- 1 Ultrasmall radical metal organic cage as cascade antioxidant nanozyme for renal
- 2 injury
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20 Abstract

Rationale: As substitutes for natural enzymes, nanozymes offer tunable enzyme-like activities and remarkable structural stability, making them potential to treat various diseases, including renal ischemia-reperfusion (I/R) injury. However, the majority of developed nanozymes suffer from unclear structures and limited activity profiles, which hinder the study of their structure-activity relationships, catalytic diversity, mass production, and clinical application.

27 **Methods:** Herein, we introduce an atomically precise and ultrasmall cascade 28 nanozyme based on a radical-functionalized metal-organic cage (MOC-R). This

nanozyme is synthesized through the coordination of radical ligands with copper ions,
resulting in a cuboctahedral structure.

Results: The MOC-R exhibits cascade antioxidant activities, mimicking the functions of superoxide dismutase (SOD) and catalase (CAT), owing to the synergism between the external radicals and internal copper clusters. The MOC-R nanozyme demonstrates exceptional radical scavenging and anti-inflammatory properties. It mitigates immune cell infiltration, promotes macrophage polarization towards the M2like phenotype, reduces inflammatory cytokine secretion, and suppresses excessive autophagy and apoptosis.

Conclusions: This study not only presents an atomically precise cascade nanozyme
 but also highlights its promising therapeutic potential for renal I/R injury.

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41 Keywords: metal organic cage, cascade nanozyme, antioxidant, ischemia reperfusion
42 injury, kidney

- 43
- 44 **Graphical Abstract**



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47 Introduction

48 Clinically, renal I/R injury is a significant concern in various medical and surgical 49 settings. This condition commonly occurs during procedures such as partial 50 nephrectomy, kidney transplantation, and aortic surgery [1]. In these scenarios, the 51 temporary cessation of blood flow to the kidneys followed by its re-establishment can 52 lead to severe damage, causing acute tubular necrosis. The underlying mechanisms 53 responsible for renal I/R injury are complex and multifaceted, involving several 54 interconnected pathways. One of the key factors is the excessive production of 55 intracellular reactive oxygen species (ROS). Under normal conditions, ROS are produced in small amounts and play essential roles in cellular signaling and 56 57 homeostasis. However, during ischemia, the lack of oxygen leads to mitochondrial 58 dysfunction, and upon reperfusion, there is a sudden influx of oxygen that triggers an 59 overproduction of ROS. Given the central role of ROS in this process, the 60 administration of antioxidants has been explored as a potential therapeutic strategy. 61 Various studies have demonstrated promising results in experimental models, where 62 antioxidants effectively mitigated tissue damage and improved renal function. However, 63 translating these findings into clinical practice has proven challenging. Drugs designed to scavenge ROS generated during reperfusion often face limitations in efficacy due to 64 65 their unsustainable one-time exhausted character and poor accumulation at the injured site [2,3]. Enzymes, as remarkable biological catalysts, play a crucial role in facilitating 66 67 chemical reactions within living organisms. One of the key characteristics of enzymes 68 is that they remain structurally and chemically unchanged before and after the 69 reactions they catalyze. This unique property allows enzymes to be reused multiple 70 times without losing their functionality [4–8].

71 However, natural enzymes possess a level of complexity and instability that 72 complicates their synthesis, large-scale production, and long-term preservation. The 73 intricate three-dimensional structures of natural enzymes are highly sensitive to 74 environmental factors such as temperature, pH levels, and organic solvents. Even 75 minor changes in these conditions can lead to denaturation, where the enzyme loses 76 its functional shape and, consequently, its catalytic ability. This sensitivity poses 77 significant challenges for industrial and clinical applications, particularly when it comes 78 to maintaining enzyme activity over extended periods or under varying conditions. Moreover, the difficulty in producing natural enzymes on a large scale and preserving 79 80 them effectively has hindered their widespread use in clinical settings. In medical

81 treatments, enzymes are often required to remain stable and active for prolonged 82 durations, which is not always feasible with natural enzymes due to their inherent 83 instability. This limitation has spurred the development of artificial enzymes designed 84 to mimic the functionality of natural enzymes while offering enhanced structural stability [9,10]. This has driven the need for artificial enzymes that offer the functionality of 85 natural enzymes and structural stability. Nanozymes, emerging from advancements in 86 87 nanoscience, represent a promising class of artificial enzymes [11,12]. They offer 88 benefits such as structural stability, multifunctionality, low cost, recyclability, and 89 feasibility for large-scale production [13,14]. After nearly a decade of evolution, various 90 nanozymes with diverse structures have been developed to alleviate various diseases 91 and manifested outstanding physiological activities [15–18]. Despite their potential, 92 several challenges persist: 1) The often ambiguous structures of nanozymes 93 complicate the study of their structure-activity relationships and their controlled large-94 scale preparation, which hinders clinical application; 2) The lack of standardized 95 protocols for evaluating catalytic activity makes it difficult to compare different 96 nanozymes; 3) Many nanozymes exhibit limited and monotonic activity, which restricts 97 their therapeutic efficacy; 4) The biological safety of nanozymes remains uncertain, 98 posing risks to clinical use [4,7,19-21]. Addressing these issues by developing 99 nanozymes with well-defined structures, versatile enzyme-like activities, and superior 100 biocompatibility is critical for advancing their practical applications.

101 To address the challenges associated with nanozymes, we employed a well-defined 102 nanoobject integrating various active components. The metal-organic cage (MOC) 103 offers an atomically precise structure, presenting a uniform, discrete, and ultrasmall 104 nanopolyhedron formed through the coordination of metal ions or clusters with organic 105 ligands [22-26]. MOCs integrate the advantages of both organic and inorganic 106 components, offering several key benefits: 1) By modifying the organic linkers and 107 metal nodes, researchers can tailor MOCs to possess varying nanoscale cavities and 108 dimensions. This tunability allows for precise control over pore size and surface area, 109 enhancing their functionality in applications such as catalysis and drug delivery. 2)

110 Direct coordination of metal ions can shorten the spatial distance between catalytic 111 sites, facilitating improved cascade catalysis. This arrangement promotes efficient 112 multi-step reactions by ensuring that intermediates are readily accessible to subsequent active sites. 3) MOCs exhibit a unique cavity structure that enables 113 114 reaction molecules to easily enter the cavities and interact with catalytically active sites 115 from both inside and outside. This accessibility enhances catalytic efficiency and 116 selectivity [27–29]. Due to these structural features, functionality, and catalytic activity 117 of metal sites, MOCs have found extensive use in the biomedical field [30,31]. Actually, 118 we have successfully constructed MOC-based cascade nanozymes recently, such as 119 heterometallic MOC and oxidized MOC, demonstrating excellent enzymatic activity 120 [32–34]. However, despite these advantages, the atomic-level structure of these MOCs 121 remains unclear, complicating detailed structural analysis and understanding of their 122 active sites. Additionally, while these MOCs exhibit adequate biocompatibility, 123 concerns remain regarding potential metallic toxicity [35-37].

