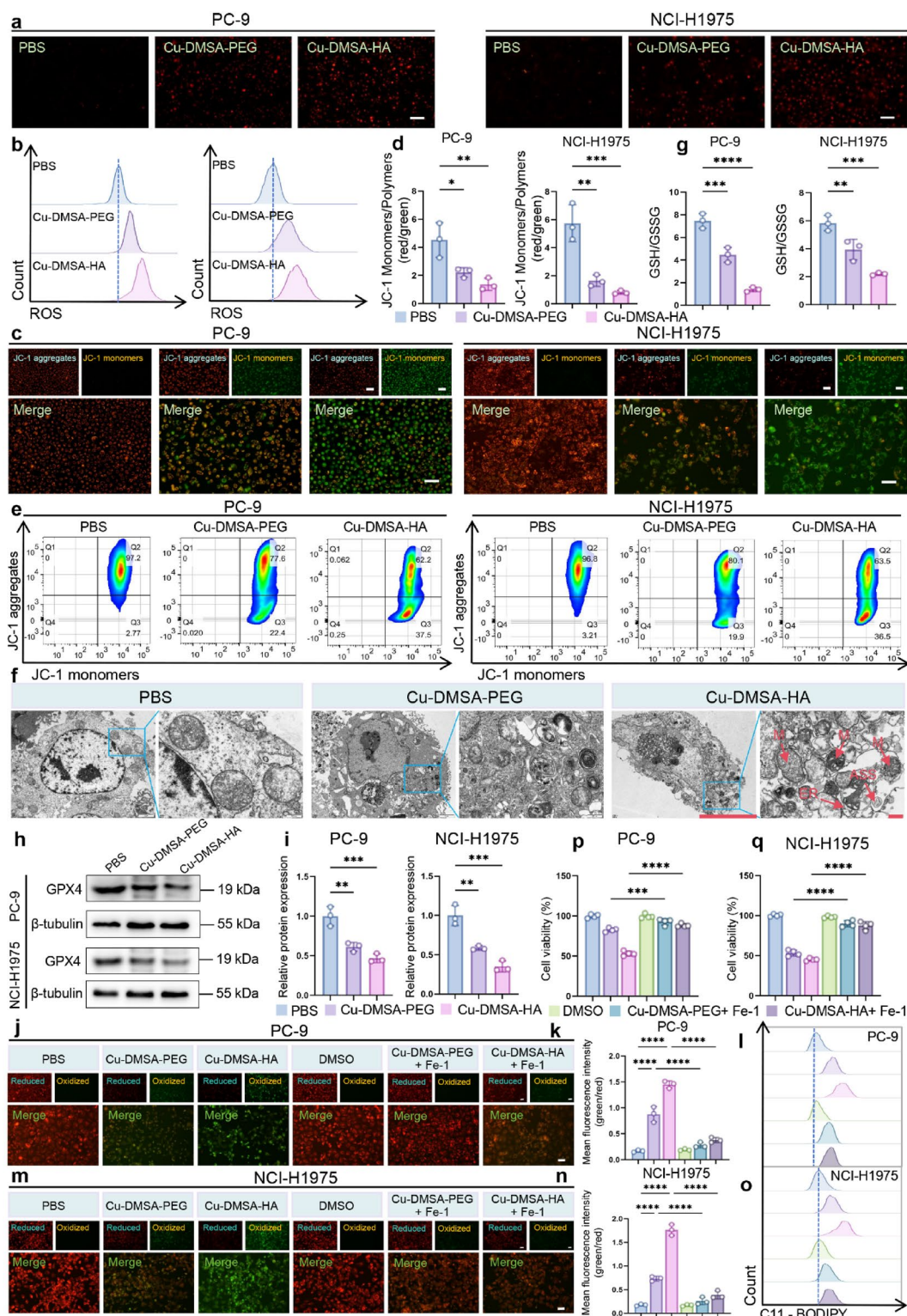


Fig. 6 Evaluation of oxidative stress and ferroptosis induction by Cu-DMSA-HA in NSCLC cells. **a** ROS staining of PC-9 and NCI-H1975 cells following treatment with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA. Scale bar: 100 μ m. **b** Flow cytometry analysis of ROS levels in PC-9 and NCI-H1975 cells treated with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA. **c** Mitochondrial membrane potential ($\Delta\psi$ m) staining of PC-9 and NCI-H1975 cells following treatment with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA, and **d** corresponding quantitative fluorescence intensity analyses ($n=3$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. Red fluorescence: JC-1 aggregates (healthy mitochondria, high membrane potential); green fluorescence: JC-1 monomers (damaged mitochondria, low membrane potential). Scale bar: 100 μ m. **e** Flow cytometry analysis of $\Delta\psi$ m in PC-9 and NCI-H1975 cells treated with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA. **f** TEM images depicting ultrastructural changes in PC-9 and NCI-H1975 cells after treatment with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA. N: nucleus; M: mitochondria; ER: endoplasmic reticulum; ASS: autolysosome. **g** Quantification of reduced GSH and GSSG levels in PC-9 and NCI-H1975 cells following treatments with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA ($n=3$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. **h** Western blot analysis and **i** quantification of GPX4 protein expression in PC-9 and NCI-H1975 cells after treatments with PBS, Cu-DMSA-PEG, or Cu-DMSA-HA ($n=3$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. C11-BODIPY staining and fluorescence microscopy analysis of lipid ROS accumulation in **j-k** PC-9 and **m-n** NCI-H1975 cells after treatments with PBS, Cu-DMSA-PEG, Cu-DMSA-HA, DMSO, Cu-DMSA-PEG + ferrostatin-1 (Fer-1), or Cu-DMSA-HA + Fer-1 ($n=3$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. C11-BODIPY staining flow cytometry quantification of lipid ROS levels in **l** PC-9 and **o** NCI-H1975 cells after treatments with PBS, Cu-DMSA-PEG, Cu-DMSA-HA, DMSO, Cu-DMSA-PEG + Fer-1, or Cu-DMSA-HA + Fer-1. CCK-8 assay of cell viability in **p** PC-9 and **q** NCI-H1975 cells after treatments with DMSO, Cu-DMSA-PEG, Cu-DMSA-HA, or in combination with Fer-1 ($n=4$; mean $\pm$ SD). Oneway ANOVA followed by Tukey's post hoc test. * $P<0.05$; ** $P<0.01$; *** $P<0.001$; **** $P<0.0001$.

The original article [1] has been corrected.

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Reference

1. Y. Fan, J. Chang, X. Qin et al., Copper-based targeted nanocatalytic therapeutics for non-small cell lung cancer. Nano-Micro Lett. **18**, 152 (2026). <https://doi.org/10.1007/s40820-025-01998-5>

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