Reduction-responsive RNAi nanoplatform for enhanced cancer sonoimmunotherapy via dual inhibition of mitophagy and Nrf2 pathways

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Abstract

Rationale: Sonodynamic therapy (SDT) has emerged as a promising non-invasive modality with deeper tissue penetration than photodynamic or chemodynamic therapies. However, its therapeutic efficacy remains limited due to inadequate reactive oxygen species (ROS) generation, largely attributed to tumor-intrinsic antioxidant systems and mitophagy. Existing combinations of SDT with immunotherapy are primarily additive and fail to address the mechanistic interplay between ROS suppression and immune evasion.

Methods: To overcome these limitations, we developed a redox-responsive RNA interference (RNAi) nanoplatform (NP) for the co-delivery of Nrf2 siRNA, the mitophagy inhibitor 3-Methyladenine (3-MA), and the sonosensitizer purpurin-18 (P-18). This NP enables tumor-specific release in high-glutathione environments and facilitates dual-pathway inhibition upon ultrasound activation.

Results: This synergistic platform simultaneously disrupted Nrf2-mediated antioxidant defenses and mitophagy-dependent mitochondrial clearance, resulting in enhanced intracellular ROS accumulation. Elevated ROS levels triggered immunogenic cell death (ICD), promoting dendritic cells maturation and antigen presentation. Concurrently, 3-MA inhibited NF-κB signaling, downregulating PD-L1 expression and mitigating T cell exhaustion. In murine breast cancer models, this dual-action approach elicited robust CD8$^+$ T cell responses and significantly suppressed tumor growth and metastasis.

Conclusions: This study introduces a mechanistically integrated sonoimmunotherapeutic strategy that concurrently overcomes ROS suppression and immune checkpoint resistance. By orchestrating redox disruption and immune reprogramming, our nanoplatform provides a compelling framework for next-generation SDT-based immunotherapy.

Keywords: sonoimmunotherapy, mitophagy inhibition, immune checkpoint blockade,

50   reactive oxygen species, redox-responsive nanoplatform
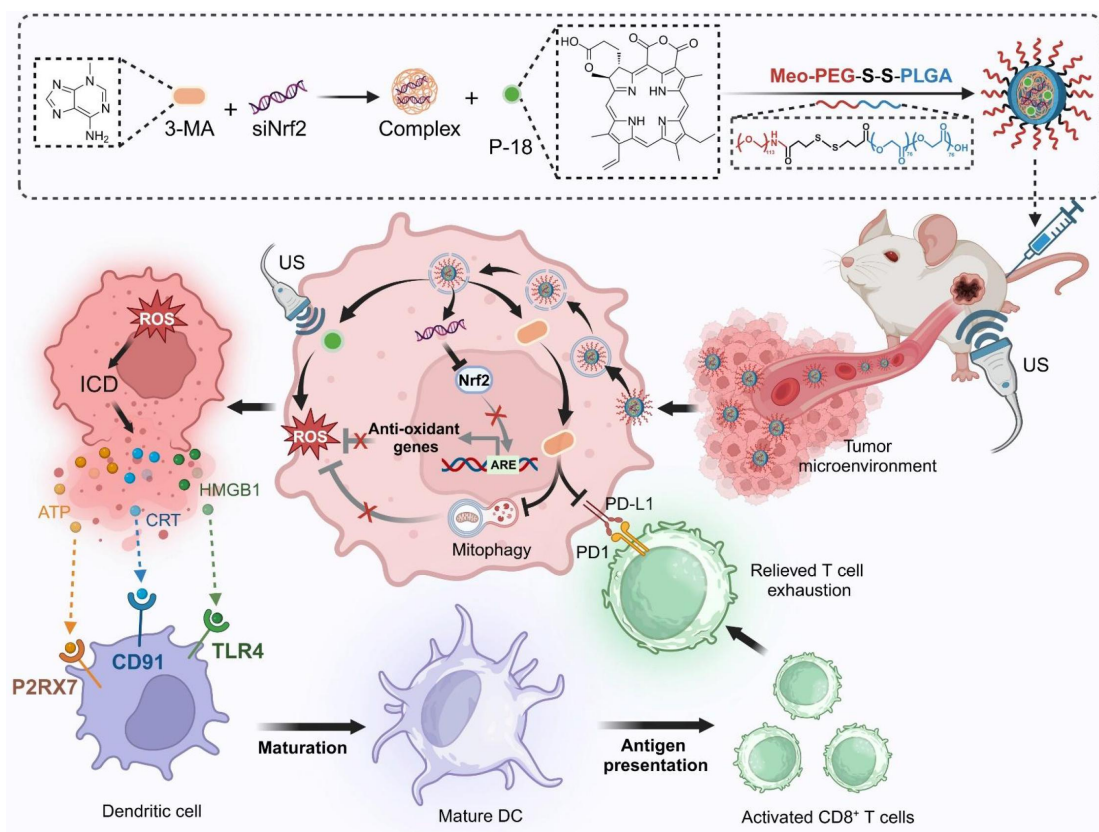

51

52   **Introduction**

53   Sonodynamic therapy (SDT) is emerging as a promising non-invasive cancer

54   treatment that leverages ultrasound (US) to activate sonosensitizers, generating

55   reactive oxygen species (ROS) for tumor ablation [1, 2]. Compared to photodynamic

56   therapy, which is hindered by limited light penetration, and chemodynamic therapy,

57   which relies on endogenous Fenton reactions, SDT offers superior tissue penetration,

58   spatiotemporal precision, and minimal off-target toxicity [3, 4]. Despite these

59   advantages, the clinical translation of SDT remains hampered by several critical

60   limitations, including inefficient ROS production, rapid ROS degradation, limited

61   intracellular delivery, and unwanted ROS consumption [5]. These challenges

62   collectively contribute to suboptimal therapeutic outcomes.

63   A primary obstacle in SDT efficacy lies in the tumor's intrinsic redox defense

64   mechanisms. The Nrf2-Keap1 pathway serves as a central regulator of antioxidant

65   responses, orchestrating the transcription of detoxifying enzymes such as HO-1,

66   NQO1, and GPX4, which mitigate oxidative stress [6]. Concurrently, mitophagy

67   selectively removes damaged mitochondria, a major intracellular source of ROS,

68   further restricting ROS accumulation [7, 8]. Together, these pathways form a

69   synergistic network that enables tumor cells to withstand oxidative insults,

70   significantly undermining ROS-dependent therapies like SDT.

