





Ultrasmall copper-based nanoparticles for reactive oxygen species scavenging and alleviation of inflammation related diseases

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Oxidative stress is associated with many acute and chronic inflammatory diseases, yet limited treatment is currently available clinically. The development of enzyme-mimicking nanomaterials (nanozymes) with good reactive oxygen species (ROS) scavenging ability and biocompatibility is a promising way for the treatment of ROS-related inflammation. Herein we report a simple and efficient one-step development of ultrasmall Cu_{5,4}O nanoparticles (Cu_{5,4}O USNPs) with multiple enzyme-mimicking and broad-spectrum ROS scavenging ability for the treatment of ROS-related diseases. Cu_{5,4}O USNPs simultaneously possessing catalase-, superoxide dismutase-, and glutathione peroxidase-mimicking enzyme properties exhibit cytoprotective effects against ROS-mediated damage at extremely low dosage and significantly improve treatment outcomes in acute kidney injury, acute liver injury and wound healing. Meanwhile, the ultrasmall size of Cu_{5,4}O USNPs enables rapid renal clearance of the nanomaterial, guaranteeing the biocompatibility. The protective effect and good biocompatibility of Cu_{5,4}O USNPs will facilitate clinical treatment of ROS-related diseases and enable the development of next-generation nanozymes.

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Inflammation is a natural defensive response to stimulating factors such as infection, injury, and toxins^{1,2}. However, excessive and uncontrolled inflammation has been demonstrated to cause numerous diseases, such as cardiovascular disease, hepatitis, nephritis, and delayed wound healing^{3,4}. It has been widely acknowledged that inflammation is intimately associated with oxidative stress^{5,6}. Reactive oxygen species (ROS) in excessive inflammatory response could aggravate localized tissue injury and lead to chronic inflammation^{7,8}. Consequently, scavenging ROS with broad-spectrum antioxidants such as *N*-acetyl cysteine and acetyl-*L*-carnitine has been regarded as a feasible strategy for the treatment of various inflammatory diseases such as acute liver injury, liver fibrosis, acute or chronic wounds, acute kidney injury (AKI)^{9,10}. However, poor bioavailability, low stability and efficacy of these drugs limit their clinical application in the treatment of ROS-related diseases^{11,12}.

Advances in nanomedicine have enabled new ways of ROS clearance and thus treatment of ROS-related diseases using various functional nanomaterials¹³, such as carbon¹⁴, ceria¹⁵, platinum¹⁶, redox polymer¹⁷, and polyphenol nanoparticles (NPs)¹⁸. Among them, one promising strategy is to develop nanozymes to maintain natural redox balance in biological system, including catalase (CAT), superoxide dismutase (SOD), and glutathione peroxidase (GPx)¹⁹. Such nanozymes shall have high ROS scavenging ability comparable to native enzymes, broad-spectrum ROS scavenging activities against various toxic ROS species, high stability in harsh disease environment, and rapid clearance from the body to guarantee excellent biocompatibility. Therefore we believe nanomaterials with ultrasmall size (hydrodynamic diameter <5.5 nm) shall be designed because this kind of nanomaterials will have high catalytic activity due to the relatively large surface to volume ratios²⁰ and rapid renal clearance because they can pass through the glomerular capillary wall in kidney^{21,22}.

Some ultrasmall ROS scavenging nanomaterials have been developed recently. For example, Cai and co-workers reported the synthesis of ultrasmall melanin-based²³, molybdenum-based¹⁰, and DNA origami-based²⁴ antioxidative NPs with preferential

renal uptake for AKI treatment. Besides, Qu and co-workers²⁵ synthesized Cu-TCPP MOF nanodots with SOD mimic property. However, the relatively low catalytic activity and/or high cost of these nanomaterials could impair their clinical translation. It remains a challenge to develop a nanomaterial with simple structure and excellent ROS scavenging ability to enable potential large-scale production and subsequently clinical applications.

Copper (Cu), an essential trace element in humans, plays an important role in many enzymes, such as tyrosinase and Cu-Zn SOD^{26,27}. Therefore, it is reasonable to infer that copper-based nanomaterials can be used to scavenge ROS. For example, Cu NPs have excellent catalytic activity to scavenge H₂O₂ and O₂⁻ due to their strong quantum confinement of electrons in the ultrasmall size regime²⁶, but cannot eliminate OH· simultaneously²⁷. Cuprous oxide (Cu₂O) NPs possess good catalytic activity and can promote electron transfer reactions to inactivate H₂O₂ or OH·, thereby partially mimicking peroxidase^{13,28}. Therefore, we presume that broader-spectrum enzymatic catalytic properties and antioxidant activities could be concurrently achieved by combining Cu₂O and Cu nanocrystals. Besides, the stability of Cu₂O coating on Cu NPs can be largely enhanced owing to the effective electron-hole separation between Cu₂O and Cu, which is also of benefit to improve the overall ROS scavenging ability^{29,30}.

Herein, we report a strategy to synthesize ultrasmall Cu_{5,4}O NPs (Cu_{5,4}O USNPs) with excellent biocompatibility, enzymatic ROS scavenging abilities, and high renal clearance properties, in order to tackle broad ROS-related diseases (Fig. 1). Cu_{5,4}O USNPs exhibit remarkable antioxidant efficiency, with a working concentration of approximately 25 ng mL⁻¹ in vitro and 2 μg kg⁻¹ for AKI in vivo. This dosage is at least two orders magnitude lower than all the other reported nanomaterials to treat ROS-related diseases. Furthermore, they possess broad-spectrum ROS scavenging activities and function as CAT, GPx, and SOD analogs. Moreover, the ultrasmall NPs show high renal clearance and achieve an outstanding therapeutic effect against broad ROS-related diseases without any noticeable toxicity. This study

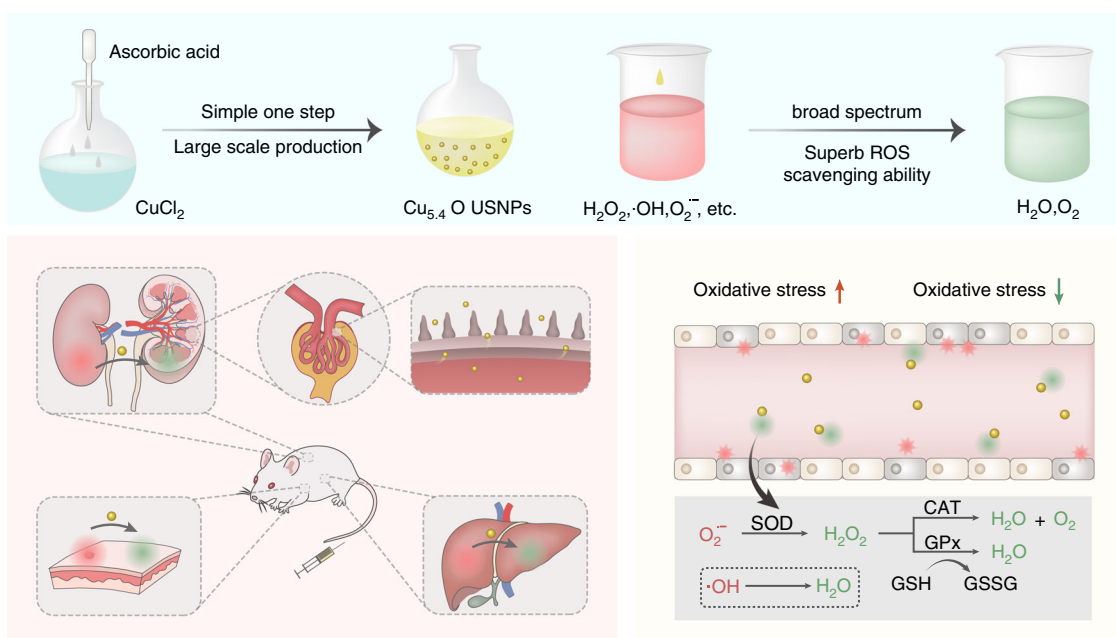


Fig. 1 Schematic illustration of Cu_{5,4}O ultrasmall nanoparticles in the treatment of ROS-related diseases. Cu_{5,4}O ultrasmall nanoparticles with multiple enzyme-mimicking and broad-spectrum ROS scavenging ability are synthesized by a simple and green method. Due to the robust ROS scavenging ability in vivo, Cu_{5,4}O ultrasmall nanoparticles exhibit therapeutic effect against broad ROS-related diseases, including acute kidney injury, acute liver injury and diabetic wound healing.

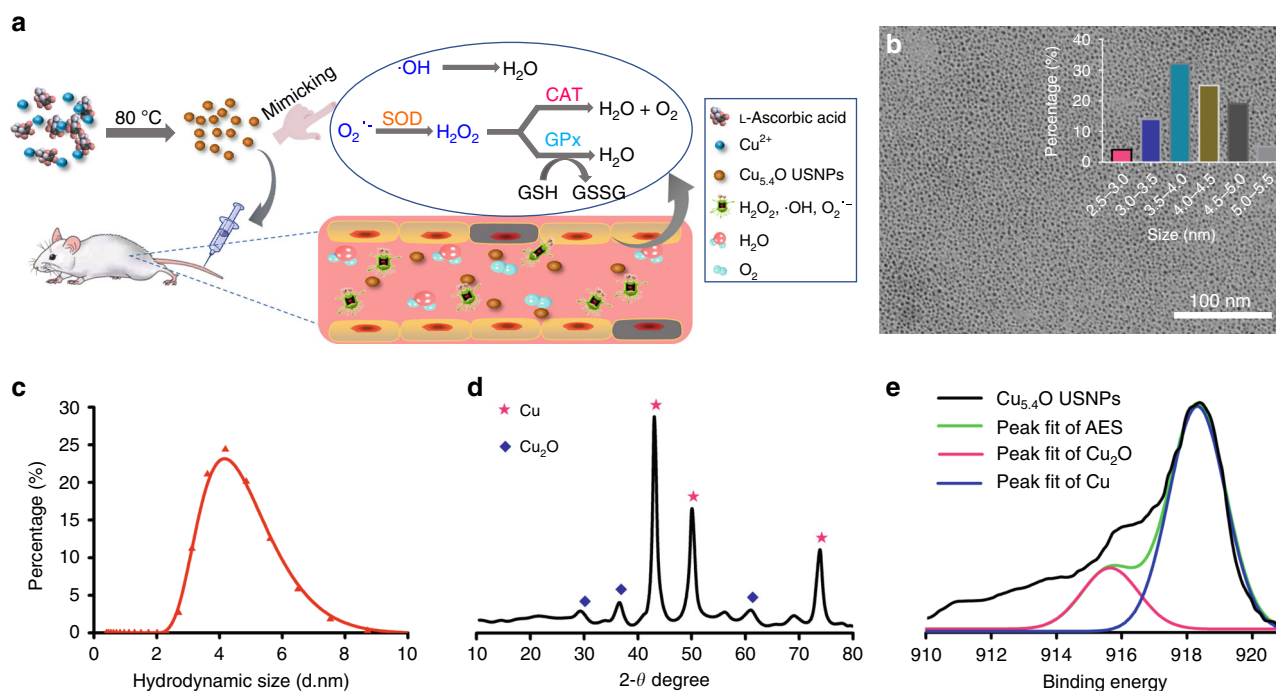


Fig. 2 Preparation and characterization of Cu_{5.4}O USNPs. **a** Schematic preparation of Cu_{5.4}O USNPs. **b** TEM image of Cu_{5.4}O USNPs. Inset is the statistical chart of particle size distribution. The cross-sectional area of each particle was measured by using ImageJ Software from the TEM images, with at least 500 particles counted per sample. **c** Hydrodynamic diameter distribution of the Cu_{5.4}O USNPs. **d** X-ray diffraction (XRD) pattern of the Cu_{5.4}O USNPs, the rhombus, and star symbols represent the characteristic peaks of Cu₂O and Cu, respectively. **e** XAES spectra of the Cu_{5.4}O USNPs. Source data are provided as a Source Data file.

provides an attractive strategy to develop ultrasmall copper-based nanozyme systems, which may serve as a blueprint for next-generation nanomedicines used in ROS-related diseases treatment and prevention.

Results and Discussion

Synthesis and characterization of Cu_{5.4}O USNPs. USNPs were synthesized by a green, rapid, and cost-effective method (Fig. 2a). The ratio of Cu²⁺ to L-ascorbic acid (AA), reaction temperature, and time were tuned to determine their impact on the particle size and catalytic activity of obtained NPs (Supplementary Figs. 1–3). A relatively long reaction time is required for obtaining uniform USNPs due to Ostwald ripening (Supplementary Fig. 2). Too low and too high temperatures are not feasible to control the reaction (Supplementary Fig. 3). The catalytic activity of the obtained copper-based USNPs was almost the same with the feeding ratio of Cu²⁺ to AA in the range of 1:10 to 1:40. Therefore, the molar ratio of Cu²⁺ to AA was fixed at 1:10 for the following studies.

Transmission electron microscopy (TEM) images showed that these Cu_{5.4}O USNPs were uniform, with an average diameter of 3.5–4.0 nm in the dry state (Fig. 2b). The average hydrodynamic diameter of Cu_{5.4}O USNPs was approximately 4.5 nm (Fig. 2c), which meets the kidney filtration threshold of 5.5 nm^{21,22}, enabling renal uptake, accumulation, and clearance. There were slight increases in the hydrophilic diameters of Cu_{5.4}O USNPs in FBS and rat serum (Supplementary Fig. 4a), possibly due to surface protein adsorption³¹. Cu_{5.4}O USNPs were uniformly dispersed, with nearly identical morphologies and particle sizes, indicating that the Cu_{5.4}O USNPs were stable in the media for at least 20 days (Supplementary Fig. 4b). The oxidation state of Cu was investigated through powder X-ray diffraction (Fig. 2d). The dominant peaks at 2θ = 41.6°, 50.7°, and 77.5°, which can be assigned to the (111), (200), and (220) lattice planes of Cu(0), are consistent with the presence of face-centered cubic copper.