124 In this study, we synthesized a novel precise and ultrasmall radical MOC nanozyme 125 with cuboctahedral topology and good water solubility, MOC-R, through robust 126 coordination between copper ions and functional isophthalic acid, incorporating 127 oligoethylene glycol and 2,2,6,6-tetramethylpiperidin-1-oxyl (TEMPO) [38,39]. The 128 mechanistic investigations demonstrated that TEMPO and the copper component drive 129 the mimicry of superoxide dismutase (SOD) [40,41] and catalase (CAT) [42,43] 130 activities, respectively, resulting in exceptional antioxidant activity due to their 131 synergistic effects. In vitro and in vivo experiments revealed that this synergy provides 132 significant radical scavenging and anti-inflammatory benefits by reducing immune cell 133 infiltration, promoting macrophage polarization to the M2-like phenotype, limiting inflammatory cytokine release, and inhibiting excessive autophagy and apoptosis 134 (Scheme 1). In summary, this atomically precise and ultrasmall cascade antioxidant 135 136 nanozyme demonstrates exceptional stability, biocompatibility, ROS scavenging 137 capacity, and anti-inflammatory performance. These properties are expected to 138 significantly advance structure-activity studies and facilitate clinical translation.



Scheme 1. Schematic illustration of preparation of precise and ultrasmall MOC-R
cascade nanozyme and its application in treating renal I/R injury. Created with
BioRender.com.

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144 **Results and Discussion**

145 **Preparation and Characterization of MOC-R Nanozyme**

146 The detailed synthetic procedures and structural characterizations are provided in the 147 Figure 1A and supporting information (Scheme S1, and Figures S1-S12). To 148 investigate the source of enzyme-like activity, water-soluble MOC lacking radical moieties is synthesized from ligand L. Analysis of the ¹H-NMR spectra for L-R and 149 150 MOC-R (Figure S13) revealed that all peaks for MOC-R were broadened and shifted 151 to lower fields compared to those of L-R, which indicate the coordination between L-R 152 and Cu²⁺, similar to previously reported results [24]. Specifically, two characteristic 153 peaks at 8.41 and 6.37 ppm, attributed to the benzene ring in L-R, shifted to 8.65 and 6.73 ppm in MOC-R. Fourier transform infrared (FT-IR) spectra showed that both 154 ligand L and L-R exhibited a carboxylic acid signal at 1700 cm⁻¹. This signal 155 disappeared in MOC and MOC-R due to coordination effects (Figure 1B). Additionally, 156 characteristic signals for TEMPO at 2940 cm⁻¹ and 1387 cm⁻¹ appeared in L-R and 157

MOC-R but were absent in L and MOC. These NMR and FT-IR results confirm the successful coordination between the ligands and Cu^{2+} to form MOC and MOC-R.

160 Typically, coordination between isophthalic acid and copper ions results in the formation of a cuboctahedral MOC with a stoichiometric Cu₂₄L₂₄ composition. Due to 161 their large molecular weights, size exclusion chromatography (SEC) was employed to 162 163 analyze the molecular weight, stability, and polydispersity of the ligands and MOCs. 164 SEC plots revealed single sharp peaks at approximately 12.90 min for L and 12.42 min for L-R, and peaks at 11.62 min for MOC and 11.15 min for MOC-R (Figure 1C). The 165 shorter elution times for MOC and MOC-R likely reflect their larger hydrodynamic 166 167 volumes compared to the ligands. The molecular weights of MOC and MOC-R, as 168 determined from the SEC plots, were 15763 g/mol and 20867 g/mol, respectively, 169 closely matching the theoretical values of 16084 g/mol and 22189 g/mol. Furthermore, 170 MOC and MOC-R exhibited narrow polydispersity indices of 1.13 and 1.06, 171 respectively, indicating high uniformity, homogeneity, and stability. These molecular 172 weights were also confirmed by matrix-assisted laser desorption ionization time-of-173 flight (MALDI-TOF) mass spectrometry (Figure 1D), where peaks at 16084 and 22189 174 g/mol were observed, consistent with the theoretical values, confirming the formation 175 of cuboctahedral MOC and MOC-R.

176 Benefiting from the surface OEG moieties, the water solubility of MOC-R is about 2.5 177 mg/mL. The size and surface charge of MOC-R were determined using dynamic light 178 scattering (DLS) and zeta potential measurements for subsequent biological 179 applications. MOC-R exhibited a hydrated particle size of approximately 7.0 nm with a 180 narrow polydispersity of 0.18 (Figure 1E) and a slightly negative charge of -3.5 mV, 181 suggesting good biosafety. The DLS results of MOC-R after being dissolved in PBS or 182 fetal bovine serum (FBS) for one day are consist with above value, indicating its 183 structural stability (Figure S14). Atomic force microscopy (AFM) was used to visualize 184 the morphology and size of MOC-R (Figure 1F). The height of MOC-R was 185 approximately 4.5 nm, showing high uniformity, which aligns with the theoretical size of the cuboctahedral MOC and confirms its structural stability, uniformity, and 186

187 ultrasmall size. Electron paramagnetic resonance (EPR) spectroscopy was employed to examine the radical characteristics of L-R and MOC-R (Figure 1G). The EPR 188 189 spectrum of L-R, showing three consistent peaks, was identical to that of unmodified 190 TEMPO, indicating that the radical properties were retained. MOC-R also displayed 191 intense radical signals, confirming the presence of radical characteristics after the 192 formation of MOC-R. The UV-vis spectra of MOC and MOC-R in PBS (0.2 mg/mL) 193 show negligible absorption in 500~650 nm, suitable for the further enzyme activity 194 evaluation (Figure S15).