71   To address this limitation, recent research has investigated SDT in combination

72   with immunotherapy, leveraging ROS-induced immunogenic cell death (ICD) to

73   activate antitumor immunity [9]. ROS can trigger the release of damage-associated

74   molecular patterns (DAMPs), such as ATP, calreticulin (CRT), and HMGB1, which

75   promote dendritic cells (DCs) maturation and enhance antigen presentation to

76   cytotoxic T lymphocytes (CTLs) [10, 11]. However, most SDT-immunotherapy

77   strategies demonstrate only additive effects. These approaches typically focus on

78   single-pathway interventions-such as Nrf2 inhibition or immune checkpoint

blockade-without addressing the intricate crosstalk between redox regulation, mitophagy, and immune evasion. Moreover, tumor cells frequently upregulate immune checkpoint proteins such as PD-L1 in response to stress-induced inflammation, establishing a negative feedback loop that suppresses T cell activation and promotes immune escape [12, 13]. Consequently, current combination strategies fall short of delivering durable and synergistic therapeutic benefits.

In this study, we present a redox-responsive RNA interference (RNAi) nanoplatform (NP) engineered to co-deliver Nrf2-targeting siRNA (siNrf2), the mitophagy inhibitor 3-Methyladenine (3-MA), and the sonosensitizer purpurin-18 (P-18). The NP features a disulfide-linked architecture that facilitates glutathione (GSH)-triggered release within the tumor microenvironment (TME), ensuring tumor-selective delivery and minimizing systemic toxicity. Upon US activation, this dual-inhibition strategy simultaneously suppresses Nrf2-driven antioxidant responses and mitophagy, resulting in robust intracellular ROS amplification and enhanced ICD. In parallel, 3-MA downregulates PD-L1 expression via NF-κB inhibition, mitigating T cell exhaustion and enhancing CD8$^+$ T cell infiltration. By synchronizing redox disruption with immune checkpoint modulation, this integrated approach addresses the core limitations of SDT and immunotherapy, offering a compelling framework for next-generation sonoimmunotherapy. Through targeted modulation of interconnected resistance pathways and TME-responsive delivery, our platform advances the therapeutic landscape toward more effective and mechanistically informed cancer immunotherapy (**Scheme 1**).

Scheme 1. Schematic illustration of the design and therapeutic mechanism of the multifunctional nanoplatform NPs(3-MA/siNrf2/P-18) for enhanced SDT and immune activation. The nanoparticle co-encapsulates 3-MA, siNrf2, and P-18, and is administered intravenously for tumor-targeted delivery. In the reductive tumor microenvironment, intracellular glutathione triggers cleavage of disulfide bonds, facilitating controlled release of the payload. Upon US irradiation, P-18 generates ROS, inducing tumor cell apoptosis. Concurrent Nrf2 silencing attenuates antioxidant defenses, while 3-MA inhibits mitophagy, disrupting mitochondrial clearance and further amplifying intracellular ROS levels. This synergistic ROS accumulation promotes ICD, characterized by the release of DAMPs that enhance DCs maturation and antigen presentation. Additionally, 3-MA downregulates PD-L1 expression via NF-κB pathway inhibition, reversing T cell exhaustion and promoting robust CD8$^+$ T cell-mediated cytotoxicity. Together, this strategy enhances SDT efficacy and elicits a potent anti-tumor adaptive immune response, offering a comprehensive approach for improved cancer therapy.

Materials and methods

Materials

3-Methyladenine (3-MA), BAY11-7082, Microcystin-LR (MC-LR) and Purpurin 18 (P-18) were purchased from MedChemExpress (MCE) and used without further modification. Dimethyl sulfoxide (DMSO) and N, N'-dimethylformamide (DMF) were acquired from Sigma-Aldrich and used as received. The cationic lipid-like compound alkyl-modified polyamidoamine (PAMAM) dendrimer (G0-C14) and Meo-PEG5k-S-S-PLGA11k copolymer were synthesized according to previously reported procedures [14]. The DAB (SA-HRP) TUNEL Cell Apoptosis Detection Kit was purchased from Servicebio®. ATP Determination Kit (#A22066), ROS probe (CM-H2DCFDA, #C6827), FIX & PERM™ Cell Permeabilization Kit (#GAS003), MitoTracker® Red CMXRos (#M7512), and LysoTrackerTM Green DND-26 (#L7526) were obtained from Thermo Fisher Scientific. Human CRT (Calreticulin) ELISA Kit (#E-EL-H0627) and Annexin V-FITC/PI Apoptosis Kit (#E-CK-A211) were purchased from Elabscience Biotechnology (Wuhan, China). siRNA targeting human and mouse Nrf2 was acquired from IGE (Guangzhou, China). The siRNA sequences were as follows: siNrf2 (human): 5'-GGC CAG CTG TGA GTG TTT CTT-3' (sense); 5'-AAG AAA CAC TCA CAG CTG GCC-3' (antisense); siNrf2 (mouse): 5'-CAA GGA GCA AUU CAA UGA A-3' (sense); 5'- UUC AUU GAA UUG CUC CUU G-3' (antisense). Cy5-labeled siNrf2 was also obtained from IGE, with the fluorescent dye Cy5 conjugated to the 5'-end of both sense and antisense strands. Dulbecco's Modified Eagle Medium (DMEM), penicillin-streptomycin, trypsin, and fetal bovine serum (FBS) were purchased from Invitrogen. All other reagents and solvents were of analytical grade and used without further purification.

Antibodies and primers

SQSTM1/p62 rabbit monoclonal antibody (mAb, A19700), LC3A/LC3B rabbit polyclonal antibody (pAb, A5618), and PD-L1/CD274 rabbit pAb (#A1645) were purchased from ABclonal Technology. GAPDH rabbit mAb (GB15004-100) and

148 Cy5-conjugated goat anti-rabbit IgG (H + L) were obtained from Servicebio (Wuhan,

149 China). Horseradish peroxidase (HRP)-conjugated anti-rabbit IgG secondary mAb

150 (#7074) and Ki67 rabbit mAb (#ab92742) were purchased from Cell Signaling

151 Technology (CST) and Abcam, respectively. Additional antibodies obtained from

152 Abcom include: Alexa Fluor® 647-conjugated recombinant anti-calreticulin (CRT)

153 antibody (#ab196159), recombinant anti-Nrf2 antibody (#ab313825), recombinant

154 anti-heme oxygenase 1 antibody (#ab189491), anti-glutathione reductase antibody

155 (#ab124995), anti-NQO1 antibody (#ab80588), anti-superoxide dismutase 3/EC-SOD