Three additional minor diffraction peaks at 2θ = 29.6°, 36.4°, and 60.1° can be assigned to the (111), (200), and (220) lattice planes of cuprous oxide (Cu₂O), respectively. The results suggest that the reaction did not generate pure copper NPs, but rather a mixture of Cu⁰ and Cu⁺ (Cu and Cu₂O NPs). As shown in Fig. 2e, the mass fractions of Cu and Cu₂O after normalization were calculated based on the peak area of Cu 2p, indicating that the proportions of Cu and Cu₂O were approximately 3.4. Therefore, the resulting ultrasmall copper-based NPs are denoted as Cu_{5.4}O USNPs.

ROS scavenging activities of Cu_{5.4}O USNPs. Three representative ROS, H₂O₂, O₂^{·-}, and ·OH, were selected to investigate the ROS scavenging activities of Cu_{5.4}O USNPs. As shown in Fig. 3, the Cu_{5.4}O USNPs exhibited a high ROS scavenging activity in a concentration-dependent manner. Approximately 80% of the total H₂O₂ was decomposed by 200 ng mL⁻¹ Cu_{5.4}O USNPs (Fig. 3a). Approximately half of the O₂^{·-} was decomposed when treated with 150 ng mL⁻¹ Cu_{5.4}O USNPs (Fig. 3b). More than 80% of the ·OH was scavenged when the Cu_{5.4}O USNP concentration was 150 ng mL⁻¹ (Fig. 3c, Supplementary Fig. 5). To further confirm the antioxidative properties of Cu_{5.4}O USNPs, a free radical scavenging experiment was performed using the classic 2,2'-azino-bis(3-ethylbenzothiazoline 6-sulfonate) (ABTS) radical assay. As presented in Fig. 3d, more than 89% of the free radicals were eliminated by a very low concentration of Cu_{5.4}O USNPs (150 ng mL⁻¹).

Compared with the working concentrations of reported metal-based nano-antioxidants for ROS scavenging at cellular level, such as Ce³⁺, Au³⁺, TiO₂ (ref. 34), MnO₂ (ref. 35), Co₃O₄ (ref. 36), and V₂O₅ (ref. 37), which are typically in the range of 20–50 μg mL⁻¹ in vitro, the working concentration of Cu_{5.4}O USNPs (25 ng mL⁻¹) was 2–3 orders of magnitude lower, indicating higher antioxidant efficacy. The working concentration of Cu_{5.4}O USNPs was 200-fold

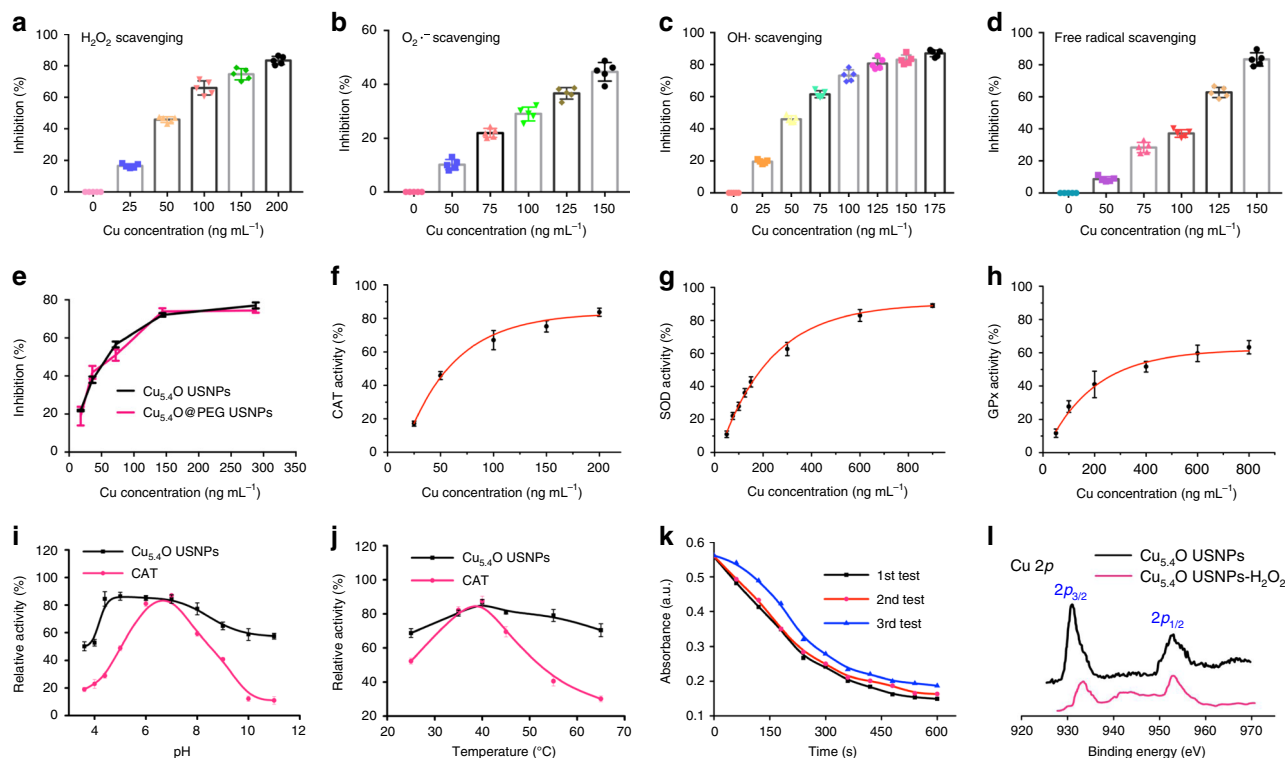


Fig. 3 ROS scavenging and multienzyme-like antioxidative activity of $\text{Cu}_{5.4}\text{O}$ USNPs. **a** H_2O_2 , **b** $\text{O}_2^{\cdot-}$, **c** $\text{OH}\cdot$, and **d** free radical scavenging ability of $\text{Cu}_{5.4}\text{O}$ USNPs. **e** H_2O_2 -scavenging capacities of $\text{Cu}_{5.4}\text{O}$ USNPs and $\text{Cu}_{5.4}\text{O}@$ PEG USNPs. **f** CAT-like, **g** SOD-like, and **h** GPX-like activity of $\text{Cu}_{5.4}\text{O}$ USNPs. **i, j** The CAT-like activity of the $\text{Cu}_{5.4}\text{O}$ USNPs after different treatments. H_2O_2 (2 mM) elimination efficiencies of $\text{Cu}_{5.4}\text{O}$ USNPs (250 ng mL^{-1}) and natural CAT (40 U mL^{-1}), respectively, pretreated with different **i** temperature and **j** pH conditions. **k** Stability test of $\text{Cu}_{5.4}\text{O}$ USNPs (250 ng mL^{-1}) when used as CAT-like catalyst at different cycles. The concentration of H_2O_2 is 2 mM. **l** XPS analysis of $\text{Cu}_{5.4}\text{O}$ USNPs before and after treatment with H_2O_2 . In **a–j**, data represent means \pm s.d. from five (**a–d**) or three (**e–j**) independent replicates. Source data are provided as a Source Data file.

lower than that of previously reported Cu_xO ($x = 1–2$)¹⁹ for the same H_2O_2 -scavenging capacity. Approximately 80–90% of H_2O_2 (1 mM) can be scavenged by 0.4 mg mL^{-1} of vitamin C³⁸, whereas 200 ng mL^{-1} of $\text{Cu}_{5.4}\text{O}$ USNPs could scavenge approximately 85% of H_2O_2 (2 mM). Therefore, $\text{Cu}_{5.4}\text{O}$ USNPs showed much stronger H_2O_2 -scavenging efficiency (approximately 2000-fold) than commonly used small molecular antioxidants, such as vitamin C. To the best of our knowledge, the working concentration of $\text{Cu}_{5.4}\text{O}$ USNPs was also the lowest among all ROS scavenging nanomaterials that have been reported so far (Supplementary Table 1).

Mechanism of ROS scavenging activities of $\text{Cu}_{5.4}\text{O}$ USNPs.

Since AA has strong antioxidant activity and can eliminate harmful free radicals both in vitro and in vivo³⁹, we initially suspected that the ROS scavenging activities of $\text{Cu}_{5.4}\text{O}$ USNPs originated from the surface-coated AA. We used chemically inert HS-PEG-OH to replace AA molecules on the surface of $\text{Cu}_{5.4}\text{O}$ USNPs (Supplementary Fig. 6). Surprisingly, the H_2O_2 -scavenging abilities of $\text{Cu}_{5.4}\text{O}$ and $\text{Cu}_{5.4}\text{O}@$ PEG USNPs were very similar (Fig. 3e), suggesting that the presence of AA is not essential for the excellent ROS scavenging abilities of $\text{Cu}_{5.4}\text{O}$ USNPs. According to the previous reports^{40,41} and our results in Supplementary Figs. 7 and 8, the AA molecules were possibly converted into dehydroascorbic acid through oxidation of Cu^{2+} , and thus lost their reduction activity.

Multienzyme-like antioxidative activity of $\text{Cu}_{5.4}\text{O}$ USNPs.

We further investigated the antioxidant enzyme-mimetic activities of $\text{Cu}_{5.4}\text{O}$ USNPs. Among biologically relevant ROS, H_2O_2 is of greatest importance because of its membrane permeability, longer half-life than $\text{O}_2^{\cdot-}$ and $\cdot\text{OH}$, and consequently highest intracellular

concentration⁴². Therefore, we focused on the CAT-like property of $\text{Cu}_{5.4}\text{O}$ USNPs, which was responsible for decomposition of H_2O_2 . As shown in Fig. 3f, the concentration-dependent CAT-like activity of $\text{Cu}_{5.4}\text{O}$ USNPs was investigated. Terephthalic acid (TPA) was used as a fluorescence probe, which reacts with $\cdot\text{OH}$ from H_2O_2 , forming highly fluorescent 2-hydroxyterephthalic acid. Gradual reduction of fluorescence intensity was observed as the concentration of $\text{Cu}_{5.4}\text{O}$ USNPs increased (Supplementary Fig. 9). Gradual reduction in the characteristic absorbance of H_2O_2 at 240 nm (Supplementary Fig. 10a) and gradual enhancement of the O_2 content (Supplementary Fig. 10b) were observed with a prolonged reaction time in the presence of $\text{Cu}_{5.4}\text{O}$ USNPs.

A steady-state kinetic assay was performed to confirm the enzymatic activity by varying the concentration of H_2O_2 in the presence of $\text{Cu}_{5.4}\text{O}$ USNPs. This reaction showed typical Michaelis–Menten kinetics (Supplementary Fig. 11). The K_m value of the $\text{Cu}_{5.4}\text{O}$ USNPs with H_2O_2 as the substrate was 0.065 mM, and the V_{max} value was $3.92 \times 10^{-6} \text{ Ms}^{-1}$. The K_m value of $\text{Cu}_{5.4}\text{O}$ USNPs was smaller than that of natural CAT (0.134 mM), indicating that our $\text{Cu}_{5.4}\text{O}$ USNPs possessed greater affinity for substrate H_2O_2 , possibly due to their ultrasmall size and greater numbers of exposed active sites.