Figure 1. Preparation and Characterization of MOC-R cascade nanozyme. (A) The
structures of ligands, MOC, and MOC-R. (B) The FT-IR spectra of ligands, MOC, and
MOC-R. (C) The SEC profiles of the ligands, MOC, and MOC-R in THF. (D) The
MALDI-TOF mass spectra of MOC and MOC-R. (E) The DLS plot of MOC-R in PBS.
(F) The AFM image of MOC-R. Scale bars represent 500 nm. (G) The EPR curves of
L-R and MOC-R.

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203 SOD-like and CAT-like Activities of MOC-R Nanozyme

204 After fully characterizing the MOC-R nanozyme, its atomically precise and ultrasmall 205 structure and radical features prompted us to explore its enzyme-like activities. Given 206 that scavenging superoxide anions (O_2) is crucial in anti-ROS cascade reactions and 207 considering the established activity of the TEMPO, we initially investigated its SOD-208 like activity by monitoring O_2^- elimination. Results from nitrotetrazolium blue chloride 209 (NBT) [44], a redox indicator for detecting O₂, demonstrated that MOC-R and its 210 components exhibited notable SOD-like activity compared to the PBS control group (Figure 2A). Specifically, Cu(OAc)₂·H₂O, L-R, and MOC all displayed evident SOD-like 211 212 activity, with the radical L-R showing superior performance. Interestingly, the SOD-like 213 activity of the assembled MOC was significantly higher than that of $Cu(OAc)_2 \cdot H_2O$, 214 likely due to the active species in MOC being the Cu₂(COO)₄ paddlewheel cluster 215 rather than free copper ions [45,46]. Notably, the integrated MOC-R exhibited the most 216 pronounced SOD-like characteristics, indicating that the formation of MOC-R 217 enhanced its SOD-like activity, primarily due to the TEMPO moieties and assisted by 218 the internal copper clusters. Further investigation revealed that the SOD-like 219 performance of MOC-R was concentration-dependent, with increased activity 220 observed as the concentration of MOC-R increased (Figure 2B). Additionally, EPR 221 spectrometry confirmed the O_2^- scavenging capability of MOC-R nanozymes, as 222 evidenced by a significant reduction in EPR signals upon the addition of MOC-R 223 despite the signal overlap of TEMPO within MOC-R and O₂ (Figure 2C). The SOD-224 like specific activity of MOC-R is calculated to be 3.16 U/mg (Figure S16) and EC₅₀ is 225 about 108 µg/mL.

Following the disproportionation of $\cdot O_2^-$, hydrogen peroxide (H₂O₂) was further catalyzed by catalase (CAT) into water and oxygen. We assessed H₂O₂ consumption and oxygen generation using a dissolved oxygen detector to validate the CATminicking activities of MOC-R nanozyme (Figure 2D) [47]. Cu(OAc)₂·H₂O, MOC, and MOC-R all exhibited significant CAT-like activities, with MOC showing better activity and MOC-R demonstrating the highest. The superior CAT-like activity of the

232 assembled MOC compared to dissolved Cu(OAc)₂·H₂O may also stem from 233 differences in active species. The MOC-R, integrating both active radicals and copper clusters [48], enhanced CAT-like activity, primarily attributed to the active copper 234 clusters and assisted by the external TEMPO moieties. The elimination rates for MOC-235 R were concentration-dependent, with higher quantities of MOC-R accelerating H_2O_2 236 237 catalysis (Figure 2E). Moreover, CAT-like kinetics, determined by oxygen generation, 238 confirmed that MOC-R nanozyme exhibited excellent CAT-like activity in converting 239 H₂O₂ into water and oxygen in a concentration-dependent manner (Figure S17). To 240 further determine the CAT-like activity of MOC-R, a kinetic analysis was carried out. 241 The Michaelis–Menten constant (Km) value of MOC-R with O₂ as substrate was 242 calculated by the Michaelis-Menten equation to be 0.516 mM, and the maximum 243 velocity (Vm) value was 7.743 mg/L/min (Figure 2F). Collectively, MOC-R 244 demonstrated excellent SOD- and CAT-like activities in vitro benefiting from the 245 synergism between external TEMPO moieties and internal copper clusters. Further, 246 we also estimated the radical scavenging ability of MOC-R (Figure S18-19). The 247 inhibition of hydroxyl radicals and ABTS radicals of MOC-R were larger than 80% at concentrations of 0.40 and 0.66 mg/mL, respectively. 248



Figure 2. The antioxidant enzyme-like activities of MOC-R nanozyme. (A) The SODlike activities of different groups. (B) Correlation between the amount of MOC-R and its SOD-like activity. (C) EPR detection of $\cdot O_2^-$ signals under varying conditions. (D) Kinetics of O_2 generation for CAT-like activity of MOC-R. (E) Impact of MOC-R amount on its CAT-like activity. (F) Kinetic analysis and Michaelis-Menten fitting of the CATlike activity of MOC-R.

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The *in vitro* Cellular Protective and ROS-Scavenging Efficacy of MOC-R Nanozyme

259 To explore the antioxidant potential of MOC-R in biomedical applications, we focused 260 on its biocompatibility firstly. Initial assessments included methyl thiazolyl tetrazolium 261 (MTT) [49] assays and hemolysis tests at the cellular level. Human kidney-2 (HK-2) 262 cells demonstrated a survival rate exceeding 90% across various MOC-R 263 concentrations (0-160 µg/mL), indicating excellent biocompatibility (Figure 3A). 264 Hemolysis rates remained exceptionally low, even at high MOC-R concentrations (160 265 µg/mL), further confirming its minimal toxicity to erythrocytes (Figure 3B). Cellular 266 uptake of MOC-R was assessed using inductively coupled plasma mass spectrometry 267 (ICP-MS), revealing increased MOC-R content in HK-2 cells with prolonged incubation 268 (Figure 3C). The ultrasmall size of MOC-R and its non-robust Cu₂(COO⁻)₄ cluster 269 facilitated gradual release from the cells in either integrated or decomposed forms.