156 antibody (#ab80946), and anti-glutathione peroxidase 4 antibody (#ab125066). Flow

157 cytometry antibodies were obtained as follows: from BD Pharmingen

158 -BV510-conjugated rat anti-mouse CD45 (563891), FITC-conjugated rat anti-mouse

159 CD11b (557396), Alexa Fluor 700-conjugated hamster anti-mouse CD3e (557984),

160 Cy5.5-conjugated Rat Anti-Mouse CD8a (551162), BV421-conjugated rat anti-mouse

161 IFN-γ (563376), FITC Rat Anti-Mouse I-A/I-E (2G9) antibody (562009), and PE

162 Hamster Anti-Mouse CD11c (HL3) antibody (557401); from BioLegend-Brilliant

163 Violet 650™anti-mouse CD86c (105036), FITC anti-mouse CD80 Antibody (104706),

164 and APC-conjugated recombinant anti-human/mouse granzyme B (372204). The

165 primers for quantitative reverse transcription polymerase chain reaction (qRT-PCR)

166 are as follows:

Gene	Forward	Reverse
Nfr2 (human)	CACATCCAGTCAGAAACCAGTGG	GGA ATGTCTGCGCCAAAAGCTG
Nfr2 (mouse)	CAGCATAGAGCAGGACATGGAG	GAACAGCGGTAGTATCAGCCAG
HO-1 (human)	CCAGGCAGAGAATGCTGAGTTC	AAGACTGGGCTCTCCTTGTTGC
HO-1 (mouse)	CACTCTGGAGATGACACCTGAG	GTGTTCCTCTGTCAGCATCACC
GR (human)	TATGTGAGCCGCCTGAATGCCA	CACTGACCTCTATTGTGGGCTTG
GR (mouse)	GTTTACCGCTCCACACATCCTG	GCTGAAAGAAGCCATCACTGGTG
NQO1 (human)	CCTGCCATTCTGAAAGGCTGGT	GTGGTGATGGAAAGCACTGCCT
NQO1 (mouse)	GCCGAACACAAGAAGCTGGAAG	GGCAAATCCTGCTACGAGCACT
SOD (human)	ACGCTGGCGAGGACGACCTG	GCTTCTTGCGCTCTGAGTGCTC
SOD (mouse)	GACCTGGTTGAGAAGATAGGCG	TGGCTGATGGTTGTACCCTGCA
GPX4 (human)	ACAAGAACGGCTGCGTGGTGAA	GCCACACACTTGTGGAGCTAGA

| GPX4 (mouse) | CCTCTGCTGCAAGAGCCTCCC | CTTATCCAGGCAGACCATGTGC |

167

Preparation and characterization of NP

Reduction-responsive RNAi NP were prepared using a modified nanoprecipitation method as previously described [8, 14, 15]. In brief, 3-MA was initially dissolved in DMF, followed by the addition of 1 nM of siPD-L1 (from a 0.1 nM/μL aqueous solution) at varying N/P molar ratios. Subsequently, P-18 in DMF and 200 μL of Meo-PEG-S-S-PLGA copolymer solution (20 mg/mL in DMF) were added to the mixture. The resulting solution was then slowly added dropwise into 5 mL of deionized water under vigorous stirring at 1000 rpm. The formed NP were transferred to an ultrafiltration device (EMD Millipore, MWCO 100 K) and centrifuged to eliminate organic solvents and unbound compounds. After rinsing with deionized water, the final formulation, designated as NPs(3-MA/siNrf2/P-18), was resuspended in deionized water at a siRNA concentration of 1 nM/mL. Control NP, denoted as NPs(G0-C14/siNrf2/P-18), were prepared using the same protocol, substituting 3-MA with the cationic lipid-like compound G0-C14 (5 mg/mL in DMF). The hydrodynamic diameter and zeta potential of all NP were measured by dynamic light scattering (DLS, Malvern, USA), and their morphology was characterized using a transmission electron microscope (TEM, FEI, USA). To evaluate encapsulation efficiency of siRNA, 3-MA, and P-18, Cy5-labeled siNrf2 was encapsulated into NP using the same procedure, yielding NPs(3-MA/Cy5-siNrf2/P-18). A 5 μL aliquot of the NP suspension was diluted 20-fold in DMSO, and the fluorescence intensity of Cy5-labeled siNrf2 was measured. Additionally, UV absorption at 413 nm (for P-18) and 279 nm (for 3-MA) were recorded using a Synergy HT multi-mode microplate reader (BioTek, USA). Encapsulation efficiencies were calculated by comparison with respective standard curves.

Cell culture

The murine BCa cell line 4T1 and the human BCa cell line MDA-MB-231 were cultured in DMEM supplemented with 10% FBS and 1% penicillin-streptomycin.

196 Cells were maintained at 37 °C in a humidified incubator containing 5% CO_2. Mouse

197 bone marrow-derived primary cells were cultured under the same conditions using

198 DMEM supplemented with 10% FBS. All animal procedures were approved by the

199 Institutional Review Board (IRB) of Sun Yat-sen Memorial Hospital.

200

201 *In vitro* cargo release

202 The NPs(3-MA/Cy5-siNrf2/P-18) were dispersed in 1 mL of PBS and transferred into

203 a Float-a-lyzer G2 dialysis device (MWCO 100 kDa, Spectrum). The device was

204 immersed in PBS with or without 10 mM GSH and incubated at 37 °C. At

205 predetermined time intervals, 5 μL of the NP solution was collected and diluted

206 20-fold with DMSO. The fluorescence intensity of Cy5-labeled siNrf2, along with UV

207 absorbance at 413 nm for P-18 and 279 nm for 3-MA, was measured using a Synergy

208 HT multi-mode microplate reader. (BioTek, USA). Cumulative release was calculated

209 using the formula: Cumulative release (%) = $(M_t / M_\infty) \times 100$, where M_t is the

210 amount of Cy5-siNrf2, 3-MA or P-18 released at a time t, and M_∞ is the total amount

211 of each cargo initially loaded into the NP.

212

213 In vitro Nrf2 silencing and inhibition of downstream genes

214 4T1 and MDA-MB-231 cells were seeded in 6-well plates at a density of 50,000 cells

215 per well and cultured in 2 mL of DMEM supplemented with 10% FBS for 24 h.