Moreover, the concentration-dependent SOD-like activity of $\text{Cu}_{5.4}\text{O}$ USNPs was also investigated. As shown in Fig. 3g, the percentage of formed formazan reduced significantly in the presence of $\text{Cu}_{5.4}\text{O}$ USNPs, indicating the SOD-like activity of $\text{Cu}_{5.4}\text{O}$ USNPs. According to the previous literature, the EC_{50} (an indicator for comparing the efficiencies of enzymes and enzyme mimics) of native Cu–Zn SOD^{25,43} was 41.6 ng mL^{-1} . The EC_{50} of $\text{Cu}_{5.4}\text{O}$ USNPs was calculated to be 191.4 ng mL^{-1} . Thus, the EC_{50} of $\text{Cu}_{5.4}\text{O}$ USNPs was approaching 21.7% of the native SOD

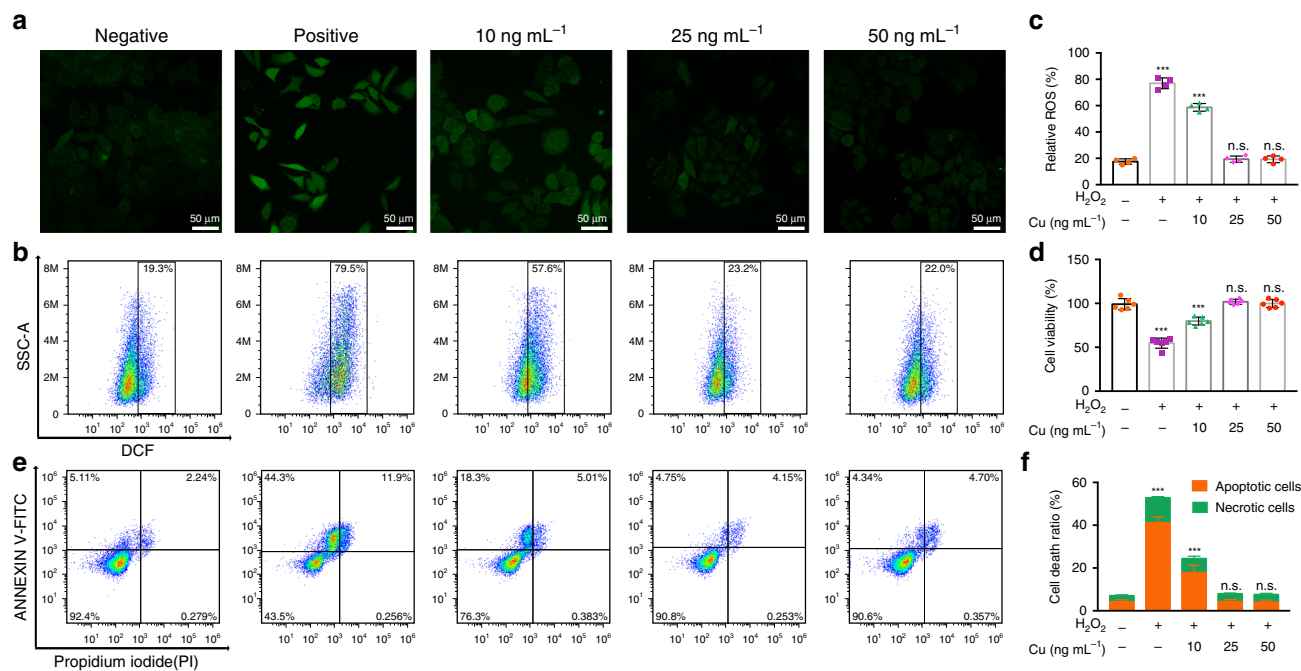


Fig. 4 Scavenging ROS with $\text{Cu}_{5.4}\text{O}$ USNPs in vitro. **a** Representative ROS staining (green fluorescence) of HEK293 cells under different treatment conditions. **b** ROS levels in untreated and $\text{Cu}_{5.4}\text{O}$ USNPs-treated HEK293 cells incubated with $250\ \mu\text{M}$ H_2O_2 . **c** Statistical analysis of ROS levels in HEK293 cells under different treatment conditions. **d** In vitro cell viabilities of HEK293 cells under different treatment conditions. **e** FACS results of cell apoptosis and necrosis distribution in untreated and $\text{Cu}_{5.4}\text{O}$ USNPs-treated HEK293 cells. **f** Statistical analysis of necrotic and apoptotic cell ratios in HEK293 cells under different treatment conditions. In **c**, **d** and **f**, data represent means \pm s.d. from four (**c**, **f**) or six (**d**) independent replicates. *** $P < 0.001$; n.s., no significance, one-way ANOVA. Source data are provided as a Source Data file.

activity, which was nearly twice as much as that of the Cu-TCPP MOF nanodots (12.6% of native SOD activity) reported by Qu and co-workers²⁵. Besides, the SOD-like activity of $\text{Cu}_{5.4}\text{O}$ USNPs was further confirmed by electron paramagnetic resonance (EPR) spectroscopy (Supplementary Fig. 12). In addition, $\text{Cu}_{5.4}\text{O}$ USNPs also displayed concentration-dependent GPx-like activity, eliminating H_2O_2 and catalyzing the oxidation of reduced glutathione (Fig. 3h).

Natural enzymes often exhibit intrinsic shortcomings, such as low operational stability, temperature and pH sensitivity, and recycling difficulties. Hence, the thermal and pH stabilities of $\text{Cu}_{5.4}\text{O}$ USNPs were investigated and compared with those of the natural enzyme CAT. The results (Fig. 3i, j) suggested that the stabilities of $\text{Cu}_{5.4}\text{O}$ USNPs when exposed to pH and temperature variations were significantly greater than those of natural CAT. Besides, the recycled $\text{Cu}_{5.4}\text{O}$ USNPs showed nearly identical catalytic activity as the original solution (Fig. 3k), indicating good stability and recyclability of $\text{Cu}_{5.4}\text{O}$ USNPs.

The X-ray photoelectron spectroscopy (XPS) spectra of the Cu $2p$ core level region for $\text{Cu}_{5.4}\text{O}$ and $\text{Cu}_{5.4}\text{O}$ oxidized by H_2O_2 showed only two intense peaks at 932.4 and 952.0 eV for $\text{Cu}_{5.4}\text{O}$ USNPs before and after oxidation, which were assigned to the binding energies of Cu $2p_{3/2}$ and Cu $2p_{1/2}$; these corresponded to Cu^+ and/or Cu^0 species, respectively (Fig. 3l). Upon interaction with H_2O_2 , the positions of the two main peaks did not shift and new peaks rarely appeared. Hence, we concluded that the ROS scavenging performance of $\text{Cu}_{5.4}\text{O}$ USNPs can be attributed to its intrinsic multienzyme-mimicking properties.

Scavenging ROS ability in vitro. Compared with the glomerulus, renal tubules are more susceptible to oxidative stress in AKI⁴⁴. Therefore, protection of renal tubules against ROS damage and subsequent initiation of a cascade of pathological processes during the early stage of AKI would significantly reduce kidney

dysfunction⁴⁴. For this analysis, the human embryonic kidney 293 (HEK293) cell line was used to examine the cytoprotective properties of $\text{Cu}_{5.4}\text{O}$ USNPs against ROS damage in vitro. As shown in Fig. 4a, the intracellular ROS level (green fluorescent signal) of HEK293 cells increased dramatically after treatment with $250\ \mu\text{M}$ H_2O_2 , leading to abnormal and shrunken cell morphology. In comparison, the intracellular ROS level obviously decreased when the cells were pretreated with $\text{Cu}_{5.4}\text{O}$ USNPs. Quantitative analysis of intracellular ROS levels via flow cytometry further confirmed this trend (Fig. 4b, c, Supplementary Fig. 13a). The results of CCK-8 analysis (Fig. 4d) revealed that an extremely low concentration ($25\ \text{ng mL}^{-1}$) of $\text{Cu}_{5.4}\text{O}$ USNPs was able to completely protect the cells against $250\ \mu\text{M}$ H_2O_2 , which is higher than the concentration in the classical AKI pathological microenvironment.

We further examined the effect of $\text{Cu}_{5.4}\text{O}$ USNPs against H_2O_2 -induced cell apoptosis and necrosis via flow cytometry. In Fig. 4e, f and Supplementary Fig. 13b, the addition of $\text{Cu}_{5.4}\text{O}$ USNPs significantly reduced the ratios of apoptotic and necrotic cells induced by H_2O_2 treatment, further confirming the ROS scavenging and cytoprotective properties of $\text{Cu}_{5.4}\text{O}$ USNPs at the cellular level.

As shown in Supplementary Fig. 14, $\text{Cu}_{5.4}\text{O}$ USNPs could be found in the mitochondria of cells, which was consistent with previous reports that ultrasized NPs could easily enter the mitochondria through the mitochondrial permeability transition pore^{45,46}, indicating that $\text{Cu}_{5.4}\text{O}$ USNPs could alleviate mitochondrial oxidative stress and maintain mitochondrial function. Besides, $\text{Cu}_{5.4}\text{O}$ USNPs could also be observed in the phagosome, suggesting that $\text{Cu}_{5.4}\text{O}$ USNPs might enter into the cells through phagocytosis.

In vitro and in vivo biocompatibility of $\text{Cu}_{5.4}\text{O}$ USNPs. The CCK-8 assay results (Supplementary Fig. 15) suggested that

Cu_{5,4}O USNPs at test concentrations did not exhibit noticeable cytotoxicity. HEK293 cells exposed to 200 ng mL⁻¹ Cu_{5,4}O USNPs for 48 h showed normal polygonal cytoskeleton morphology (Supplementary Fig. 16), indicating good biocompatibility. Such results are in accordance with previously findings that copper NPs are generally non-toxic at relatively low concentrations (<5 μg mL⁻¹)⁴⁷.

Additionally, hemolysis rate of nanomaterials must be less than 5% to ensure safety during intravenous administration^{48,49}. As shown in Supplementary Fig. 17, the hemolysis rate of 500 ng mL⁻¹ Cu_{5,4}O USNPs, which is 10-fold greater than the concentration applied in vivo to treat AKI (2 μg kg⁻¹, corresponding to 50 ng mL⁻¹ for a 25 g mouse with 1 mL of blood), was less than 5%.

Next, we evaluated the impacts of Cu_{5,4}O USNPs (4 μg kg⁻¹, twofold greater than the concentration used to treat AKI) on blood chemistry, inflammatory cytokine levels, and major organ histopathology in healthy mice to reveal their in vivo biocompatibility. As shown in Fig. 5a and Supplementary Fig. 18, no necrosis, congestion, or hemorrhage was observed in the heart, liver, spleen, and lung at 1 and 30 days after single dose intravenous injection of Cu_{5,4}O USNPs. Moreover, no distinguishable inflammatory lesions or tissue damage were observed in the glomerulus, tubules, collecting ducts, and urethra at 24 h after single dose intravenous injection of Cu_{5,4}O USNPs (Fig. 5b).

As shown in Fig. 5c, the serum levels of interleukin-6 (IL-6) and tumor necrosis factor-α (TNF-α) in the Cu_{5,4}O USNPs-treated group were identical to the levels in the control group ($P > 0.05$), indicating that Cu_{5,4}O USNPs would not trigger obvious immune responses in vivo at the tested concentration. The serum biochemistry analysis results (Fig. 5d, e) showed that serum concentrations of liver function indicators (aspartate transaminase (AST) and alanine transaminase (ALT)) and kidney function indicators (BUN and CRE) in the Cu_{5,4}O USNPs-treated group were similar to those in the control group ($P > 0.05$), revealing good biocompatibility in the liver and kidney. Moreover, the results of complete blood panel analysis (Fig. 5f–j) showed no obvious differences in the hematology of the Cu_{5,4}O USNPs-treated group when compared to that of the control group ($P > 0.05$).

We also investigated the accumulation of Cu_{5,4}O USNPs in the major organs after repeated daily administration of Cu_{5,4}O USNPs for seven consecutive days. As shown in Supplementary Fig. 19, the liver and kidneys exhibited higher normalized dosage distribution of Cu_{5,4}O USNPs, reaching 7.92 and 7.97 %ID g⁻¹ respectively, while the distributions of NPs in the heart, spleen, lung, and blood reached 2.25, 5.07, 3.02, and 2.84 %ID g⁻¹, respectively. As shown in Supplementary Fig. 20, Cu_{5,4}O USNPs could be found in the tissues of major organs by TEM, which is consistent with the above results.

The in vivo toxicity of Cu_{5,4}O USNPs accumulated in the major organs after repeated intravenous administration was also evaluated. As shown in Supplementary Fig. 21, no necrosis, congestion, or hemorrhage was found in the heart, liver, spleen, lung, and kidney of the mice after intravenous repeated administration. Besides, the serum biochemistry analysis and complete blood panel analysis (Supplementary Fig. 22) results further indicated no obvious toxicity of Cu_{5,4}O USNPs. All the results confirmed that the synthesized Cu_{5,4}O USNPs exhibited negligible short-term and long-term in vivo toxicity.

Pharmacokinetics and biodistribution of Cu_{5,4}O USNPs in mice. The time-dependent blood circulation profiles of Cu_{5,4}O USNPs in Fig. 6a demonstrated a classical two-compartment pharmacokinetic model. The terminal elimination half-lives of the central component and peripheral component were 0.77 and

71.2 h, respectively. The biodistribution of Cu_{5,4}O USNPs in the major organs of AKI mice at 24 h post injection was detected by inductively coupled plasma-atomic emission spectrometry. As shown in Fig. 6b, the kidney exhibited the highest normalized dosage distribution of Cu_{5,4}O USNPs (4.4 %ID g⁻¹), while the distributions of NPs in the heart, liver, spleen, and lung reached 0.89, 3.2, 2.1, and 0.94 %ID g⁻¹, respectively. The accumulation of Cu_{5,4}O USNPs in the kidneys might be attributed to the high renal uptake resulting from their ultrasmall hydrodynamic diameter (approximately 4.5 nm) and excellent hydrophilicity. Previous studies showed that the size threshold of the glomerular basement membrane (GBM) was approximately 5.5 nm, and nanomaterials with a diameter less than 5.5 nm could be effectively cleared from the blood to the renal tubules through the GBM^{50–52}. As shown in Supplementary Fig. 23, TEM analysis revealed Cu_{5,4}O USNPs in the urine of AKI mice after injection, further confirming that the Cu_{5,4}O USNPs could pass through the GBM to the tubules and then undergo excretion in the urine. The time-dependent accumulation of Cu_{5,4}O USNPs in the kidneys of normal and AKI mice indicated that Cu_{5,4}O USNPs could effectively accumulate in the kidneys, regardless of the permeability of the GBM (Supplementary Fig. 24). Moreover, time-dependent accumulation of Cu_{5,4}O USNPs in the urine revealed that renal clearance is the main excretion pathway for these particles, rather than intestinal excretion (Fig. 6c). Approximately 70% of Cu_{5,4}O USNPs could be excreted within 48 h post injection (60% through the kidneys and 10% through the intestine), suggesting relatively rapid clearance and potentially low long-term toxicity of Cu_{5,4}O USNPs.

To further analyze the distribution of Cu_{5,4}O USNPs in the kidneys, renal tissues were collected, ultrathin sectioned, and observed under TEM. As shown in Fig. 6d and Supplementary Fig. 25, Cu_{5,4}O USNPs (arrowheads) were found in the GBM, urinary space, and epithelial cell cilia of renal tubules, confirming that the Cu_{5,4}O USNPs could pass through the GBM and reach tubules containing urine.