270 Subsequently, we evaluated the cellular protective capacity of MOC-R nanozyme 271 against stressors induced by H₂O₂ or hypoxia/reoxygenation (H/R) exposure (Figure 272 3D-E). MTT assay results showed a significant reduction in HK-2 cell viability with H₂O₂ 273 treatment or H/R exposure compared to controls. MOC-R treatment, however, notably 274 mitigated these adverse effects and preserved cell viability, highlighting its potential as 275 a protective agent against oxidative stress. To elucidate MOC-R's mechanism, we 276 assessed it's in vitro antioxidant efficacy using the DCFH-DA [50] probe through flow 277 cytometry (Figure 3F). H/R treatment elevated ROS levels in HK-2 cells, with L-R, MOC, 278 and MOC-R all demonstrating ROS scavenging abilities. MOC-R proved significantly

279 more effective than the other groups, underscoring its potent antioxidant performance. We subsequently quantified the mRNA expression levels of inflammation-related 280 cytokines in HK-2 cells following H/R injury. Our results demonstrated that the mRNA 281 expression of pro-inflammatory cytokines (TNF- α , IL-2, and IL-6) significantly 282 increased after H/R injury, while the mRNA expression of the anti-inflammatory 283 284 cytokine IL-10 significantly decreased. As expected, treatment with the MOC-R 285 nanozyme markedly reversed these changes (Figure 3G). Moreover, Annexin V/PI staining revealed that MOC-R significantly reduced H/R-induced apoptosis in HK-2 286 cells, indicating that MOC-R not only attenuates inflammation but also enhances cell 287 288 survival under conditions of oxidative stress (Figure 3H).



Figure 3. The *in vitro* cellular protective and ROS-scavenging efficacy of MOC-R nanozyme. (A) The impact of MOC-R on HK-2 cell viability assessed using the MTT assay. (B) Assessment of hemolytic activity of double-distilled water (DDW) or MOC-

R on HK-2 cells. (C) Cellular uptake of MOC-R in HK-2 cells. (D) MTT assay of MOC-R on H₂O₂ treated HK-2 cells. (E) MTT assay of MOC-R on H/R assaulted HK-2 cells. (F) Flow cytometry and fluorescence intensity analysis to evaluate the influence of various treatments on HK-2 cells post H/R. (G) The mRNA levels of TNF-α, IL-2, IL-6, and IL-10 in HK-2 cells following H/R. (H) Flow cytometry analysis using Annexin V/PI staining to determine apoptosis of HK-2 cells after H/R. Data are presented as mean ± s.d. ns p > 0.05, * p < 0.05, ** p < 0.01.

300

301 MOC-R Nanozyme Protected Against Renal I/R Injury

302 Firstly, the *in vivo* biocompatibility of MOC-R nanozyme was evaluated by intravenous 303 injection of twice the usual dose. Two days later, the major organs were harvested and 304 underwent histological analysis via H&E staining (Figure S20) [51]. The outcomes 305 demonstrated that the histological injury of all groups of mice were difficult to detect, 306 thereby indicating that MOC-R nanozyme did not inflict any conspicuous damage on 307 the mice, ascertaining the favorable in vivo biocompatibility of MOC-R. Next, ICP-MS 308 was employed to analyze the biodistribution at diverse time points subsequent to the 309 injection of MOC-R solution, serving as an approach to explore the protective 310 mechanism of MOC-R nanozyme in renal I/R injury (Figure 4A). Evidently, the 311 concentration of MOC-R nanozyme in the bloodstream decreased over time, and the 312 spleen and I/R kidney displayed the highest levels of MOC-R accumulation among the 313 organs analyzed. The accumulation of MOC-R in the I/R-injured kidney can be 314 attributed to the increased vascular permeability resulting from I/R injury, which 315 disrupts endothelial cell junctions and triggers the release of pro-inflammatory 316 mediators, thereby compromising the integrity of the vascular barrier. Moreover, owing to the ultrasmall particle size of MOC-R, a considerable portion had been metabolized 317 318 within 24 h across all the examined tissues.

Oxidative stress arises from an imbalance between the production of ROS and the
body's ability to detoxify these reactive intermediates or repair the resulting damage.
This imbalance can lead to cellular dysfunction, tissue damage, and ultimately, organ

322 failure. We therefore evaluated the protective role of MOC-R nanozyme in renal I/R 323 injury. All mice in the study were subjected to 30 min of ischemia. Renal tissues of 324 mice in the I/R group all manifested substantial tubular necrosis and impairment after 325 I/R injury. In contrast, mice treated with the L-R and MOC groups exhibited lessened tubular necrosis and damage after I/R injury; however, the effect was rather low. This 326 327 was likely because molecular L-R was so prone to be cleared out and difficult to exert 328 its effect. Nevertheless, mice treated with MOC-R exhibited significant alleviation of 329 tubular and renal damage following I/R injury, as evidenced by minimal loss of brush 330 border, limited intratubular debris, and reduced glomerular congestion. In contrast, the 331 other groups displayed more pronounced tubular necrosis (Figure 4B-C). The renal 332 protective effects of MOC-R were further examined by assessing creatinine and blood 333 urea nitrogen (BUN) levels following I/R injury (Figure 4D). The outcomes indicated 334 that both creatinine and BUN levels progressively increased and peaked at 24 h after 335 I/R injury, then began to decline after 24 h. In contrast to the I/R group, L-R or MOC 336 treatments slightly reduced the levels of creatinine and BUN, while the levels dropped 337 significantly after MOC-R treatment. Given that MOC-R nanozyme exhibited its anti-338 oxidative capability in vitro, we next verified whether the amelioration of oxidative 339 stress constitutes the underlying mechanism through which MOC-R nanozyme confers 340 renal protection against I/R injury. By utilizing dihydroethidium (DHE), an intracellular 341 superoxide indicator capable of being oxidized by superoxide to yield a red fluorescent product, we discovered that the fluorescence intensity of DHE-stained kidney tissue 342 was augmented subsequent to I/R insult, indicating a high degree of ROS generation 343 344 (Figure 4E-F). This increase was mitigated by L-R or MOC treatment, and was further 345 diminished by MOC-R nanozyme treatment, demonstrating that MOC-R nanozyme 346 inhibited I/R-induced oxidative stress in the kidneys [52,53].