216 Following incubation, NPs(3-MA/siNrf2/P-18) were added to the cells at a final

217 siRNA concentration of 30 nM. After an additional 24 h of incubation, the culture

218 medium was replaced with fresh medium, and cells were further incubated for an

219 additional 48 h. Subsequently, total RNA and protein were extracted from the cells to

220 assess the mRNA and protein expression levels of Nrf2, HO-1, NQO1, GR, SOD,

221 GPX4, and GAPDH, employing qRT-PCR and western blotting, respectively.

222

223 qRT-PCR

224 Total RNA was isolated from treated cells utilizing Trizol reagent, and 1 μg of total

RNA was reverse-transcribed into complementary DNA (cDNA) using the Superscript First-Strand cDNA Synthesis Kit (catalog number 18080-051, Invitrogen, USA), following the manufacturer's protocol. qRT-PCR was carried out employing the SYBR® Premix Ex Taq™ II kit (product code DRR081A, Takara, Japan) on a LightCycler 480 System (Roche, Switzerland).

Western blot

Protein samples were quantified using a bicinchoninic acid (BCA) protein assay kit (Pierce/Thermo Scientific) according to the manufacturer's protocol. Equal amounts of protein were separated by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred onto a polyvinylidene difluoride (PVDF) membranes. Membranes were blocked in PBS containing 3% bovine serum albumin (BSA) and 0.1% Tween 20 (PBST) for 1 h at room temperature. After blocking, membranes were incubated overnight at 4 °C with primary antibody diluted in PBS containing 1% BSA. Following three washes with PBST, membranes were incubated with HRP-conjugated anti-rabbit IgG secondary antibody for 1 h at 4 °C. After additional washing with PBST, protein bands were visualized using an enhanced chemiluminescence (ECL) detection system. Target proteins included Nrf2, HO-1, NOQ1, GR, SOD, GPX4, GAPDH, PD-L1, p62, LC3, and HMGB1.

Immunofluorescence (IF)

4T1 and MDA-MB-231 cells were seeded in 6-well plates at a density of 50 000 cells per well and treated with NPs(3-MA/siNrf2/P-18) at a final siRNA concentration of 30 nM as described above. After treatment, cells were fixed with paraformaldehyde (PFA) and permeabilized with 0.2% Triton X-100 in PBS solution for 5 min. Cells were then washed three times with PBS and blocked with PBS containing 3% BSA for 1 h at room temperature. Primary PD-L1 antibody, diluted in PBS solution containing 1% BSA, was added and incubated with the cells for 1 h at 4 °C. After three PBS washes, cells were incubated with Alexa Fluro 647-conjugated secondary

antibody for 1 h at 4 °C. After another set of PBS washes, nuclei were stained with Hoechst 33342. Fluorescence imaging was performed using a ZEISS 800 confocal laser scanning microscope (CLSM).

In vitro inhibition of mitophagy

4T1 and MDA-MB-231 cells were seeded into 6-well plates at a density of 50 000 cells per well and treated with NPs(3-MA/siNrf2/P-18) at a final siRNA concentration of 30 nM as previously described. After treatment, cells were stained with MitoTracker® Red CMXRos and LysoTracker™ Green DND-26 to evaluate the co-localization of lysosomes and mitochondria using CLSM. Following imaging, cells were harvested by trypsinization and total protein was extracted for western blot analysis of p62 and LC3 I/II expression.

Detection of intracellular ROS levels

4T1 and MDA-MB-231 cells were seeded into 6-well plates at a density of 50000 cells per well and treated with NPs(3-MA/siNrf2/P-18) under US irradiation, using a final siRNA concentration of 30 nM, as previously outlined. After treatment, cells were incubated with 5 μM CM-H2DCFDA, a ROS-sensitive fluorescent probe, for 15 min at 37 °C. Intracellular ROS levels were visualized using CLSM. Following imaging, cells were harvested and analyzed by flow cytometry utilizing a BD FACSAria™ III flow cytometry to quantitatively assess ROS accumulation.

Detection of ATP and CRT release

4T1 and MDA-MB-231 cells were seeded in 6-well plates at a density of 50 000 cells per well and treated with NPs(3-MA/siNrf2/P-18) under US irradiation at a final siRNA concentration of 30 nM, accordingly to the protocol described above. After treatment, the cell culture supernatants were collected for quantification of ATP and CRT levels. ATP concentration was measured using the ATP Determination Kit, while CRT levels were assessed using both the human CRT ELISA Kit (#E-EL-H0627) and

283 Recombinant Alexa Fluor® 647 anti-CRT antibody, according to the manufacturer's
284 protocol.

285

286 ***In vitro* proliferation and colony formation**
287 4T1 and MDA-MB-231 cells were seeded in 6-well plates at a density of 20,000 cells
288 per well and cultured in 2 mL of DMEM supplemented with 10% FBS for 24 h. Cells
289 were then treated with NPs(G0-C14/siNrf2/P-18), NPs(3-MA/siCTL/P-18), or
290 NPs(3-MA/siNrf2/P-18) at a siRNA concentration of 30 nM, with or without US
291 irradiation (3 min, 1 W/cm², 3 MHz, and 50% duty cycle). After 24 h of incubation,
292 cells were rinsed with PBS, and cell viability was assessed using the Alamar Blue
293 assay according to the manufacturer's protocol. After the viability measurement, the
294 Alamar Blue reagent was removed and cells were cultured in fresh medium. For the
295 colony formation assay, MDA-MB-231 and 4T1 cells were seeded in 6-well plates at
296 a density of 2,000 cells per well. The cells were treated with the same nanoparticle
297 formulations and conditions as described above. After seven days of incubation in
298 complete medium, colonies were fixed and stained with crystal violet. Colony
299 formation was observed using an MVX10 Macro View Dissecting Scope equipped
300 with an Olympus DP80 camera.

301

302 **Apoptosis analysis**
303 MDA-MB-231 and 4T1 cells were seeded into 6-well plates at a density of 50,000 per
304 well and cultured in 2 mL of DMEM supplemented with 10% FBS for 24 h. Cells
305 were then treated with NPs(G0-C14/siNrf2/P-18), NPs(3-MA/siCTL/P-18), or
306 NPs(3-MA/siNrf2/P-18) at a final siRNA concentration of 30 nM with or without US
307 irradiation, as described previously. After 24 h, cells were washed with PBS,
308 incubated in fresh medium for an additional 24 h, harvested, and stained using the
309 Annexin V-FITC/PI Apoptosis Detection Kit (AK12637, Elabscience). Apoptosis was
310 quantified by flow cytometry using a CytoFlex LX Flow Cytometry Analyzer
311 (Beckman Coulter).