In vivo therapeutic efficacy of Cu_{5,4}O USNPs on AKI mice. The therapeutic effect of Cu_{5,4}O USNPs on AKI mice was tested (Fig. 7a). As shown in Fig. 7b, compared with the control group (90% of mice died within 5 days following establishment of the AKI model), all AKI mice treated with Cu_{5,4}O USNPs survived for more than 14 days. In addition, AKI mice treated with Cu_{5,4}O USNPs exhibited a weight increase similar to that of healthy mice (Fig. 7c), while control AKI mice underwent dramatic body weight loss within 24 h. Furthermore, the serum levels of two important kidney function indicators, BUN and CRE levels, of AKI mice treated with Cu_{5,4}O USNPs were significantly lower than those of AKI mice in the control group ($P < 0.001$, Fig. 7d, e). There were no detectable differences in BUN and CRE levels between healthy mice and Cu_{5,4}O USNPs-treated AKI mice ($P > 0.05$), confirming the excellent therapeutic efficacy of Cu_{5,4}O USNPs in the treatment of glycerol-induced AKI.

In kidney diseases, the precipitation of denatured proteins in tubules forms a cast structure, which is regarded as an important diagnostic marker of pathological changes. As shown in Fig. 7f, many casts (marked as triangles) could be found in the renal tissues of AKI mice, whereas only a few casts could be observed in AKI mice that were treated with Cu_{5,4}O USNPs, suggesting that renal tissue integrity was maintained via adoption of the NPs.

To further demonstrate the therapeutic activity of Cu_{5,4}O USNPs acting as a ROS scavenger in vivo, NAC molecules with different concentrations (8, 40, and 160 mg kg⁻¹) were intravenously injected into AKI mice as controls. As shown in Supplementary Fig. 26, the CRE, BUN levels and survival percentage of AKI mice

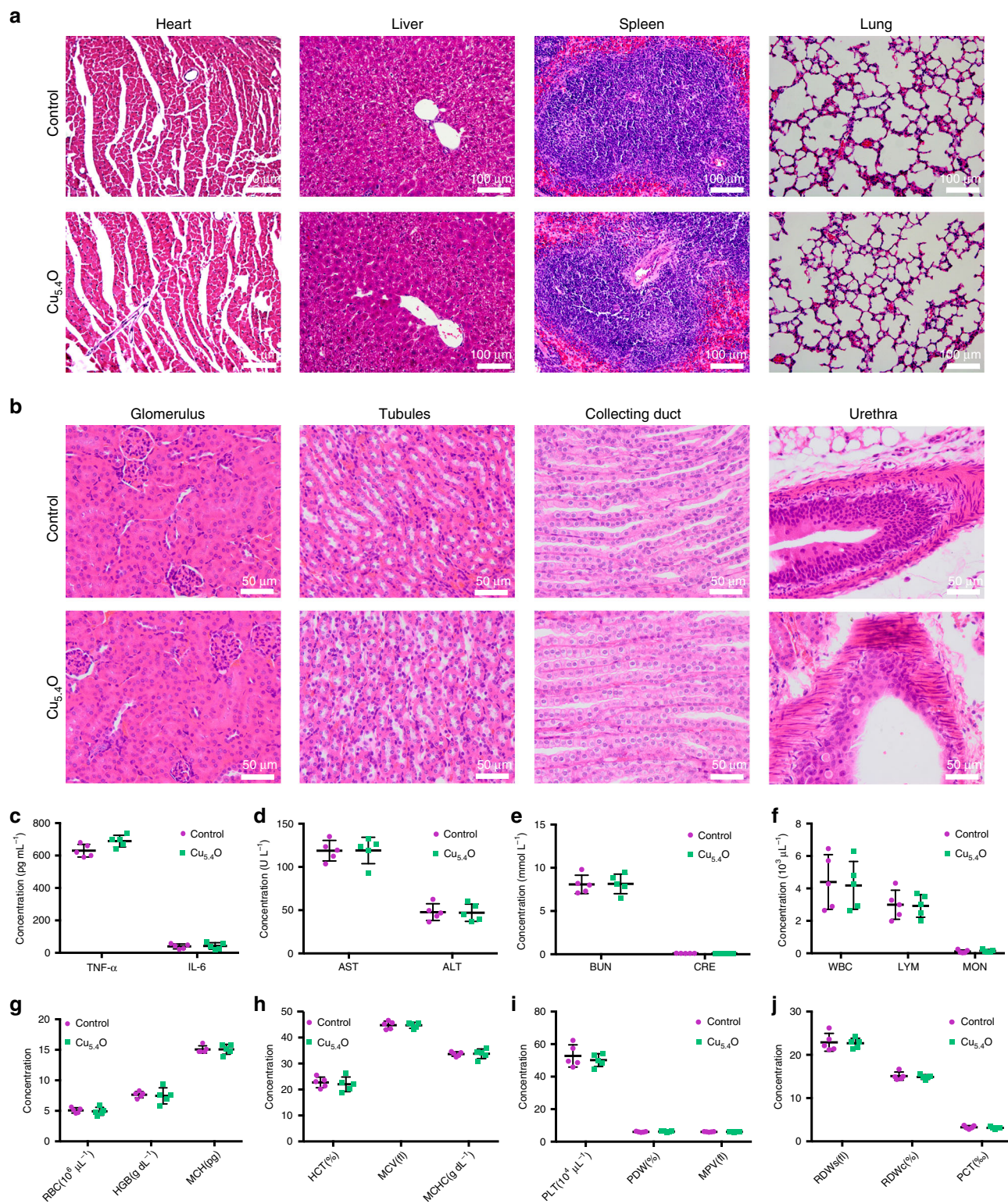


Fig. 5 **In vivo** biocompatibility assessment of $\text{Cu}_{5.4}\text{O}$ USNPs. **a** Evaluation of in vivo toxicity of $\text{Cu}_{5.4}\text{O}$ USNPs to major organs (heart, liver, spleen, and lung) at 24 h after intravenous administration. **b** Assessment of in vivo toxicity of $\text{Cu}_{5.4}\text{O}$ USNPs to the kidney at 24 h after intravenous administration. **c** Serum levels of inflammatory factors of interleukin-6 (IL-6) and tumor necrosis factor alpha (TNF- α). **d** Serum levels of liver function indicators: aspartate transaminase (AST) and alanine transaminase (ALT). **e** Serum levels of kidney function indicators: blood urea nitrogen (BUN) and creatinine (CRE). **f–j** Blood parameters in normal mice (control group), and mice intravenously injected with $\text{Cu}_{5.4}\text{O}$ USNPs, 24 h after injection. In **c–j**, data represent means \pm s.d. from five independent replicates. Source data are provided as a Source Data file.

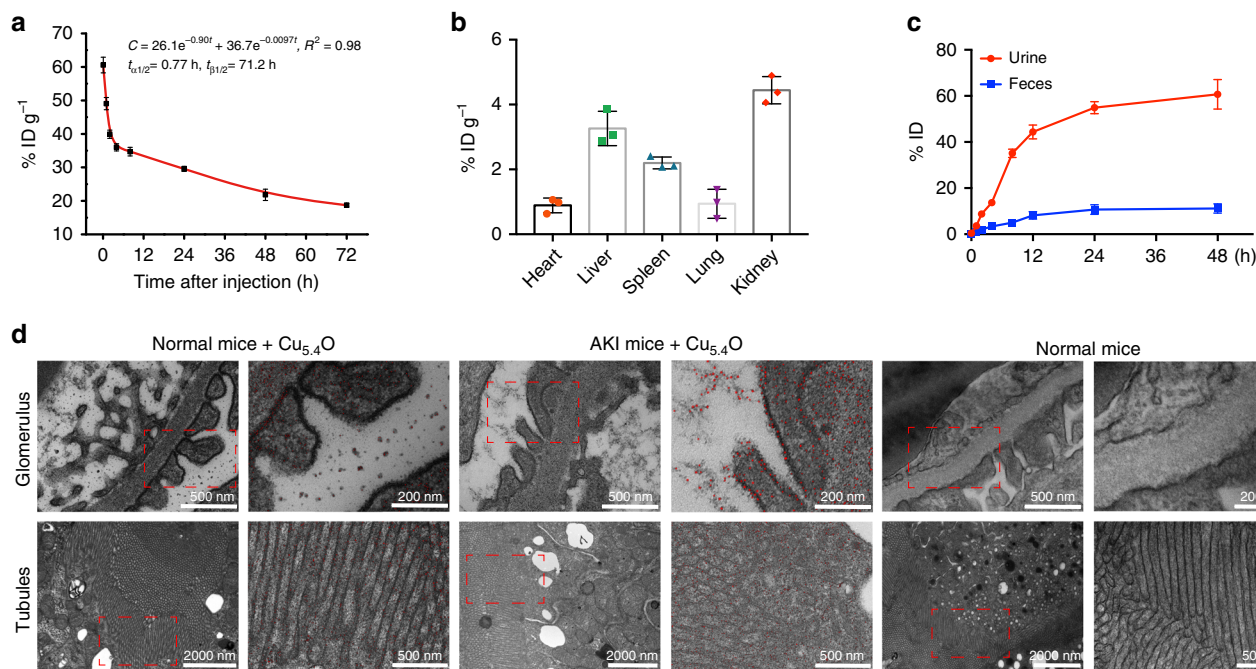


Fig. 6 Pharmacokinetics, biodistribution, and clearance of $\text{Cu}_{5.4}\text{O}$ USNPs in AKI mice. **a** In vivo blood pharmacokinetic curves after intravenous injection of $\text{Cu}_{5.4}\text{O}$ USNPs. **b** Biodistribution of $\text{Cu}_{5.4}\text{O}$ USNPs in major organs at 24 h post injection. **c** Cumulative urine and feces excretion at different time points. **d** Biodistribution of $\text{Cu}_{5.4}\text{O}$ USNPs in renal tissues by TEM observation. Red dashed lines indicate the magnified area. Dots with red pseudocolor indicate $\text{Cu}_{5.4}\text{O}$ USNPs. In **a–c**, data represent means \pm s.d. from three independent replicates. Source data are provided as a Source Data file.

treated with $\text{Cu}_{5.4}\text{O}$ USNPs (at $2 \mu\text{g kg}^{-1}$ concentration) were similar to those of AKI mice treated with 160 mg kg^{-1} of NAC, which was also in accordance with the histological analysis of renal tissues. The overall results indicated the in vivo ROS scavenging activity of $\text{Cu}_{5.4}\text{O}$ USNPs.

The ROS levels and SOD activity in the kidneys were further investigated to understand the molecular mechanism underlying the protection process. Compared with PBS-treated AKI mice, the levels of superoxide in $\text{Cu}_{5.4}\text{O}$ USNPs-treated AKI mice were significantly reduced, reaching the levels exhibited in normal mice (Fig. 7g, Supplementary Fig. 27). As shown in Fig. 7h, the renal SOD activity in $\text{Cu}_{5.4}\text{O}$ USNPs-treated AKI mice was similar to that of healthy mice, whereas a significant reduction in SOD activity was observed in AKI mice in the control group ($P < 0.001$). This result suggested that $\text{Cu}_{5.4}\text{O}$ USNPs could protect renal cells by functioning as antioxidants to scavenge ROS and maintain SOD activity in vivo. Furthermore, the renal expression levels of two important kidney injury biomarkers^{53,54}, heme oxygenase-1 (HO-1) and kidney injury molecule-1 (KIM-1), were detected. As shown in Fig. 7i, j, compared with AKI mice in the control group, $\text{Cu}_{5.4}\text{O}$ USNPs-treated AKI mice exhibited significantly reduced levels of KIM-1 and HO-1 ($P < 0.001$), consistent with the results we found for CRE and BUN.

The cisplatin-induced AKI (Cis-AKI) mouse model was also used to demonstrate the broad application of $\text{Cu}_{5.4}\text{O}$ USNPs for ROS-mediated AKI diseases. As shown in Supplementary Fig. 28, kidney function examination and histological analysis revealed that kidney damage in Cis-AKI mice treated with $\text{Cu}_{5.4}\text{O}$ USNPs was significantly lower than that in Cis-AKI mice treated with PBS ($P < 0.001$; $P < 0.05$), confirming the therapeutic effect of $\text{Cu}_{5.4}\text{O}$ USNPs against Cis-AKI.

Therapeutic effect on acute liver injury and wound healing. It is intriguing for us to discover whether the $\text{Cu}_{5.4}\text{O}$ USNPs could be applicable to ROS-related pathological conditions other than AKI. Therefore, acetaminophen (APAP)-induced acute liver

injury (AILI) and full-thickness skin defect were chosen as the representative ROS-related disease models as well. As shown in Fig. 8a, b, the ALT and AST levels of $\text{Cu}_{5.4}\text{O}$ USNPs-treated AILI mice were significantly lower than those of AILI mice without treatment ($P < 0.001$), indicating the excellent therapeutic effect of $\text{Cu}_{5.4}\text{O}$ USNPs on AILI. Histological observation of liver tissues further confirmed the therapeutic effect of $\text{Cu}_{5.4}\text{O}$ USNPs on AILI (Fig. 8c).