Kidney injury molecule-1 (KIM-1), a biomarker of kidney injury, is expressed at an
extremely low level in healthy kidneys and promptly upregulated following damaging
events [54]. Neutrophil gelatinase-associated lipocalin (NGAL), a small protein
belonging to the lipocalin family, is physiologically produced at a minimal level but rises

351 under harmful circumstances. We then tested KIM-1 and NGAL using immunofluorescence to evaluate the biomarkers expression in injured kidneys. 352 353 Subsequently, we employed immunofluorescence to visualize KIM-1 and NGAL for 354 evaluating the molecular changes occurring in injured kidneys (Figure 4G-I). The 355 results revealed that the expression of NGAL and KIM-1 was significantly up-regulated 356 in I/R mice. However, no significant effect was observed after treatment with L-R and 357 MOC, and a marked decrease in the expression of NGAL and KIM-1 was detected 358 after treatment with MOC-R, indicating that MOC-R significantly changed the expression patterns of I/R induced renal damage. 359





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Figure 4. The protective effect of MOC-R nanozyme against I/R injury. (A) The
biodistribution of MOC-R in diverse tissues as indicated by the copper concentration.
(B) Representative Hematoxylin and eosin staining of the kidney at 24 h post
reperfusion (10x and 20x). Vascular congestions are indicated by arrows. Scale bars

366 represent 100 µm. (C) Assessment of kidney injury based on tubular injury scores. (D) 367 Serum creatinine and BUN levels in renal I/R injury. (E) Representative 368 immunofluorescence images of DHE expression at 24 h post reperfusion. Scale bars 369 represent 100 µm. (F) The count of the DHE positive cells. (G) Representative 370 immunofluorescence images showing KIM-1 expression at 24 h post reperfusion. (H) 371 Representative immunofluorescence images showing NGAL expression at 24 h post 372 reperfusion. Scale bars represent 100 µm. (I) The count of the KIM-1 and NGAL positive cells. Data are presented as mean \pm s.d. * p < 0.05, ** p < 0.01. 373

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375 MOC-R Nanozyme Restricted I/R Induced Inflammatory Milieu and Programmed 376 Cell Death

377 ROS are essential for the induction and maintenance of M1 macrophage polarization. 378 Numerous studies have highlighted the role of ROS in activating both the NF-kB and 379 p38/MAPK signaling pathways, which in turn promote pro-inflammatory gene 380 expression in macrophages [55,56]. Immunofluorescence assays demonstrated that 381 I/R insult augmented the expression of the M1 macrophage phenotype (CD86), yet L-382 R or MOC treatment marginally reversed the M1 phenotype to the M2 macrophage 383 marker (CD206) [57]. Intriguingly, the expression of CD86 substantially declined 384 concomitant with a notable increase in the expression of CD206 following MOC-R 385 nanozyme treatment, which was precisely the converse of that in I/R kidneys (Figure 386 5A-B). Given the significant role of M2 macrophages in kidney fibrosis, we utilized 387 Masson's trichrome staining to assess collagen deposition in renal tissues and 388 evaluate the long-term effects of MOC-R on renal fibrosis (Figure S21). Our results 389 indicated that MOC-R did not increase collagen deposition, suggesting it does not 390 promote fibrotic changes. Additionally, neutrophil infiltration after reperfusion was 391 quantified by Immunofluorescence assays (with positive cells indicated by red 392 fluorescence normalized to DAPI-stained nuclei). A considerable increase in 393 neutrophils was noted in I/R-injured kidney compared to the sham group. While neither 394 L-R nor MOC treatment proved effective in lowering neutrophil levels, a propensity that

395 was effectively alleviated by treatment with MOC-R nanozyme, which significantly 396 decreased neutrophil levels (Figure 5C-D). We then measured the mRNA expression 397 of inflammatory-related cytokines in kidney tissues following I/R injury (Figure 5E), and 398 discovered that the mRNA expression of proinflammatory cytokines (TNF-a, IL-2, and 399 IL-6) significantly increased after I/R injury, while the MOC-R nanozyme markedly 400 decreased them. In contrast, upon I/R injury, the mRNA expression of the anti-401 inflammatory cytokine (IL-10) declined, whereas the MOC-R nanozyme elevated it to 402 a considerable extent, albert missing the significance. These findings demonstrated 403 that the MOC-R nanozyme can not only inhibit the polarization of macrophages to the 404 M1 phenotype and facilitate its polarization to the M2 phenotype but also enhance the 405 suppression of the I/R-induced inflammatory response.

Autophagy is a regulated cellular process of self-degradation and is typically regarded 406 407 as an inducible adaptive response to cellular stress. However, dysregulation or 408 excessive activation of autophagy, such as that occurring during I/R injury, triggers 409 autophagic cell death without caspase involvement. Herein, western blotting was 410 performed to detect the expression levels of autophagy-related proteins (Beclin-1 and 411 P62) in order to evaluate autophagic flux in kidneys following I/R injury (Figure 5F). As 412 expected, MOC-R treatment significantly reduced the expression levels of Beclin-1 and 413 P62 (Figure 5G). In line with the western blotting data, immunostaining targeting LC3 414 manifested a pronounced increase in the number of autophagosomes within the kidney 415 tissue after I/R and a decrease subsequent to MOC-R nanozyme treatment. However, 416 L-R or MOC treatment had no effect on the autophagy level in I/R kidneys (Figure 5H-417 I). Apoptosis is a tightly-regulated ATP-dependent process of programmed cell death 418 that is activated by hypoxia due to ischemia and during ROS generation in reperfusion. 419 The level of cellular apoptosis in I/R kidney was assessed using the TUNEL assay in 420 this study (Figure 5J-K). Our findings indicated that the proportions of TUNEL-positive 421 cells were significantly lower in mice treated with MOC-R than in the control group. 422 These results indicated that the MOC-R nanozyme modulated the anti-inflammatory

423 response by alleviating oxidative stress, thereby reducing programmed cell death in



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Figure 5. MOC-R nanozyme restricted I/R induced inflammatory milieu and 426 427 programmed cell death. (A) Representative immunofluorescence images showing CD86 and CD206 expression at 24 h post reperfusion. Scale bars represent 100 µm. 428 429 (B) The count of the CD86 and CD206 positive cells. (C) Representative 430 immunofluorescence images showing MPO expression at 24 h post reperfusion. Scale 431 bars represent 100 µm. (D) The count of the MPO positive cells. (E) The mRNA levels 432 of TNF- α , IL-2, IL-6, and IL-10 in renal tissue. (F) Expression of Beclin-1 and P62 detected by western blotting assay. (G) The relative densities of the bands in each lane 433 were analyzed and normalized to β -actin. (H) Representative immunofluorescence 434

images showing LC3 expression at 24 h post reperfusion. (I) The count of the cells with LC3 positive autophagosomes. (J) Representative immunofluorescence images showing TUNEL expression at 24 h post reperfusion. Scale bars represent 100 μ m. (K) The count of the TUNEL positive cells. Data are presented as mean ± s.d. * *p* < 0.05.