Animals

Healthy female BALB/c mice (4-5 weeks old) were purchased from Sun Yat-sen University Experimental Animal Center (Guangzhou, China). All animal experiments were conducted in accordance with protocols approved by the Institutional Animal Care and Use Committee of Sun Yat-sen Memorial Hospital (#AEP20240215).

Pharmacokinetics

Healthy female BALB/c mice were randomly divided into three groups (n = 3) and administered an intravenous injection of one of the following formulations: (i) naked Cy5-labled siNrf2, or (ii) NPs(3-MA/Cy5-labled siNrf2/P-18) at a dose of 5 mg/kg P-18 and/or 6 mg/kg 3-MA and/or 1 nM siNrf2 per mouse. At predetermined time intervals post-injection, 20 μL of blood was collected via the orbital vein. The fluorescence intensity of Cy5-labeled siNrf2 in the blood samples was measured using fluorescence spectroscopy to assess circulation kinetics.

Orthotopic and lung metastatic tumor model

To establish the 4T1 orthotopic tumor model, 200 μL of a 4T1 cell suspension-comprising a 1:1 volume mixture of DMEM and Matrigel with a cell concentration of 1×10^7 cells/mL-was subcutaneously injected into the second pair of mammary fat pads of healthy female BALB/c mice. Once tumors reached approximately 100 mm^3 in volume, these tumor-bearing mice were enrolled in subsequent *in vivo* experiments. For the lung metastasis model, 2×10^5 luciferase-expressing 4T1 cells suspended in 100 μL of PBS were administered intravenously into healthy mice. Tumor progression was tracked by bioluminescence imaging using an IVIS Lumina III system (PerkinElmer, USA). Prior to imaging, D-luciferin was administered via intraperitoneal injection at a dose of 150 mg/kg. Average radiance values at tumor sites were used to quantify tumor burden.

Biodistribution

4T1 orthotopic tumor-bearing mice were randomly divided into two groups (n = 3) and administered intravenous injections of either: (i) unencapsulated Cy5-labeled siNrf2, or (ii) NPs(3-MA/Cy5-labeled siNrf2/P-18) at a dosage of 5 mg/kg P-18 and/or 6 mg/kg 3-MA and/or 1 nM siNrf2 per mouse. At 24 h post-injection, tumors and major organs were harvested and imaged using the IVIS Lumina III system (PerkinElmer, USA). The accumulation of Cy5-labeled siNrf2 in tumor and organ tissues was quantified based on fluorescence intensity using Image J software.

In vivo PD-L1 downregulation and Nrf2 silencing

4T1 orthotopic tumor-bearing mice were randomly divided into six groups (n = 5) and treated with daily intravenous injections of the following formulations: (i) PBS, (ii): NPs(3-MA/siCTL/P-18), (iii) NPs(G0-C14/siNrf2/P-18), (iv) NPs(3-MA/siCTL/P-18) + US, (v) NPs(G0-C14/siNrf2/P-18) + US, and (vi) NPs(3-MA/siNrf2/P-18) + US. Each injection was administered at a dose of 5 mg/kg P-18 and/or 6 mg/kg 3-MA and/or 1 nM siNrf2 per mouse. After three consecutive injections, mice were sacrificed 24 h following the final dose. Tumors were excised, and total protein was extracted for western blot analysis of PD-L1 and Nrf2 expression. In parallel, tumor tissues were homogenized into single-cell suspensions for flow cytometry analysis to evaluate DCs maturation, CD8$^+$ T cell infiltration, and granzyme B and IFN-γ production by CD8$^+$ T cells, according to the manufacturer's protocol.

Inhibition of orthotopic tumor growth

4T1 orthotopic tumor-bearing mice were randomly divided into six groups (n = 5) and treated with four consecutive intravenous injections of the following formulations: (i) PBS, (ii) NPs(3-MA/siCTL/P-18), (iii) NPs(G0-C14/siNrf2/P-18), (iv) NPs(3-MA/siCTL/P-18) + US, (v) NPs(G0-C14/siNrf2/P-18) + US, and (vi) NPs(3-MA/siNrf2/P-18) + US. Injections were administered once every two days at a dose of 5 mg/kg P-18 and/or 6 mg/kg 3-MA and/or 1 nM siNrf2 per mouse. Tumor growth was monitored every two days by measuring the shortest (W) and longest (L)

diameters with a caliper. Tumor volume was calculated using the formula: $V = W^2 \times L/2$. At the experimental endpoint, tumors were collected and sectioned for TUNEL and Ki67 immunohistochemical staining according to the manufacturer's protocol.

Inhibition of lung metastatic tumor growth

4T1 lung metastasis tumor-bearing mice were randomly assigned into six groups (n = 5) and treated with four consecutive intravenous injections of the following: (i) PBS, (ii) NPs(3-MA/siCTL/P-18), (iii) NPs(G0-C14/siNrf2/P-18), (iv) NPs(3-MA/siCTL/P-18) + US, (v) NPs(G0-C14/siNrf2/P-18) + US, and (vi) NPs(3-MA/siNrf2/P-18) + US. The injections were administered once every two days at a dose of 5 mg/kg P-18 and/or 6 mg/kg 3-MA and/or 1 nM siNrf2 per mouse. Lung tumor progression was monitored on days 0, 7, and 14 using a bioluminescence imaging system, following the procedures previously described. At the conclusion of the experiment, tumors were excised and sectioned for TUNEL and Ki67 staining, in accordance with the manufacturer's instructions.

Immunohistochemistry (IHC)

IHC staining was conducted on formalin-fixed, paraffin-embedded tumor sections. In summary, tumor slides were initially heated to 60 ° C for 1 h, followed by deparaffinization using xylene (three washed, 5 min each) and rehydration through a graded ethanol alcohol series. Antigen retrieval was performed using DAKO Target Retrieval Solution at 95-99 °C for 40 min, followed by washing in distilled water. Endogenous peroxidase activity was blocked using DAKO peroxidase blocking buffer for 5 min. After washing, slides were incubated with the appropriate primary antibody diluted in DAKO antibody diluent for 1 h at room temperature. Slides were then washed and incubated with a peroxidase-conjugated polymer for 30 min. Following a final wash, staining was developed using DAB^+ substrate-chromogen solution and counterstained with hematoxylin. Stained slides were mounted and imaged using an MVX10 MacroView Dissecting Scope equipped with an Olympus DP80 camera.