Elevated ROS production in the wound site has also been implicated in delaying wound healing^{32,55}, especially in chronic diabetic wounds where high glucose and proinflammatory environment caused the large production of ROS in the wound bed^{56,57}. As shown in Fig. 8d, e, the diabetic wound healing rate of $\text{Cu}_{5.4}\text{O}$ USNPs group was always significantly faster than that of the corresponding control group on days 4, 7, 9, and 15 post-surgery ($P < 0.01$). The results clearly indicated that $\text{Cu}_{5.4}\text{O}$ USNPs could accelerate the diabetic wound healing process. As shown in Fig. 8f–i, the length of newly regenerated epidermis and the thickness of granulation tissue in the $\text{Cu}_{5.4}\text{O}$ USNPs group were significantly greater than those in the control group, further confirming that $\text{Cu}_{5.4}\text{O}$ USNPs could promote diabetic wound healing.

Therapeutic mechanisms of $\text{Cu}_{5.4}\text{O}$ USNPs on AKI. To further elucidate the underlying therapeutic mechanisms, AKI was chosen as the representative disease model for further transcriptomics analysis. An unguided principal component analysis (PCA) of the data revealed substantially different transcriptomic profiles between $\text{Cu}_{5.4}\text{O}$ - and PBS-treated AKI mice kidneys (Fig. 9a). The Venn diagram in Fig. 9b showed that 13025 genes were co-expressed by two groups, while 584 genes were exclusively expressed by $\text{Cu}_{5.4}\text{O}$ -treated group. Volcano plots (Fig. 9c) showed 5819 significantly differentially expressed genes (DEGs), of which 2813 and 3006 genes were upregulated and down-regulated, respectively.

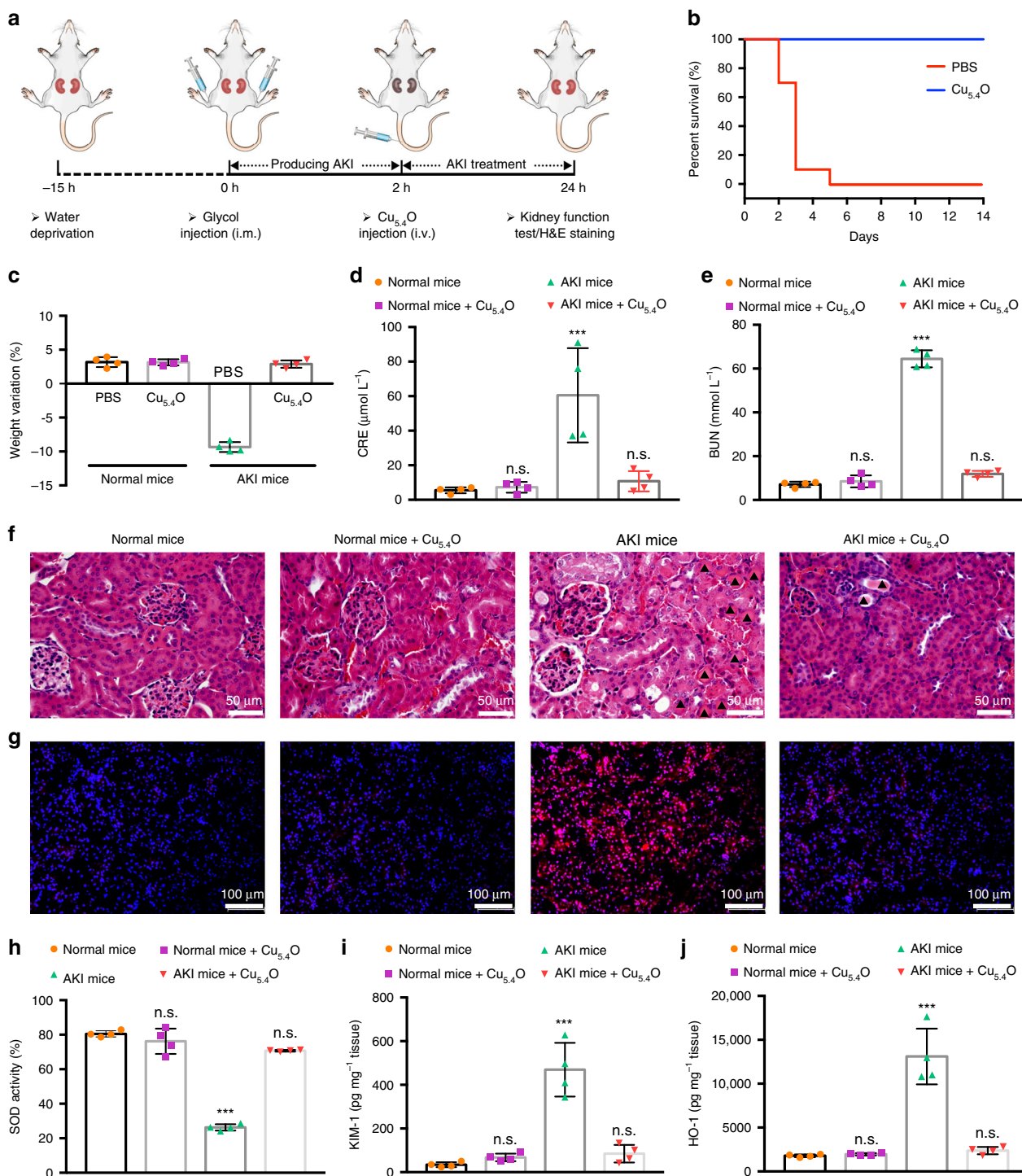


Fig. 7 Therapeutic efficiency of $\text{Cu}_{5,4}\text{O}$ USNPs on AKI mice. **a** Schematic illustration of the establishment and treatment schedule of AKI mice. **b** Survival curves of AKI mice with different treatment. **c** Weight variation of AKI mice at 24 h after treatment with $\text{Cu}_{5,4}\text{O}$ USNPs. Serum levels of **d** CRE and **e** BUN in AKI mice at 24 h after different treatment. **f** H&E staining of kidney tissues from each group. Triangles indicate the formation of casts. **g** Dihydroethidium (red fluorescence) and DAPI (blue fluorescence) staining of kidney tissues from each group. **h** SOD, **i** KIM-1, and **j** HO-1 levels measured in renal tissue homogenates from each group. In **c-e** and **h-j**, data represent means \pm s.d. from four independent replicates ($***P < 0.001$; n.s., no significance, One-way ANOVA). Source data are provided as a Source Data file.

Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway enrichment analysis (Fig. 9d) indicated that the glutathione (GSH) metabolism, MAPK signaling pathway, and TNF signaling pathways were highly associated with the therapeutic mechanisms of $\text{Cu}_{5,4}\text{O}$ USNPs. Literature has reported that ROS could activate the MAPK signaling pathway to induce renal cell apoptosis and

the release of local or systemic inflammatory mediators to aggravate renal injury^{58–60}. Notably, the MAPK signaling pathway (Supplementary Fig. 29) was significantly inhibited after $\text{Cu}_{5,4}\text{O}$ USNPs treatment, indicating that $\text{Cu}_{5,4}\text{O}$ USNPs could alleviate renal injury via inhibiting the MAPK signaling pathway through decreasing the ROS level. In addition, it has been reported that

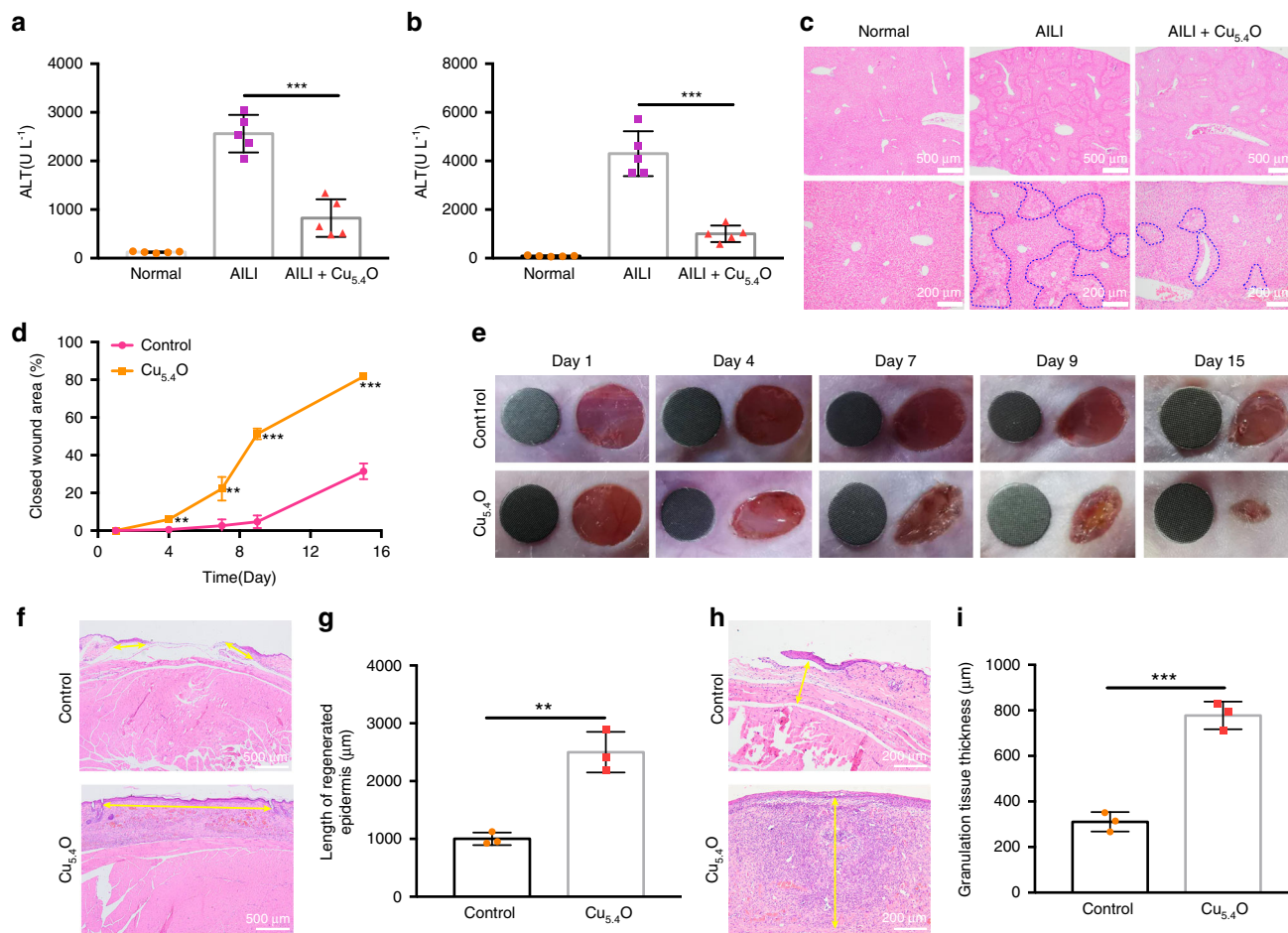


Fig. 8 Therapeutic efficiency of $\text{Cu}_{5,4}\text{O}$ USNPs on AILI and wound healing. Serum levels of **a** AST and **b** ALT in AILI mice at 24 h after different treatment. **c** H&E staining of liver tissues from each group. Blue dashed lines indicate the range of hepatic necrosis. The percentage of **d** closed diabetic wound area and representative macroscopic appearance of **e** diabetic wound at different time points. A 6-mm-diameter standard green disc was used as the reference when taking photos. **f** Representative histological images and **g** quantitative determination of the length of regenerated epidermis on day 15 post-surgery. The yellow double-headed arrows indicate the regenerated epidermis. **h** Representative histological images and **i** quantitative measurement of granulation tissue thickness on day 15 post-surgery. The yellow double-headed arrows indicate the granulation tissue. In **a**, **b**, data represent means \pm s.d. from five independent replicates (one-way ANOVA). In **d**, **g** and **i**, data represent means \pm s.d. from three independent replicates (Student's *t*-test, ** $P < 0.01$; *** $P < 0.001$). Source data are provided as a Source Data file.

ROS could promote the production of proinflammatory cytokines (e.g. $\text{TNF-}\alpha$, $\text{IL-1}\beta$, IL-6)^{58,61} and that $\text{TNF-}\alpha$ could further trigger a strong cascade inflammatory response throughout the $\text{TNF-}\alpha$ /MAPK and $\text{TNF-}\alpha$ /NF- κB signaling pathways, resulting in an excessive inflammation response and more pronounced renal damage⁶². Notably, the TNF signaling pathway (Supplementary Fig. 30) was also significantly inhibited after $\text{Cu}_{5,4}\text{O}$ USNP treatment, suggesting that other therapeutic mechanisms of renal protection occur through inhibiting the TNF signaling pathway.

We also investigated the impact of $\text{Cu}_{5,4}\text{O}$ USNPs on the expression of genes related to oxidative stress. As shown in Fig. 9e, f, several important antioxidant genes, including *SOD1*, *SOD2*, *SOD3*, *GPX1*, *GPX3*, *GPX4*, *GPX6*, and *CAT*, were significantly upregulated after the $\text{Cu}_{5,4}\text{O}$ USNPs treatment. The expression trend of the *SOD* and *HMOX1* genes was consistent with the aforementioned result regarding the *SOD* (Fig. 7h) and HO-1 proteins level (Fig. 7j) in AKI mice, respectively. Moreover, the genes associated with oxidative stress that significantly changed after $\text{Cu}_{5,4}\text{O}$ USNP treatment were used in the protein–protein interactions network analysis (Fig. 9g). We discovered that the neighboring proteins connected to the leading proteins contained *SOD1*, *SOD3*, *CAT*, etc., indicating that these genes play an

important role in ROS scavenging after $\text{Cu}_{5,4}\text{O}$ treatment. As shown in Fig. 9h, the mRNA expression levels of antioxidant genes in $\text{Cu}_{5,4}\text{O}$ -treated mice kidneys were significantly higher than those of the corresponding control group, confirming that $\text{Cu}_{5,4}\text{O}$ USNPs could maintain a high expression of antioxidant genes in AKI by protecting renal cells from ROS damage.