440 **Conclusion**

441 In this study, an atomically precise, ultrasmall, and water-soluble cascade antioxidant 442 nanozyme was constructed based on a metal organic cage integrating active organic 443 radicals and metal clusters. Through the assembly of ligands containing radicals and 444 copper ions, the water-soluble MOC-R could be obtained in a high yield. The particle 445 size of MOC-R is approximately 7.0 nm, and it is negatively charged; these properties 446 of MOC-R are beneficial for the clinical application of antioxidants in vivo. The MOC-R 447 compound not only demonstrates SOD-like activity and CAT-like activity, but also 448 showcases remarkable potential in the field of antioxidant therapy. Its SOD-like activity 449 is primarily attributed to the presence of TEMPO, a key component that contributes to 450 its free radical scavenging abilities. Meanwhile, the CAT-like activity of MOC-R arises 451 from the unique properties of its copper clusters, which enable it to efficiently neutralize 452 ROS. As a result, this cascading nanozyme is capable of efficiently scavenging harmful 453 ROS, particularly $\cdot O_2^-$ and H_2O_2 , under stressful conditions. MOC-R also displays 454 excellent biocompatibility and significant ROS-scavenging ability at both in vitro and in 455 vivo levels, exerting exceptional renoprotective potency by influencing the anti-456 inflammatory status of the immune response and regulating programmed cell death. In 457 conclusion, the multifaceted capabilities of MOC-R make it a compelling candidate for 458 further exploration in biomedical research aimed at combating oxidative stress-related 459 disorders.

460

461 Materials and Methods

462 **SOD-like activity**

463 SOD-like activity was determined using the nitrogen blue tetrazolium (NBT) method. 464 Under UV irradiation, riboflavin and methionine can react to produce O_2^- , which 465 reduces NBT to generate blue methylhydrazone with a maximum absorption wavelength of 560 nm. However, SOD can scavenge O2 and inhibit the generation of 466 methylhydrazone. Therefore, the higher the SOD-like activity, the lower the amount of 467 468 the reduction product (methylhydrazone), and the lower the absorbance at 560 nm. 469 Thus, the decrease in absorbance at 560 nm confirms the role of SOD-like activity. 470 The PBS solution of samples (2 mg/mL, 90 µL) were added to the mixture of NBT (1 471 mg/mL, 0.5 mL), Met (30 mg/mL, 0.5 mL), EDTA-Na₂ (0.4 mg/mL, 0.3 mL), riboflavin 472 (1 mg/mL, 0.5 mL) and PBS solution (pH = 7.4, 2 mL) and incubated together under 473 constant UV irradiation for 60 s to test the SOD-like activity. The experiments were 474 divided into five groups due to the different samples: (1) PBS; (2) Cu(OAc)₂ H₂O; (3) 475 L-R; (4) MOC; (5) MOC-R. The concentration-dependent SOD-like activities were also 476 determined by varying the concentrations of MOC-R under the same conditions and 477 incubated under UV irradiation for 60 s.

478 To further investigate the SOD-like activity of MOC-R, we employed EPR spectroscopy 479 for a more intuitive assessment. 5,5-Dimethyl-1-pyrroline N-oxide (DMPO) was used 480 as a potent free radical trapping agent, capable of forming spin adducts upon 481 interaction with the $\cdot O_2^-$ radical. The spin adducts, which can be detected by EPR, 482 provide a direct measurement of the $\cdot O_2^-$ radical concentration in the reaction system. 483 The reaction mixture consisted of the following components: Met (130 mM, 0.25 mL), 484 EDTA-Na₂ (0.1 mM, 0.25 mL), H₂O (0.20 mL), MOC-R (2 mg/mL, 0.05 mL), riboflavin (0.02 mM, 0.25 mL), DMPO (100 mM, 70 µL), and phosphate buffer (pH 7.4, 50 mM, 485 486 1.68 mL). The mixture was thoroughly mixed and then placed into a nuclear magnetic 487 resonance tube for subsequent UV irradiation at two time points: 0 s and 20 s, followed 488 by EPR spectrum analysis.

489

490 •O₂⁻ Inhibition Rate

- 491 Mixture of NBT (1 mg/mL, 0.4 mL), Met (30 mg/mL, 0.4 mL), EDTA-Na₂ (0.4 mg/mL, 492 0.25 mL), riboflavin (1 mg/mL, 0.4 mL) and PBS solution (pH = 7.4, 1.85 mL) to obtain 493 the detection working solution. Subsequently, MOC-R solutions (0.35-2.45 mg/mL, 200 494 µL) or PBS (200 µL) was added to above detection working solution and immediately 495 irradiated it with ultraviolet light. Record the changes in absorbance between the 496 sample and blank, calculate the inhibition rate, and obtain the enzyme activity unit. 497 Inhibition rate (%) = (A_b - A_s)/A_b × 100
- 498 where A_b is the absorbance of the blank and A_s is the absorbance of the sample.
- 499

500 CAT-like activity

501 CAT-like activity was determined using a dissolved oxygen meter. Oxygen produced 502 was measured at room temperature by using a specific oxygen electrode on a split 503 dissolved oxygen meter (SW9403). Sample (2 mg/mL, 100 µL) and H₂O₂ solution (30%, 504 200 µL) was added to water (14.94 mL), and the concentration (in mg/L) of produced 505 oxygen was recorded from 0 to 5 min. To assess the CAT-like activity, five samples 506 were set up: (1) PBS; (2) Cu(OAc)₂·H₂O; (3) L-R; (4) MOC; and (5) MOC-R. In addition, 507 different amounts (50, 100, 150, and 200 µL) of MOC-R solution (2 mg/mL) were added 508 to the final volume of the buffer solution (15 mL), and their corresponding CAT-like 509 activity was determined.

510

511 CAT-like Kinetics

A series of solutions of hydrogen peroxide (0.10-12.68 mM in PBS, 25 mL) was prepared, then MCO-R solution (1 mg/mL in PBS, 0.10 mL) was added. The dissolved oxygen in the above mixture was immediately monitored for 5 min using a dissolved oxygen analyzer. The corresponding reaction rate and hydrogen peroxide concentration were then fitted to the Michaelis-Menten equation to obtain the relevant reaction kinetics parameters.