Blood and histological analysis

Healthy female BALB/c mice were randomly divided into six groups (n = 3) and treated with intravenous injections of the following formulations: (i) PBS, (ii) NPs(3-MA/siCTL/P-18), (iii) NPs(G0-C14/siNrf2/P-18), (iv) NPs(3-MA/siCTL/P-18) + US, (v) NPs(G0-C14/siNrf2/P-18) + US, and (vi) NPs(3-MA/siNrf2/P-18) + US. Following three consecutive daily injections, blood samples were collected 24 h after the final dose, and serum was isolated for analysis of standard hematological and biochemical parameters. Major organs, including the heart, liver, spleen, lung, and kidneys, were harvested and processed for histological examination.

Statistical analysis

All quantitative data are presented as mean ± standard deviation (SD) from at least three independent experiments. Graphpad Prism software (version 8.0) was used for data visualization, statistical analysis, and figure generation. Specific sample sizes used for each experiment are detailed in the corresponding figure legends. Statistical comparisons between two groups were performed using two-tailed Student's *t*-test, while comparisons among multiple groups were conducted using one-way ANOVA. A P-value < 0.05 was considered statistically significant.

Results and discussion

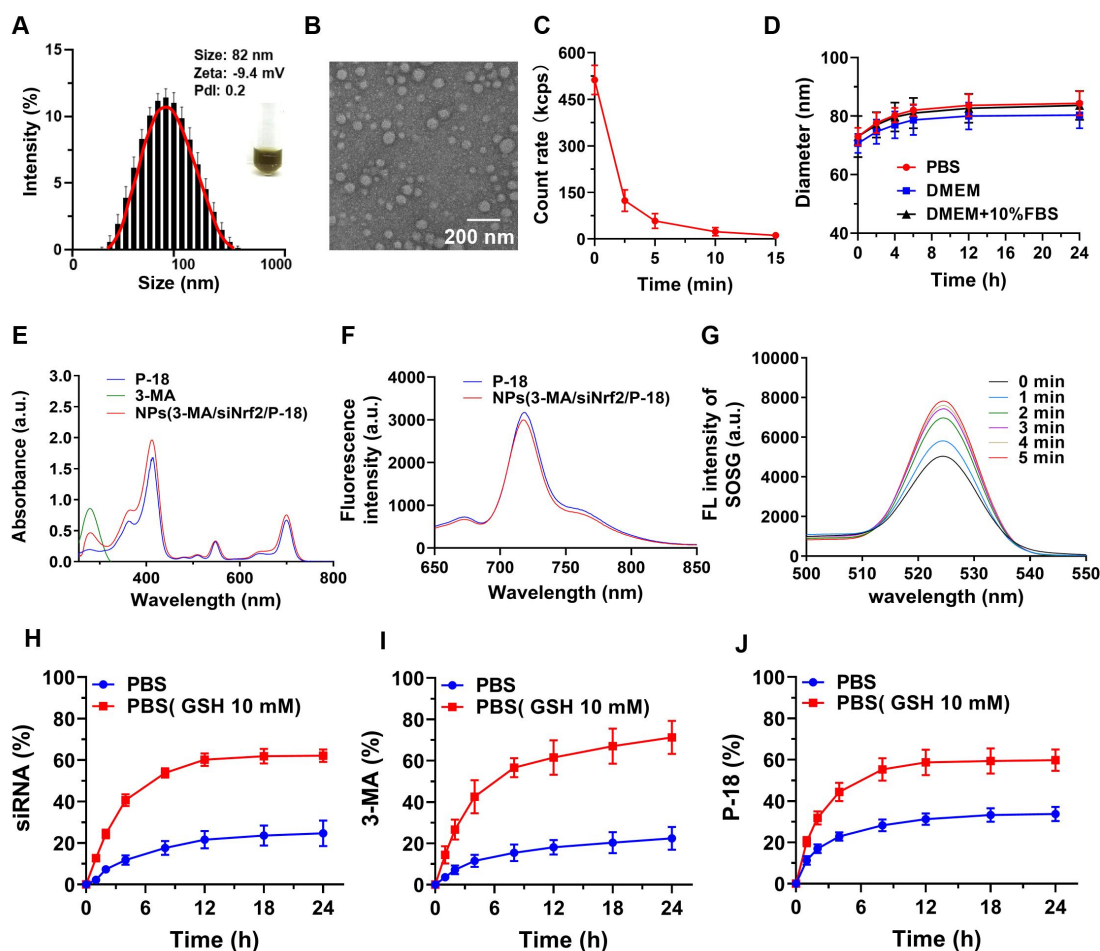**Preparation and characterizations of NPs(3-MA/siNrf2/P-18)**

To facilitate clinical translation, a simplified and efficient method was employed for the synthesis of a reduction-responsive RNAi NP. The NP were prepared using a modified nanoprecipitation method [16-18], in which the amphiphilic copolymer Meo-PEG-S-S-PLGA and the sonosensitizer P-18 were dissolved in dimethyl formamide (DMF) and subsequently mixed with aqueous solutions of siNrf2 and 3-MA. The mixture was then added dropwise into deionized water under vigorous stirring. In aqueous conditions, the Meo-PEG-S-S-PLGA polymer self-assembled into spherical NP featuring a hydrophobic PLGA inner core and hydrophilic PEG outer

shell [17]. Within this formulation, 3-MA formed electrostatic complexes with siNrf2, enabling co-encapsulation within the PLGA core alongside P-18 (Scheme 1). By adjusting the feed ratio of 3-MA to siNrf2 (**Figure S1, Supporting Information**), an optimal nitrogen-to-phosphate (N/P) molar ratio of 95:1 was selected, yielding well-defined spherical NPs(3-MA/siNrf2/P-18) with an average diameter of ~82 nm (**Figure 1A-1B; Figure R, Supporting Information**). These NP demonstrated high encapsulation efficiencies: ~76% for siNrf2, ~93% for 3-MA, and ~62% for P-18. UV absorbance spectra confirmed successful loading, with characteristic peaks at 413 nm, 548 nm, and 700 nm for P-18, and 279 nm for 3-MA (**Figure 1E**). Fluorescence (FL) analysis revealed an emission peak at 720 nm for the NP, corresponding to the free P-18 spectrum, further confirming its successful incorporation (**Figure 1F**). As a control, NPs(G0-C14/siNrf2/P-18) were synthesized by replacing 3-MA with the amphiphilic cationic compound G0-C14, developed in a previous study (**Figure S2, Supporting Information**) [19-21]. The redox-responsiveness of the NP was validated by its structural disassembly in the presence of 10 mM glutathione (GSH), which mimics the TME (**Figure 1C**), while stability was retained under normal physiological conditions (**Figure 1D**). This disassembly triggered the rapid release of siNrf2 (**Figure 1H**), 3-MA (**Figure 1I**), and P-18 (**Figure 1J**). The NPs(3-MA/siNrf2/P-18) exhibited robust colloidal stability in PBS, DMEM, and 10% FBS over 24 h, with minimal variation in particle size, attributed to the PEGylated surface providing steric hindrance that mitigates protein adsorption and aggregation-a well-documented strategy for prolonged circulation [22]. The disulfide linker in Meo-PEG-S-S-PLGA enabled selective drug release under TME conditions, where elevated GSH levels (~10 mM) trigger rapid NP disassembly. This dual-functionality design-PEG-mediated circulation stability and TME-responsive release-minimizes off-target leakages while maximizing tumor-specific drug delivery, consistent with previous reports [17]. Plasma protein interaction studies showed only slight increases in NP size after 24 h of incubation in mouse plasma, indicating strong anti-fouling properties (**Figure S3, Supporting Information**). The low protein binding was