The phosphorylation of NF- κB and I κB were significantly enhanced in the AKI mice (Fig. 9i, Supplementary Fig. 31), indicating the activation of NF- κB signaling pathway in AKI. Besides, the phosphorylation of NF- κB and I κB were significantly decreased in the $\text{Cu}_{5,4}\text{O}$ USNPs group, indicating that the NF- κB signaling pathway was inhibited after $\text{Cu}_{5,4}\text{O}$ USNPs treatment, which was also in accordance with the aforementioned transcriptomics analysis result (Supplementary Fig. 30). We also detected the downstream inflammatory factors of the NF- κB signaling pathway. As shown in Fig. 9j–m, $\text{Cu}_{5,4}\text{O}$ USNPs could significantly reduce the serum and tissue levels of $\text{TNF-}\alpha$ and $\text{IL-1}\beta$, indicating that $\text{Cu}_{5,4}\text{O}$ USNPs could protect kidney tissues from oxidative stress by inhibiting the production of excessive proinflammatory factors.

Additionally, we found that several important genes related to tissue repair, including fibroblast growth factor 10 (*FGF10*),

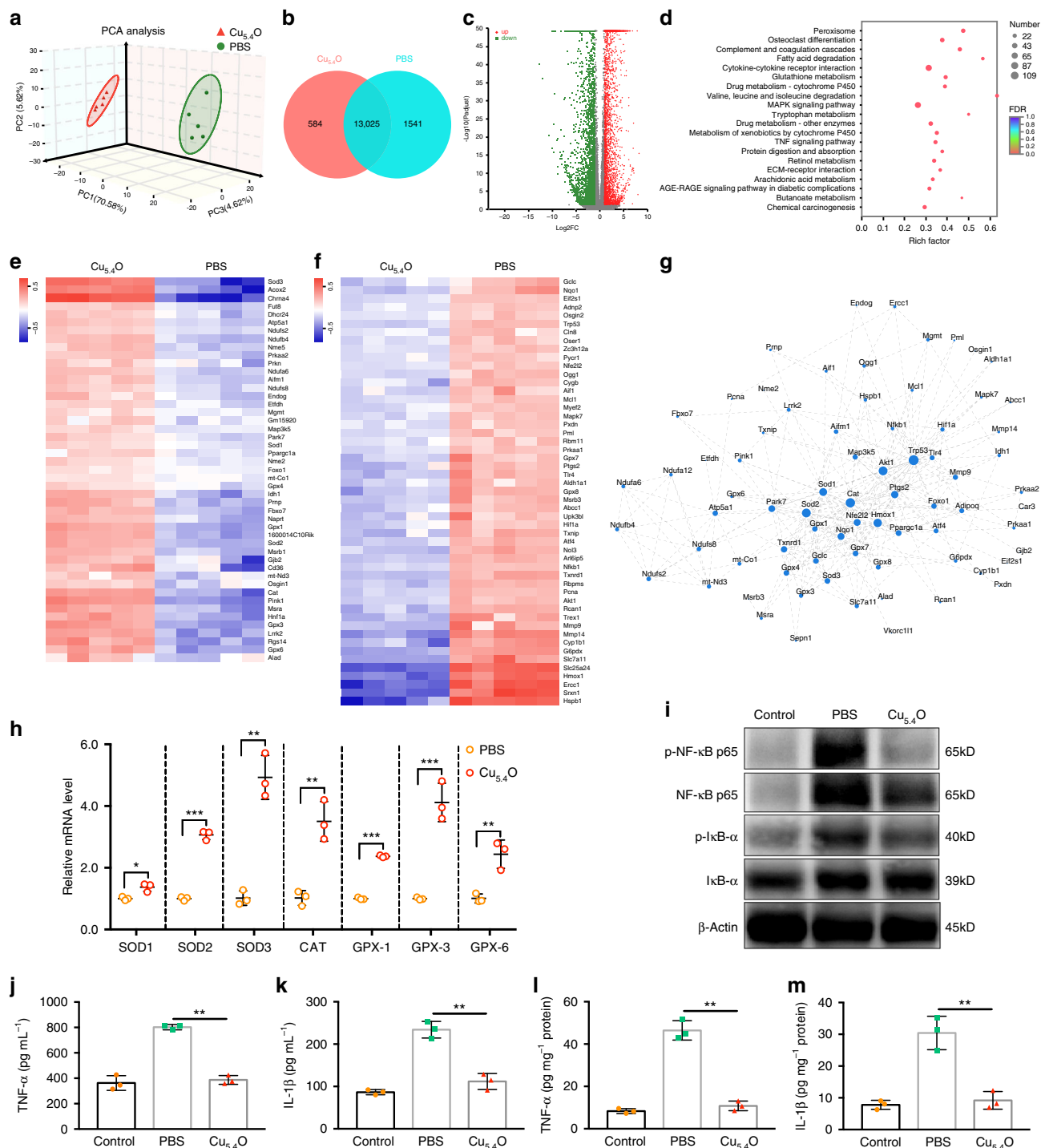


Fig. 9 Therapeutic mechanisms of Cu_{5.4}O USNPs on AKI. **a** Principal component analysis (PCA) was performed based on differentially expressed genes from the kidneys of two groups. Each data point corresponds to the PCA analysis of each sample. **b** Venn diagram of the transcriptomic profiles between Cu_{5.4}O USNPs and PBS groups. **c** Volcano plots showing the identified upregulated and downregulated genes by Cu_{5.4}O USNPs. **d** KEGG pathway enrichment analysis of the identified differentially expressed genes. The 20 most significantly enriched pathways are shown. Heat maps of significantly **e** upregulated and **f** downregulated genes involved in the oxidative stress after Cu_{5.4}O USNPs treatment (fold change ≥ 2 and $P < 0.05$). **g** Protein-protein interaction network of differentially expressed genes involved in the oxidative stress. **h** qRT-PCR analysis of the mRNA expression levels of antioxidant genes. **i** Western blot analysis of the expression of phospho-NF- κ B p65, total NF- κ B p65, phospho-I κ B- α , and total I κ B- α in kidney tissues of AKI mice. Serum levels of **j** TNF- α and **k** IL-1 β . Renal levels of **l** TNF- α and **m** IL-1 β . In **h**, **j**–**m**, data represent means \pm s.d. from three independent replicates, and P values were calculated by Student's t -test (**h**) and one-way ANOVA (**j**–**m**). * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$. Source data are provided as a Source Data file.

hepatocyte growth factor (*HGF*), *NOTCH1*, and wingless-type MMTV integration site family member 7A (*WNT7A*), were significantly upregulated after the Cu_{5,4}O USNPs treatment (Supplementary Fig. 32a, b). The protein–protein interaction network further confirmed the importance of these genes in tissue repair and regeneration, as shown in Supplementary Fig. 32c. The overall results indicated in addition to the ROS scavenging ability, Cu_{5,4}O USNPs may also promote the expression of genes related to renal repair and regeneration.

In summary, we have presented uniform and stable Cu_{5,4}O USNPs that mimic an intracellular antioxidant enzyme-based defense system. The distinctive advantages of Cu_{5,4}O USNPs are the ultrasmall particle size, rapid renal clearance, high biocompatibility, and broad ROS scavenging abilities. Both in vitro and in vivo experiments demonstrated excellent biocompatibility and cytoprotective effects of Cu_{5,4}O USNPs against ROS-mediated damage. Cu_{5,4}O USNPs could be applicable to various ROS-related pathological conditions such as AKI, ALI, and diabetic wound. Taken together, the synthesized ultrasmall Cu_{5,4}O USNPs with robust ROS scavenging abilities and excellent biocompatibility could represent a promising antioxidant for the treatment of AKI and other oxidative stress-related diseases. We expect that our findings will promote the development of nanomaterials with multiple enzyme-mimicking properties and enable further clinical applications of copper-based ROS scavengers in biomedical treatment and research.

Methods

Materials. Cupric chloride (CuCl₂), L-ascorbic acid (AA), and sodium hydroxide (NaOH) were purchased from J&K Scientific (Beijing, China). 3,3',5,5'-Tetramethylbenzidine (TMB) and 2,2'-azino-bis(3-ethylbenzthiazoline-6-sulfonate) (ABTS) was bought from Macklin (Shanghai, China). Acetic acid (HAc), sodium acetate (NaAc), and potassium persulfate (K₂S₂O₈) was obtained from Sigma-Aldrich (Shanghai, China). Thiol-polyethylene glycol-OH (SH-PEG-OH) (Mw: 1000 Da) was purchased from Yare Biotech. Inc. (Shanghai, China). Catalase (purified powder from bovine liver), superoxide dismutase (purified powder from bovine erythrocytes), and peroxidase (purified powder from horseradish roots) were from Sangon Biotech (Shanghai, China). Phosphate-buffered saline (PBS, pH 7.4, Na₂HPO₄–NaH₂PO₄, 10 mM) solution was prepared in the laboratory. All chemicals and reagents were of analytical grade and used as received without further purification. Ultrapure water (18.2 MΩ cm⁻¹ at 25 °C) purified by a Milli-Q system was used throughout the experiment.

Instrumentation. TEM imaging was conducted on a Tecnai G220 (Shimadzu, Japan) at 200 keV. The sample was prepared by dispersing a small amount of freeze-dried powder in the PBS. Then, the suspension was dropped on 230 mesh copper TEM grids covered with thin amorphous carbon films. Fourier Transform infrared spectroscopy (FTIR) spectra were measured by a FTIR-8300 series spectrometer (Shimadzu, Japan) in the range of 4000–400 cm⁻¹. The fluorescence spectra were determined by using an F-2500 spectrofluorometer (Hitachi, Japan). The UV-vis spectra were obtained using a Hitachi U-3010 spectrometer (Hitachi, Japan). A dynamic light scattering (DLS) particle size analyzer (Malvern 2000, USA) was used to determine the hydrophilic diameters of the particles. XPS and X-ray Auger electron spectroscopy measurement were performed by an ESCALAB 250 Xi Mg (Thermo Scientific, Japan) X-ray resource. Crystal structure and oxidation state of Cu_{5,4}O USNPs were analyzed using X-ray diffractometer (XRD, Bruker AXS D8) with a scan rate of 1 s per step with a step size of 0.02. The concentration of Cu was detected by inductively coupled plasma-atomic emission spectrometry (ICP-AES) (Thermo Scientific, iCAP 7400, USA). The concentration of Cu_{5,4}O USNPs used in following studies is calculated based on Cu element. The EPR spectroscopy signal was obtained on a Bruker A300 (X-band) spectrometer (Bruker, Germany). All the measurements were performed at room temperature if not specially mentioned.

Synthesis of fluorescent Cu_{5,4}O USNPs. The Cu_{5,4}O USNPs was synthesized according to previous report with some modifications⁴⁰. In a typical preparation process, 10 mM CuCl₂ powders were dissolved in 50 mL deionized water and stirred for 10 min at 80 °C in an oil bath with magnetic stirring. Then, L-ascorbic acid aqueous solution (100 mM, 50 mL) was added slowly to the above CuCl₂ solution. Afterwards, the pH of the solution was adjusted to 8.0–9.0 using NaOH solution (1 M). The mixture was kept at 80 °C for 12 h with constantly stirring. After reaction, the larger aggregates were removed by centrifugation (6577 × g, 15 min), and then the supernatant was dialyzed against water (Mw cutoff: 10,000

Da) for 2 days to remove small molecules. Purified Cu_{5,4}O USNPs were concentrated with centrifugation.

Surface functionalization of Cu_{5,4}O USNPs by SH-PEG-OH. In total, 0.1 g Cu_{5,4}O USNPs were dispersed into 5 mL water under ultrasonication for 1 min and then stirred for 1 h. In all, 0.1 wt% of SH-PEG-OH was added. The reaction mixture was stirred for another 24 h, dialyzed to remove unreacted SH-PEG-OH molecules, and then concentrated by centrifugal ultrafiltration.

H₂O₂ scavenging activity of Cu_{5,4}O USNPs. H₂O₂ scavenging capacity of Cu_{5,4}O USNPs was tested by the Hydrogen Peroxide Detection Kit (Nanjing Jiancheng Bioengineering Institute, China). H₂O₂ reacts with ammonium molybdate to form a stable yellow complex, which displays an absorbance peak at 405 nm. Various concentrations of Cu_{5,4}O USNPs (25–200 ng mL⁻¹) were incubated with 2 mM H₂O₂ at 37 °C for 2 h, respectively. After reaction, the concentration of remaining H₂O₂ was determined according to the manufacturer's instructions, and the H₂O₂-eliminating capacity was calculated.

·OH scavenging activity of Cu_{5,4}O USNPs. The TMB chromogenic method was performed for ·OH scavenging activity test. The ·OH was generated by the classical Fenton reaction between H₂O₂ and Fe²⁺, which can convert the TMB to an oxidized TMB (oxTMB) with a characteristic absorption at 652 nm. Therefore, the concentration of remaining ·OH can be determined via monitoring the absorption at 652 nm of oxTMB. In detail, The working test solutions containing 250 μM TMB, 2 mM H₂O₂, 1 mM FeSO₄ and different concentrations of Cu_{5,4}O USNPs (25–175 ng mL⁻¹) in HAc/NaAc buffer (0.5 M, pH 4.5) were prepared in the dark and rest for 5 min. Afterwards, the absorbance peak in 652 nm of the solution was monitored with a UV-vis spectroscopy.