518

519 Hydroxyl radical scavenging

- The reaction mixture contains FeSO₄ (6 mM, 0.1 mL), hydrogen peroxide (6 mM, 0.1 mL), and sample (0.15-2.4 mg/mL in PBS, 0.1 mL). The mixture was incubated at 37 °C for 10 min, covered with Salicylic acid ethanol solution (6 mM, 0.1 mL), and then further incubated at 37 °C for 30 min. Measure the mixture at 510 nm using an enzymelinked immunosorbent assay reader. Ultrapure water was used as a blank instead of the sample.
- 526 Hydroxyl radical scavenging activity (%) = $(A_b A_s)/A_b \times 100$
- 527 where A_b is the absorbance of the blank and A_s is the absorbance of the sample.
- 528

529 **ABTS radical scavenging activity**

The solutions of ABTS (7.0 mM in DDW), potassium persulfate (9.4 mM in DDW), and MOC-R (0.25-2.0 mg/mL in PBS) were prepared firstly. Mixing the solutions of ABTS and potassium persulfate in equal volume at room temperature in the dark for 12 h and then diluted for 20 times with PBS to obtain the ABTS working solution. Add ABTS working solution (0.2 mL) and samples (0.15-2.4 mg/mL in PBS, 0.1 mL) to each well on a 96 well plate, incubate for 30 min, and measure the absorbance at 734 nm using an enzyme-linked immunosorbent assay reader.

- 537 ABTS radical scavenging activity(%) = $(A_b A_s)/A_b \times 100$
- 538 where A_b is the absorbance of the blank and A_s is the absorbance of the sample. 539

540 **Cytotoxicity assay**

541 To assess the cytotoxic effect of MOC-R, HK-2 (7000 cells/well) cells were added into a 96 well plate, and cultured at 37 °C for 12 h. Different concentrations (0, 5, 10, 20, 542 543 40, 80, 160 µg/mL, 10 µL) of MOC-R solution in PBS were added in cells and cultured 544 for another 24 h. Afterwards, the cells were gently washed by PBS (pH 7.4) for three times. The Thiazolyl Blue Tetrazolium Bromide (MTT) solution (Beyotime, Shanghai) 545 was added to each well individually and incubated for 4 h in the dark. Subsequently, 546 the cell supernatant was collected using a micropipettor, followed by addition of 150 547 548 µL of DMSO. After thorough agitation for 10 min to completely dissolve MTT crystals,

549 cell viability was assessed using a microplate reader at absorbance wavelengths of550 570 nm.

551

552 Hemolysis assay

The whole blood of healthy mice was collected in a 2 mL EP tube, and the blood was centrifuged at 2000 rpm for 5 min to remove the upper serum and obtain erythrocytes. Then erythrocytes were washed three times with PBS, and 0.5 mL erythrocytes were diluted with 10 mL PBS. Then 0.5 mL of MOCs solution (5, 10, 20, 40, 80, 160 µg/mL), or DDW were added to 0.5 mL erythrocytes suspension, respectively. All sample tubes were kept at room temperature for 2 h and then centrifuged at 2000 rpm for 5 min. Finally, the experimental result was recorded by a camera.

560

561 **Cellular uptake**

562 ICP-MS was used to accurately measure the uptake of MOC-R by HK-2 cells. In the *in vitro* cell uptake assay, HK-2 cells were inoculated into 10 mm tissue plates at a density of 10,000 cells/mL and cultured for 24 h. After 24 h, MOC-R (10 µg/mL, final concentration) was added to each tissue plate and cultured for 2, 4, 8, or 12 h.

566 In the *in vitro* cell retention assay, HK-2 cells were inoculated onto 10-mm tissue plates at a density of 10,000 cells/mL, and after 24 h of incubation, MOC-R (10 µg/mL, final 567 568 concentration) was added to each tissue plate and incubated for another 12 h. After 569 that, the medium was extracted and replaced with fresh medium (without MOC-R). 570 Cells were subsequently collected at the "12 + 4" h and "12 + 8" h time points. After trypsinization and centrifugation steps, the cell pellet was washed three times with 571 572 phosphate buffer solution. The samples were digested using a solution containing HNO₃ (68%, 0.25 mL) and HCI (38%, 0.75 mL), and the digestion process lasted for 4 573 h at a temperature of 110 °C. After cooling, the sample was diluted with HCl (2%) to a 574 575 volume of 10 mL. The copper concentration was then determined by ICP-MS.

576

577 Intracellular ROS scavenging ability

578 Intracellular ROS levels were assessed using the fluorescent probe DCFH-DA. HK-2 579 cells (7000 cells per well) were cultured in 6-well microtiter plates at 37 °C for 12 h. To 580 induce hypoxia, Na₂S₂O₄ solution (2 mM) was added to the culture medium and the 581 cells were placed in a CO₂ incubator. The five experimental groups were designed as 582 follows: (1) Control; (2) H/R; (3) H/R + L-R; (4) H/R + MOC; (5) H/R + MOC-R. The 583 experimental groups (3-5) were added with samples (10 µg/mL, final concentration) and cultured at 37 °C for another 6 h. The results were analyzed using flow cytometry 584 (FACSymphony, BD Biosciences, NJ). 585

586

587 Real-time PCR analysis

The total cellular or renal RNA was extracted using TRIzol reagent (Invitrogen, 588 589 Shanghai) according to the manufacturer's instructions. cDNA was transcribed using 590 a Superscript III Reverse Transcriptase Kit (Invitrogen) and oligo d(T) (Applied 591 Biosystems, Waltham, MA, USA). Quantitative RT-PCR analysis was performed with a SYBR RT-PCR kit (Takara, Tokyo) and the StepOne Real-Time PCR System 592 593 (Applied Biosystems). All reactions were conducted in a 20 µL reaction volume in triplicate. The relative expression levels for a target gene were normalized against 594 595 GAPDH. Primers used for RT-PCR analysis are as follows: TNF-a (5'-AAG CCT GTA 596 GCC CAC GTC GTA-3', 5'-GGC ACC ACT AGT TGG TTG TCT TTG-3'), IL-2 (5'-CCA 597 TGA TGC TCA CGT TTA AAT TTT-3', 5'-CAT TTT CCA GGC ACT GGA GAT G-3'), IL-6 (5'-ACA ACC ACG GCCTTC CCT ACT T-3', 5'-CAC GAT TTC CCA GAG AAC 598 599 ATG TG-3'), IL-10 (5'-GCT TTA CTG ACT GGC ATG AG-3', 5'-CGC AGC TCT AGG AGC ATG TG-3'), and GAPDH (5'-TGA CCA CAG TCC ATG CCA TC-3', 5'-GAC GGA 600 601 CAC ATT GGG GGT AG-3').