attributed to the hydrophilic PEG corona of Meo-PEG-S-S-PLGA, which forms a "stealth" protective layer through steric hindrance and hydrogen bonding, thereby limiting opsonin adsorption (e.g., immunoglobulins and complement proteins) and reducing recognition by the reticuloendothelial system [22]. These results collectively support the suitability of the NP as a stable and effective delivery vehicle for cancer therapy.

To evaluate the sonodynamic performance of the P-18-loaded NP, singlet oxygen generation was assessed under US irradiation (3 min, 1 W/cm^2, 3 MHz, and 50% duty cycle). The singlet oxygen sensor green (SOSG) probe was employed to monitor singlet oxygen levels. Upon US exposure, a sharp increase in SOSG absorbance was observed over time, indicating progressive generation of singlet oxygen by NPs(3-MA/siNrf2/P-18) (**Figure 1G**). These findings confirmed that the NP possesses excellent sonodynamic properties and can effectively induce high levels of ROS under US stimulation.
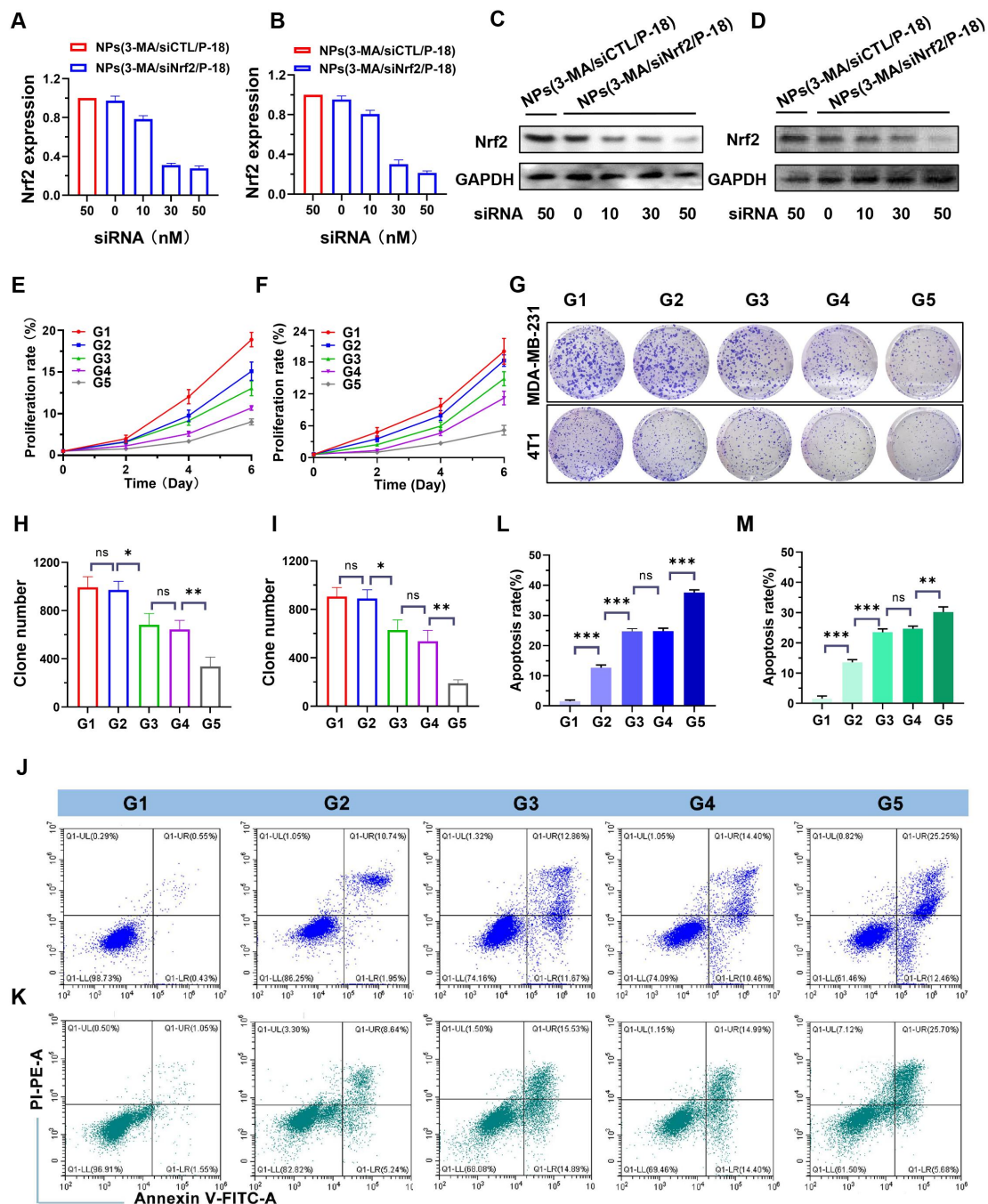
Figure 1. Characterization and stability of NPs(3-MA/siNrf2/P-18). (A) DLS analysis showing the hydrodynamic size distribution of NPs(3-MA/siNrf2/P-18). (B) TEM image illustrating the morphology and uniform size of the nanoparticles (scale bar: 200 nm). (C) Quantitative analysis of nanoparticle degradation over time in the presence of 10 mM glutathione (GSH), simulating reductive tumor microenvironment conditions. (D) Assessment of nanoparticle size stability following incubation in PBS, DMEM, and DMEM supplemented with 10% FBS over various time points. (E) UV-Vis absorption spectra confirming successful encapsulation of P-18 and characterization of optical properties. (F) Fluorescence emission spectra of free P-18 versus P-18-loaded nanoparticles in DMSO, indicating retained photophysical properties post-encapsulation. (G) Singlet oxygen generation, measured via SOSG fluorescence, from NPs(3-MA/siNrf2/P-18) subjected to ultrasound (US) irradiation for varying durations. (H-J) Cumulative release profiles of siRNA, 3-MA, and P-18

485 from the nanoplatform over 24 hours in PBS and GSH-containing PBS (10 mM),

486 demonstrating redox-responsive drug release behavior.