The EPR spectroscopy signal was measured by a Bruker A300 spectrometer (Bruker, Germany). Typically, 2 mM H₂O₂, 20 μM FeSO₄, 100 mM DMPO, and different concentration of Cu_{5,4}O USNPs (0, 50, 150 ng mL⁻¹, respectively) were added into the HAc/NaAc buffer (0.5 M, pH 4.5); EPR signal was detected immediately.

O₂⁻ scavenging activity of Cu_{5,4}O USNPs. The superoxide anion (O₂⁻) scavenging activity was assessed using a superoxide anion assay kit (Nanjing Jiancheng Bioengineering Institute, Nanjing, China) according to the manufacturer's instructions. Different concentrations of Cu_{5,4}O USNPs (0–150 ng mL⁻¹) were added to the working solution. The absorbance at 550 nm was measured using a multiple plate reader after standing for 10 min.

ABTS radical scavenging activity of Cu_{5,4}O USNPs. The evaluation of ABTS radical scavenging activity was based on the method reported by Wang et al.⁶³. Briefly, the ABTS radicals were generated by reacting 7 mM ABTS stock solution with 2.45 mM potassium persulfate in the dark for 16 h. Then, the ABTS radical solution was diluted by PBS to reach a proper absorbance at 734 nm. Two milliliters Cu_{5,4}O USNPs solutions (0, 50, 75, 100, 125, 150 ng mL⁻¹, respectively) were mixed with 2 mL ABTS solution and placed in dark for 10 min. Then the absorbance peak at 734 nm was monitored with a UV-vis spectroscopy. The ABTS radical scavenging abilities were calculated as follows:

$$\text{ABTS scavenging ratio(\%)} = \left(\frac{A_{\text{control}} - A_{\text{sample}}}{A_{\text{control}}} \right) \times 100, \quad (1)$$

where A_{control} is the absorbance of a standard without any radical scavengers, and A_{sample} is the absorbance after the reaction with the radical scavengers, respectively.

CAT-like activity of Cu_{5,4}O USNPs. The CAT-like activity of the Cu_{5,4}O USNPs and the steady-state kinetic were assayed using a fluorescence spectroscopy^{19,36}. Typically, TPA is a non-fluorescent compound, which can react with ·OH, decomposed from H₂O₂, to produce a fluorescent aromatic hydroxylated product (2-hydroxyterephthalic acid) with excitation/emission peaks at 315/425 nm respectively. In the presence of CAT or CAT analogs, the H₂O₂ decomposed into H₂O and O₂ could not produce the fluorescent 2-hydroxyterephthalic acid. Hence, the CAT-like activity of the Cu_{5,4}O USNPs was investigated by monitoring the fluorescence signal of the system. The steady-state kinetic assay of Cu_{5,4}O USNPs and CAT were performed to confirm the enzymatic catalysis mechanism, by varying the concentration of H₂O₂ (0–6 mM) in the presence of Cu_{5,4}O USNPs (1000 ng mL⁻¹) and H₂O₂ (0–1 mM) in the presence of CAT. Firstly, 200 μL 1000 ng mL⁻¹ of Cu_{5,4}O USNPs solution or 20 U mL⁻¹ CAT was added into the PBS (10 mM, pH 7.4), different concentrations of H₂O₂ were added and incubated at 40 °C for 5 h. Finally, the TPA in 2 mM NaOH was added and the fluorescence spectra of the resultant solution were recorded under an excitation of 315 nm, and the variation of fluorescence emission at 425 nm was adopted to quantify remaining H₂O₂. Apparent kinetic parameters were calculated on the flowing basis of the Michaelis–Menten equation⁶⁴.

$$V_0 = \frac{V_{\text{max}}[S]}{[S] + K_m}, \quad (2)$$

where V_0 is the initial catalytic rate, V_{\max} is the maximum rate conversion, which is obtained when the catalytic sites on the enzyme are saturated with substrate concentration, and K_m is the apparent Michaelis–Menten equation. Maximum initial velocity (V_{\max}) and Michaelis–Menten constant (K_m) were obtained using Lineweaver–Burk plots⁶⁵.

SOD-like activity of Cu_{5,4}O USNPs. The SOD-like activity of Cu_{5,4}O USNPs was determined by formazan formation using a SOD assay kit (WST-1 method) (Nanjing Jiancheng Bioengineering Institute, Nanjing, China)¹⁹. Briefly, O₂^{•-} was generated through the oxidation of xanthine by xanthine oxidase (XO), which can convert the WST-1 into WST-1 formazan with a characteristic absorption at 450 nm. The formazan concentration was determined at 450 nm using a multiple plate reader. The SOD-like activity of Cu_{5,4}O USNPs was further confirmed with EPR spectroscopy. Briefly, a series of samples containing xanthine (5 mM) and xanthine oxidase (0.5 U mL⁻¹) in 10 mM PBS and incubated for 10 min at 37 °C. Different Cu_{5,4}O USNPs solutions (0, 20, 100, 1000 ng mL⁻¹) were added and then the EPR signals were recorded immediately.

GPx-like activity of Cu_{5,4}O USNPs. The GPx-like activity of Cu_{5,4}O USNPs was estimated using a GPx assay kit (Solarbio, Shanghai, China) according to the manufacturer's instructions. Glutathione (GSH) can react with 5,5'-Dithiobis-2-nitrobenzoic acid (DTNB) to form a compound with characteristic absorption at 412 nm, which could be monitored using a UV-Vis spectroscopy. During the test, GSH was oxidized to oxidized glutathione (GSSG). The decrease of GSH concentration is proportional to the catalytic activity of Cu_{5,4}O USNPs.

O₂ generation from H₂O₂ catalyzed by Cu_{5,4}O USNPs and CAT. A commercial O₂ probe (Oxford-optronix, Oxford, UK) was utilized to measure the real-time O₂ level according to the manual instructions. Ten milliliters of PBS was poured into a small three-necked flask and stirred at 37 °C. The flask was then sealed with parafilm and rubber stoppers, and inserted with two needles for gassing. 95% N₂/5% CO₂ gas mixture was bubbled into the flask for 20 min to create hypoxic condition. Cu_{5,4}O USNPs (500 μL, 200 ng mL⁻¹) and CAT (500 μL, 20 U mL⁻¹) solutions were injected to the mixture, while PBS was used in the control group. Then H₂O₂ (500 μL, 2 mM) was injected and the amount of generated O₂ was monitored at predetermined time intervals.

Cell culture. The human embryonic kidney 293 (HEK293) cell line was purchased from the American Type Culture Collection (ATCC). The HEK293 cells were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum (FBS), 100 μg mL⁻¹ streptomycin, and 100 U mL⁻¹ penicillin at 37 °C in an incubator supplied with a humidified atmosphere of 5% CO₂.

In vitro ROS scavenging using Cu_{5,4}O USNPs. To investigate the ROS scavenging ability of Cu_{5,4}O USNPs in cells, HEK293 cells were seeded into 96-, 48-, and 24-well plates at the density of 1 × 10⁴ cells per well, 5 × 10⁴ cells per well, and 10 × 10⁴ cells per well, respectively. After 24 h incubation, Cu_{5,4}O USNPs with different concentrations (0–50 ng mL⁻¹) were added to each group of wells and incubated for 30 min. Then, cells were treated with 250 μM H₂O₂ and further incubated at 37 °C for 24 h. Cells seeded in 96-well plates were incubated with cell counting kit-8 (CCK-8, Dojindo, Japan) to detect the cell viability. Wells without the addition of H₂O₂ were regarded as the negative control.

Furthermore, cells seeded in 48-well plates were stained with Annexin V-FITC apoptosis detection kit (C1062, Beyotime, China) to detect the ratio of apoptotic and necrotic cells⁶⁶. In brief, HEK293 cells in a well were collected, washed with cold PBS, and re-suspended in 195 μL binding buffer after the aforementioned incubation with H₂O₂ for 24 h. Then, 5 μL Annexin V-FITC and 10 μL PI were sequentially added to the cell suspension and incubated at room temperature in dark for 15 min. After that, cells were analyzed by the Attune Acoustic Focusing Cytometer (Life Technologies, USA). At least 50,000 cells were analyzed in each sample.

2',7'-Dichlorofluorescein diacetate (DCFH-DA, D6883, Sigma-Aldrich, USA), an oxidation sensitive fluorescent dye, was used to detect the intracellular ROS level according to the literature²⁵. Briefly, DCFH-DA is a non-fluorescent chemical compound which could diffuse through cell membrane freely and could be hydrolyzed by intracellular esterase to DCFH. The non-fluorescent DCFH could be oxidized by the intracellular ROS to fluorescent DCF. Therefore, the quantity of intracellular ROS is correlated with the fluorescent intensity of DCF. After the aforementioned incubation with H₂O₂ for 24 h, cells were gently rinsed thrice with serum-free medium to remove the free Cu_{5,4}O USNPs. Then, a final concentration of 10 μM of DCFH-DA in serum-free medium was added to the cells and incubated in dark at 37 °C for 30 min. Afterwards, the cells were washed with serum-free medium thrice to remove unloaded DCFH-DA probe, then were imaged using a laser confocal microscope (Zeiss LSM780, Germany), and were subjected to a flow cytometry analysis to quantify the intracellular ROS levels respectively.

To determine the location of Cu_{5,4}O USNPs in the cells, HEK293 cells (1 × 10⁶ cells) were collected after incubation with 50 ng mL⁻¹ of Cu_{5,4}O USNPs at 37 °C for 24 h. The collected cells were fixed with 2.5% glutaraldehyde at 4 °C for 2 h, then post-fixed with 1% osmium tetroxide and prepared for TEM observation.

In vitro biocompatibility evaluation of Cu_{5,4}O USNPs. The cytotoxicity of Cu_{5,4}O USNPs was determined by the CCK-8 assay in vitro. Briefly, HEK293 cells were seeded into 96-well culture plates at the density of 1 × 10⁴ cells per well and incubated at 37 °C in an incubator with 5% CO₂ for 24 h. Afterwards, the cell culture medium was aspirated and fresh culture media containing various concentrations of Cu_{5,4}O USNPs (0–200 ng mL⁻¹) were added. After 24 or 48 h incubation, cells were gently washed once with sterile PBS and then treated with 100 μL fresh culture medium and 10 μL CCK-8 solution, and further incubated at 37 °C for 2 h. The cell viability was then quantified by measuring the absorbance value at 450 nm by a microplate reader (Thermo Varioskan Flash, USA).

The hemolysis assay was performed on the basis of previously reported methods with some modifications⁶⁷. All the animal experiments were carried out under the approval of the Institutional Animal Care and Use Committee of the Third Military Medical University (Army Medical University). Fresh whole-blood samples were collected from the orbital venous of healthy Sprague-Dawley rats. The collected blood samples were centrifuged for 15 min at 231 × g to collect erythrocytes and gently washed thrice with saline solution. Then, 3.67 mL of saline solution was added to erythrocytes collected from 1 mL blood. Afterwards, 100 μL of diluted erythrocytes suspension was mixed with 1 mL Cu_{5,4}O USNP dispersion at various concentrations (50–5000 ng mL⁻¹). The mixed dispersions were incubated for 3 h at 37 °C and then centrifuged for 15 min at 13800 × g before observing and recording the hemolysis phenomenon. The hemolysis ratio was quantified by measuring the absorbance value of supernatant at 540 nm with a microplate reader. Deionized water and saline solution were used as the positive and negative control, respectively.

In vivo biocompatibility evaluation of Cu_{5,4}O USNPs. To evaluate the biocompatibility of Cu_{5,4}O USNPs in vivo, BALB/c mice (aged 8–10 weeks, 20–25 g) were intravenously administrated with Cu_{5,4}O USNPs at a single dose of 4 μg kg⁻¹. The mice injected with PBS were used as the control group. One day post injection, the blood samples were collected for complete blood panel analysis and serum biochemistry test. The serum biochemistry test included two important indicators of hepatic function as aspartate aminotransferase (AST) and alanine aminotransferase (ALT), and two indicators of kidney function as blood urea nitrogen (BUN) and creatinine (CRE). Serum IL-6 and TNF-α levels were quantified by the ELISA assay. One day and thirty days post injection, the mice were sacrificed to harvest major organs (including heart, liver, spleen, lung, and kidney) for hematoxylin and eosin (H&E) staining and histological analysis.

To evaluate the in vivo toxicity of Cu_{5,4}O USNPs after repeated administration, BALB/c mice (aged 8–10 weeks, 20–25 g) were intravenously administrated with Cu_{5,4}O USNPs at a dose of 4 μg kg⁻¹ every day for seven consecutive days. On eighth day, mice were sacrificed to harvest major organs for H&E staining and collect blood samples for complete blood panel analysis and serum biochemistry test.

To detect the accumulation of Cu_{5,4}O USNPs in the major organs after repeated administrations, BALB/c mice (aged 8–10 weeks, 20–25 g, $n = 8$) were intravenously administrated with Cu_{5,4}O USNPs at a dose of 4 μg kg⁻¹ every day for seven consecutive days. On the eighth day, mice were sacrificed to harvest major organs. The organs of four mice were weighed, homogenized, and then dissolved in aqua regia to calculate the percentage of injected dose per gram of tissue (%ID g⁻¹) by ICP-AES. The organs of the other four mice were fixed with the mixture of 2.5% glutaraldehyde and 4% formaldehyde for 24 h, and then post-fixed with 1% osmium tetroxide and prepared for TEM observation.