602

603 Cellular apoptosis assay

An Annexin V-FITC/PI Apoptosis Detection Kit (Beyotime) was utilized to evaluate cell
 apoptosis. Cells were washed twice with cold PBS and subsequently harvested using
 trypsin without EDTA. The cells were resuspended in 195 μL of binding buffer. Next, 5

μL of Annexin V-FITC and 10 μL of propidium iodide staining solution were added,
followed by incubation in an ice bath protected from light for 15 min. The percentage
of apoptotic cells was quantified by flow cytometry (FACSymphony, BD Biosciences,
NJ).

611

612 Animal experimentation

Animal and Ethics Statement: C57BL/6 mice (female, 7-8 weeks old) were obtained from Shanghai Laboratory Animal Center, China. All animal experiments were performed in accordance with the guidelines of the National Institute of Health for the Care and Use of Laboratory Animals and approved by the Scientific Investigation Committee of Shanghai Changzheng Hospital (No. 202403013A).

618

619 **Pharmacokinetics analysis**

The mice were sacrificed at the designated time points. Blood and all organs were collected, weighted and digested in a solution containing 0.25 mL HNO₃ (68%) and 0.75 mL HCl (38%) for 12 h at 110 °C. After cooling, the samples were diluted with HCl (2%) to 10 mL. Cu contents were then detected by ICP-MS.

624

625 Induced renal ischemia/reperfusion model

626 Mice were anesthetized with sodium pentobarbital (50 mg/kg, intraperitoneal injection), 627 and after a right flank incision, the right renal hilum was clamped with a noninvasive 628 microaneurysm clamp (Shanghai Medical Devices Co., Ltd., Shanghai, China) for 30 629 min. The left contralateral kidney was considered as a sham operation. The incision 630 was temporarily closed during ischemia. After removal of the microaneurysm clip, the lateral abdominal incision was closed after visual confirmation of reperfusion. Body 631 temperature was maintained using an adjustable heating pad. All mice were 632 633 intraperitoneally injected with 0.5 mL of isotonic saline after surgery, and L-R, MOC, or 634 MOC-R (250 µg/mL, 400 µL) were injected into different mice through the tail vein and 635 were executed at the indicated reperfusion time points.

636

637 Histopathological assessment

638 Mice were executed at the indicated time points, and mouse kidneys were coronal 639 sectioned, fixed in 10% buffered formalin, paraffin-embedded, and sectioned at a 640 thickness of 3 µm sections were stained with hematoxylin and eosin to assess tissue 641 damage. Kidney sections were blindly labeled and randomized for observation by two 642 researchers. The extent of renal tubular injury was assessed semiguantitatively and 643 pathologically on a scale of 0 to 4: 0 = no recognizable injury; 1 = single cell necrosis; 644 2 = necrosis of all cells in the adjacent convoluted tubule, with surrounding tubules 645 surviving; 3 = necrosis confined to the distal one-third of the proximal convoluted tubule, 646 with a band of necrosis spanning the endothelial layer; 4 = necrosis affecting all three 647 segments of the proximal convoluted tubule.

648

649 Immunofluorescence assay

Kidney frozen sections were processed according to standard protocols. Subsequently, the sections were incubated overnight with anti-DHE, KIM-1, NGAL, CD86/CD206, MPO, LC3 or TUNEL primary antibody provided by Beyotime Biotech. After that, the sections were rinsed three times with PBS and then incubated with secondary antibody for 1 h at room temperature. Cell nuclei were stained using DAPI. Finally, the slides were examined and imaged under a fluorescence microscope (Nikon80i, Tochigi, Japan).

657

658 Western blotting

Tissues were rinsed with PBS twice and lysed in ice-cold RIPA buffer (Roche, Basel, Switzerland) containing phosphatase and protease inhibitors. Sample proteins were then subjected to 12% sodium dodecyl sulfate-polyacrylamide gel electrophoresis and transferred to nitrocellulose membranes. Membranes were probed with Beclin-1 and P62 antibodies from Beyotime Biotech. The relative quantity of proteins was determined by a densitometer software (ImageJ, NY). 665

666 Masson staining

The kidney tissue was fixed in 4% paraformaldehyde. Slices of the paraffin-coated kidney samples were taken. Masson's Trichrome Stain Kit (Beyotime) was used on kidney tissues. Kidney sections were blindly labeled and randomized for observation by two researchers. The Masson staining was utilized to semi-quantitatively analyze the collagen volume fraction, which is defined as the proportion of collagen-positive blue area to the total tissue area. The Image J software was employed for this purpose.

673

674 Statistical analysis

Statistical significance was determined utilizing an ANOVA followed by Bonferroni's test correction using GraphPad Prism 10 (La Jolla, USA). The results are expressed as the mean \pm standard deviation (SD). In every case, p < 0.05 was considered statistically significant.

679

680 Abbreviations

681 AFM: atomic force microscopy; BUN: blood urea nitrogen; CAT: catalase; DAPI: 4',6-682 diamidino-2-phenylindole; DCFH-DA: 2',7'-dichlorodihydrofluorescein diacetate; DLS: 683 dynamic light scattering; DHE: dihydroethidium; DMPO: 5,5-dimethyl-1-pyrroline N-684 oxide; EPR: electron paramagnetic resonance; FBS: fetal bovine serum; FT-IR: flourier 685 transform infrared spectroscopy; H/R: hypoxia/reoxygenation; H&E: hematoxylin and 686 eosin; HK-2: Human kidney-2; I/R: ischemia-reperfusion; ICP-MS: inductively coupled 687 plasma mass spectrometry; KIM-1: kidney injury molecule-1; MALDI-TOF: matrix 688 assisted laser desorption/ionization time of flight; MOC: metal-organic cage; MOC-R: 689 radical-functionalized metal-organic cage; MTT: methyl thiazolyl tetrazolium; NBT: 690 nitrotetrazolium blue chloride; NGAL: neutrophil gelatinase-associated lipocalin; ROS: 691 reactive oxygen species; SEC: size exclusion chromatography; SOD: superoxide 692 dismutase; TEMPO: 2,2,6,6-tetramethylpiperidin-1-oxyl; WB: western blot.

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- 698

699 **Competing Interests**

- 700 The authors have declared that no competing interest exists.
- 701

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