AKI models in mice. The glycerol-induced AKI model was established according to the previously reported protocol¹⁰. Briefly, female BALB/c mice (aged 8–10 weeks, 20–25 g) were deprived of water but given free access to food for 15 h. After water deprivation, the two hindlimbs of mice were equally intramuscularly injected with 50% glycerol at a dose of 8 mL kg⁻¹. After that, all the mice had free access to water and food. Symptoms of AKI, such as a lack of activities and decreased urine output, could be observed in a few hours after glycerol injection.

The cisplatin-induced AKI model was also established according to previously reported method⁶⁸. In brief, female BALB/c mice (aged 8–10 weeks, 20–25 g) were given one time intraperitoneal injection of cisplatin (BP809, Sigma-Aldrich, USA) at a dose of 20 mg kg⁻¹. Mice in the treatment group were intravenously injected with Cu_{5,4}O USNPs at 2 μg kg⁻¹ dose 2 h after intraperitoneal cisplatin injection. Mice received saline injection was used as control. Three days post injection, mice were sacrificed to collect blood and renal tissues for kidney function analysis and histological analysis, respectively.

Pharmacokinetics and biodistribution of Cu_{5,4}O USNPs. To evaluate the blood circulation half-life of Cu_{5,4}O USNPs, BALB/c mice ($n = 3$) with established glycerol-induced AKI model were intravenously injected with Cu_{5,4}O USNPs at a dose of 2 μg kg⁻¹. At different time points post injection (10 min, 1, 2, 4, 8, 24, 48, and 72 h), 10 μL of whole-blood samples were collected from the mouse tail vein. The collected blood samples were dissolved in aqua regia and the concentrations of NPs were quantified by ICP-AES. A two-compartment pharmacokinetic model was utilized to calculate the pharmacokinetics parameters of Cu_{5,4}O USNPs. Simultaneously, the urine and feces of mice were collected at different time points and

dissolved in aqua regia to quantify the content of Cu by ICP-AES. Furthermore, the collected urine was also diluted with PBS and dropped on a carbon-coated copper grid to detect the presence of Cu_{5,4}O USNPs in the urine under TEM (JEOL JEM-1400, Japan).

To detect the biodistribution of Cu_{5,4}O USNPs in the major organs, BALB/c mice ($n = 3$) with established glycerol-induced AKI model were intravenously injected with Cu_{5,4}O USNPs at a dose of 2 $\mu\text{g kg}^{-1}$. One day after injection, mice were sacrificed to harvest major organs including heart, liver, spleen, lung, and kidney. The tissues were weighed, homogenized, and then dissolved in aqua regia to calculate the percentage of injected dose per gram of tissue (%ID g^{-1}) by ICP-AES.

To study the accumulation of Cu_{5,4}O USNPs in the kidney at different time periods, healthy female BALB/c mice or the ones with established AKI model ($n = 3$) were intravenously injected with Cu_{5,4}O USNPs at a dose of 2 $\mu\text{g kg}^{-1}$. At desired time points (6, 12, 24 h), three mice in each group were sacrificed. Kidney tissues were harvested, weighed, homogenized, and dissolved in aqua regia to calculate the %ID g^{-1} of Cu by ICP-AES. The same experiment setup was applied for another batch of mice to investigate the biodistribution of Cu_{5,4}O USNPs in the GBM. At different time points (2, 24, 72 h), three mice in each group were sacrificed and a small piece of kidney tissues was fixed with the mixture of 2.5% glutaraldehyde and 4% formaldehyde for 24 h, then post-fixed with 1% osmium tetroxide and prepared for TEM observation.

In vivo therapeutic outcome of Cu_{5,4}O USNPs on AKI mice. BALB/c mice with established AKI model were randomly divided into five groups ($n = 4$): received PBS injection (control group), NAC (8 mg kg^{-1}), NAC (40 mg kg^{-1}), NAC (160 mg kg^{-1}), and received with Cu_{5,4}O USNPs at a dosage of 2 $\mu\text{g kg}^{-1}$, respectively. The drug or NPs were administrated once. The body weight variations in each group after treatment were monitored for 24 h.

After 24 h post injection, mice were sacrificed to collect blood samples for detecting the BUN and CRE levels. The left kidneys were harvested and homogenized for the detection of renal biomarkers. The level of SOD in the kidney was detected with a SOD assay kit (19160; Sigma-Aldrich, USA). The expression levels of two important kidney injury biomarkers, heme oxygenase-1 (HO-1) and kidney injury molecule-1 (KIM-1), were detected with HO-1 (ab204524, Abcam, USA) and KIM-1 (ab213477, Abcam, USA) ELISA kits, respectively.

The right kidneys were cut into two equal sections. One section was fixed with 4% paraformaldehyde and embedded in paraffin for H&E staining. The other section was frozen and embedded in optimum cutting temperature (O.C.T.) specimen matrix (Sakura, Leiden, The Netherlands) for cryostat sectioning at -20°C . The renal tissue was sectioned into approximately 5 μm thickness. Frozen renal tissue sections were stained with DAPI (C1005, Beyotime, China) and dihydroethidium (DHE, D7008, Sigma-Aldrich, USA) at 37°C for 30 min, gently washed thrice with PBS to remove excessive dyes and then imaged under a fluorescence microscope to qualitatively detect the ROS levels in the renal tissues.

The same experiment setup was applied for another batch of mice ($n = 10$) to draw the survival curves of AKI mice within 2 weeks. For therapeutic mechanism investigation of Cu_{5,4}O USNPs, after 24 h post treatment, another batch of mice ($n = 3$) were sacrificed to collect serum and kidney tissues for measurement of inflammatory factors by ELISA kit.

In vivo therapeutic effect of Cu_{5,4}O USNPs on AILI mice. The acetaminophen (APAP)-induced acute liver injury model was established according to the previously reported protocol⁶⁹. Briefly, female C57BL/6 mice (aged 8–10 weeks, 18–20 g) were deprived of food but given free access to water for 15 h. Then, 10 mice were intraperitoneally injected with APAP (dissolved in warm saline, 15 mg mL^{-1}) at a dose of 300 mg kg^{-1} and randomly divided into two groups ($n = 5$). Mice ($n = 5$) without APAP injection served as control. Afterwards, all the mice had free access to water and food. Mice in the treatment group were intravenously injected with Cu_{5,4}O USNPs at 100 $\mu\text{g kg}^{-1}$ dose 2 h after APAP injection. At 24 h post injection, mice were sacrificed to collect blood and liver tissues for liver function test and histological analysis, respectively.

In vivo therapeutic effect of Cu_{5,4}O USNPs on wound healing. To investigate the effect of Cu_{5,4}O USNPs on diabetic wound healing, the STZ-induced diabetic mice model was established according to the previous literature⁷⁰. Briefly, male BALB/c mouse (aged 8–10 weeks, 20–25 g, $n = 3$) were intraperitoneal injected with 100 mg kg^{-1} streptozotocin (STZ, Sigma-Aldrich, USA) for six consecutive days. Then all mouse were provided with normal food and water. The glucose level of mice was then monitored everyday from the tail venous blood using a blood glucose meter (Roche Diagnostics, Shanghai, China). Mice with sustained blood glucose levels exceeding 250 mg dL^{-1} were considered diabetic mice. These diabetic mice were used for further wounding model at 2 weeks after the initiation of STZ treatment. Full-thickness wounds were simultaneously created in the dorsal skin using a sterile 6-mm diameter punch. A green round marker (6-mm diameter) was placed beside each wound to represent the initial wound area and the wounds were photographed immediately using a digital camera. Afterwards, each wound in the treatment group was topically administrated with 20 μL of Cu_{5,4}O USNPs at 400 ng mL^{-1} . Wounds in the control group were treated with 20 μL of PBS instead.

Then, the wounds were covered with a piece of biological membrane (NPWT-1, Negative Pressure Wound Therapy Kit, China). The wounds were photographed and Cu_{5,4}O was topically administrated at days 1, 4, 7, 9, 15 post-surgery, respectively. Wound areas were measured using ImageJ software. The wound healing rate was calculated based on the following formula:

$$\text{Wound healing rate}\% = (I - R) / I \times 100\%, \quad (3)$$

where I represented the initial wound area and R represented the remaining wound area on the determined day post-surgery.

At day 15 post-surgery, mice were sacrificed to harvest wound tissues for histological analysis. The wound tissues were fixed with 4% paraformaldehyde and embedded in paraffin for H&E staining. The length of regenerated epidermis and the thickness of granulation tissues were quantified using ImageJ software.

Transcriptome analysis of AKI mice. BALB/c mice with established AKI model were randomly divided into two groups: received PBS injection (control group, $n = 5$) and received with Cu_{5,4}O USNPs at a dosage of 2 $\mu\text{g kg}^{-1}$ (experimental group, $n = 5$), respectively. After 24 h post injection, mice were sacrificed to collect the kidneys. Total RNA of kidney tissues were prepared using Trizol Reagent (Invitrogen, USA) according to the manufacturer's instructions and genomic DNA was removed using DNase I (TaKaRa, Japan). Then RNA quality was determined by 2100 Bioanalyser (Agilent) and quantified using the ND-2000 (NanoDrop Technologies). Only high-quality RNA sample ($\text{OD}_{260/280} = 1.8\text{--}2.2$, $\text{OD}_{260/230} \geq 2.0$, $\text{RIN} \geq 6.5$, $28\text{S}:18\text{S} \geq 1.0$, $>2 \mu\text{g}$) was used to construct sequencing library. The RNA purification, reverse transcription, library construction, and sequencing were performed at Majorbio Bio-pharm Biotechnology Co., Ltd (Shanghai, China) using Illumina HiSeq X10 (Illumina, San Diego, CA) according to the manufacturer's instructions. The raw paired end reads were trimmed and quality controlled by SeqPrep (<https://github.com/jstjohn/SeqPrep>) and Sickle (<https://github.com/najoshi/sickle>) with default parameters. Then clean reads were separately aligned to reference genome with orientation mode using TopHat (<http://tophat.cbcb.umd.edu/>, version 2.1.1) software.

For bioinformatics analysis, the expression level of each transcript was calculated according to the fragments per kilobase of exon per million mapped reads (FPKM) method. RSEM (<http://deweylab.biostat.wisc.edu/rsem/>) was used to quantify gene abundances. DEGs were identified using R statistical package software DESeq2 (<http://bioconductor.org/packages/stats/bioc/DESeq2.html>) (fold change ≥ 2 and P value < 0.05) with a false discovery rate (FDR) cutoff < 0.05 . KEGG functional enrichment analysis was performed to identify which DEGs were significantly enriched in KEGG signaling pathways at Bonferroni-corrected P value < 0.05 compared with the whole-transcriptome background. KEGG enrichment analysis was performed by KOBAS 2.1.1 (<http://kobas.cbi.pku.edu.cn/download.php>). Besides, protein–protein interactions of genes were analyzed by Search Tool for the Retrieval of Interacting Genes/Proteins (STRING) algorithm (<http://www.string-db.org/>).

Quantitative real-time PCR and Western blot analysis. The total RNA from kidney tissue was extracted using Trizol Reagent. cDNA was synthesized using a reverse transcription system kit according to the manufacturer's instructions (PrimeScript™ RT reagent Kit with gDNA Eraser, RR047A, TaKaRa, Japan). Real-time PCR was performed using the TB Green Premix Ex Taq™ II kit (RR820A, Takara, Japan) following the manufacturer's protocol. Gene expression levels were normalized to GAPDH and analyzed using the comparative cycle threshold ($F = 2^{-\Delta\Delta\text{Ct}}$) method. Primer sequences for qRT-PCR are listed in Supplementary Table 2.

The total proteins from kidney tissue were extracted using ice-cold RIPA lysis buffer containing phosphatase and protease inhibitor cocktail (KeyGen, China). The concentrations of extracted proteins were determined using a BCA protein assay kit (Thermo Scientific, USA). An equal amount of protein from each sample was run in 8% SDS-PAGE gel, then transferred to polyvinylidene difluoride (PVDF) membranes (Millipore, USA). PVDF membranes were blocked with 5% skim milk at room temperature for 2 h and then incubated with primary antibodies at 4°C overnight, followed by incubation with secondary horseradish peroxidase-conjugated antibodies (1:1000, Sungene Biotech, China) for 1 h at room temperature. The intensity of bands was visualized and determined using a ChemiDoc™ XRS detection system (Bio-Rad, USA). Primary antibodies used were: NF- κB p65 (L8F6), phospho-NF- κB p65 (Ser536, E1Z1T), I κB - α (L35A5), phospho-I κB - α (Ser32/36, 5A5) (1:1000, Cell Signaling Technology, USA) and β -actin (1:1000, Sungene Biotech, China). The primary images (Fig. S31) were cropped for presentation.

Reporting summary. Further information on research design is available in the Nature Research Reporting Summary linked to this Article.

Data availability

The main data supporting the findings of this study are available within the Article and its Supplementary Information. Extra data are available from the corresponding authors upon reasonable request. The source data underlying Figs. 2b–e, 3a–l, 4c, d, f, 5c–j, 6a–c,

7b–e, h–j, 8a, b, d, g, i, 9e, f, h, j–m as well as Supplementary Figs. 1f–h, 2, 3a, 4a, 5–12, 15, 17, 19, 22, 24, 26a–c, 28a, b, 32a, b are provided as a Source Data file.

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Author contributions

T.L., Z.M., G.L., X.C., and J.D. conceived and designed the research. T.L. conducted the experiments with assistance from B.X. F.X., J.T., and Z.C.; X.Z. assisted in the Western blot and qRT-PCR experiments. C.W. performed the BioTEM and contributed to the discussion. T.L., Z.M., G.L., X.C., and J.D. co-wrote the paper.

Competing interests

The authors declare no competing interests.

Additional information

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