487

488 *In vitro* **Nrf2 silencing and cellular functional assessment of reduction-responsive**

489 **RNAi NP**

490 Following the successful development of the reduction-responsive RNAi NP, its

491 ability to silence Nrf2 expression was evaluated. NPs(3-MA/siNrf2/P-18) were

492 incubated with human-derived BCa cells (MDA-MB-231) and mouse-derived BCa

493 cells (4T1) to assess Nrf2 knockdown efficiency. Nrf2 is often overexpressed in

494 cancer cells and has been associated with the promotion of angiogenesis, drug

495 resistance, cancer stem cell formation, and metastasis [23]. Aberrant expression of

496 Nrf2 contributes to decreased therapeutic efficacy and confers cytoprotective

497 advantages to tumor cells. Therefore, Nrf2 knockdown is considered a promising

498 strategy to disrupt these cancer-promoting pathways. Quantitative results indicated

499 that treatment with NPs(3-MA/siNrf2/P-18) led to a dose-dependent decrease in Nrf2

500 mRNA levels in both MDA-MB-231 (**Figure 2A**) and 4T1 cells (**Figure 2B**). A

501 corresponding reduction in Nrf2 protein expression was also observed in

502 MDA-MB-231 (**Figure 2C**) and 4T1 cells (**Figure 2D**) following

503 NPs(3-MA/siNrf2/P-18) treatment, confirming efficient RNA interference after

504 cellular uptake of the NP. In addition to Nrf2 silencing, functional studies

505 demonstrated that these nanoparticles significantly inhibited cell proliferation and

506 induced apoptosis under US irradiation. As shown in **Figure 2E** and **Figure 2F**,

507 treatment with NPs(3-MA/siNrf2/P-18) markedly reduced proliferation in

508 MDA-MB-231 (**Figure 2E**) and 4T1 cells (**Figure 2F**), by approximately fivefold

509 relative to controls. The results of the colony formation assay (**Figure 2G**) further

510 supported these findings, showing that US-activated NPs(3-MA/siNrf2/P-18)

511 treatment significantly suppressed long-term growth potential in MDA-MB-231

512 (**Figure 2H**) and 4T1 cells (**Figure 2I**). Furthermore, flow cytometry analysis

513 indicated a pronounced increase in apoptosis upon treatment with

514 NPs(3-MA/siNrf2/P-18) in both cell lines, attributed to combined mitophagy

515 inhibition and Nrf2 silencing. This effect was enhanced under US-induced

516 sonodynamic therapy conditions, as evidenced by apoptosis rates in MDA-MB-231

517 (**Figure 2J** and **Figure 2L**) and 4T1 cells (**Figure 2K** and **Figure 2M**). Collectively,

518 these findings suggest that the reduction-responsive RNAi NP effectively silences

519 Nrf2 expression, suppresses cellular proliferation, and enhances apoptosis in BCa

520 cells through the synergistic effects of RNA interference, mitophagy inhibition by

521 3-MA, and sonodynamic ROS induction via P-18.

522

Figure 2. *In vitro* evaluation of Nrf2 silencing, proliferation inhibition, and apoptosis induction by NPs(3-MA/siNrf2/P-18). (A, B) Quantitative real-time PCR (qRT-PCR) analysis of relative Nrf2 mRNA expression in MDA-MB-231 and 4T1 cells, respectively, following treatment with NPs(3-MA/siNrf2/P-18). (C, D) Western blot analysis of Nrf2 protein levels in MDA-MB-231 and 4T1 cells after the indicated treatments. (E, F) Cell proliferation assays showing growth inhibition in MDA-MB-231 and 4T1 cells treated with NPs(3-MA/siNrf2/P-18) with or without

22

ultrasound (US) irradiation. (G) Representative colony formation assay images and (H, I) Quantitative analysis of colony numbers in MDA-MB-231 and 4T1 cells, respectively, following treatment with NPs(3-MA/siNrf2/P-18) under US irradiation. (J-M) Flow cytometry analysis of apoptosis rates in MDA-MB-231 and 4T1 cells after treatment with NPs(3-MA/siNrf2/P-18) with US activation. G1: Blank; G2: NPs(3-MA/siNrf2/P-18); G3: NPs(3-MA/siCTL/P-18) + US; G4: NPs(G0-C14/siNrf2/P-18) + US; G5: NPS(3-MA/siNrf2/P-18) + US. Data are presented as mean ± SD (n = 3). Error bars represent standard deviation. Statistical analysis was performed using one-way ANOVA for multiple comparisons. Significance is indicated as *P < 0.05, **P < 0.01 and ***P < 0.001, ns, no significance.

Enhancing sonodynamic therapy and ICD through dual inhibition of mitophagy and Nrf2 pathways

Mitochondrial autophagy (mitophagy) serves as an important quality control mechanism that eliminates damaged mitochondria and excess ROS, thereby preserving cellular homeostasis [7]. Beyond this protective role, mitophagy helps limit the deleterious effects of ROS accumulation. Inhibition of mitophagy has been shown to enhance ROS levels within cells [24, 25]. Therefore, the potential of the reduction-responsive RNAi NP to simultaneously disrupt mitophagy and the Nrf2-associated antioxidant system in BCa cells was evaluated to determine its capacity to potentiate SDT-mediated ROS accumulation. To explore the impact of Nrf2 silencing on mitochondrial function, JC-1 staining and MitoSOX Red assays were performed. JC-1 staining revealed a significant reduction in the red/green fluorescence ratio in both MDA-MB-231 and 4T1 cells following Nrf2 knockdown (**Figure S4, Supporting Information**), indicating mitochondrial membrane potential ($\Delta\Psi$m) collapse. This effect was further exacerbated under US irradiation, confirming that Nrf2 deficiency sensitizes mitochondria to SDT-induced damage. MitoSOX Red staining demonstrated that mtROS levels increased substantially upon Nrf2 silencing,
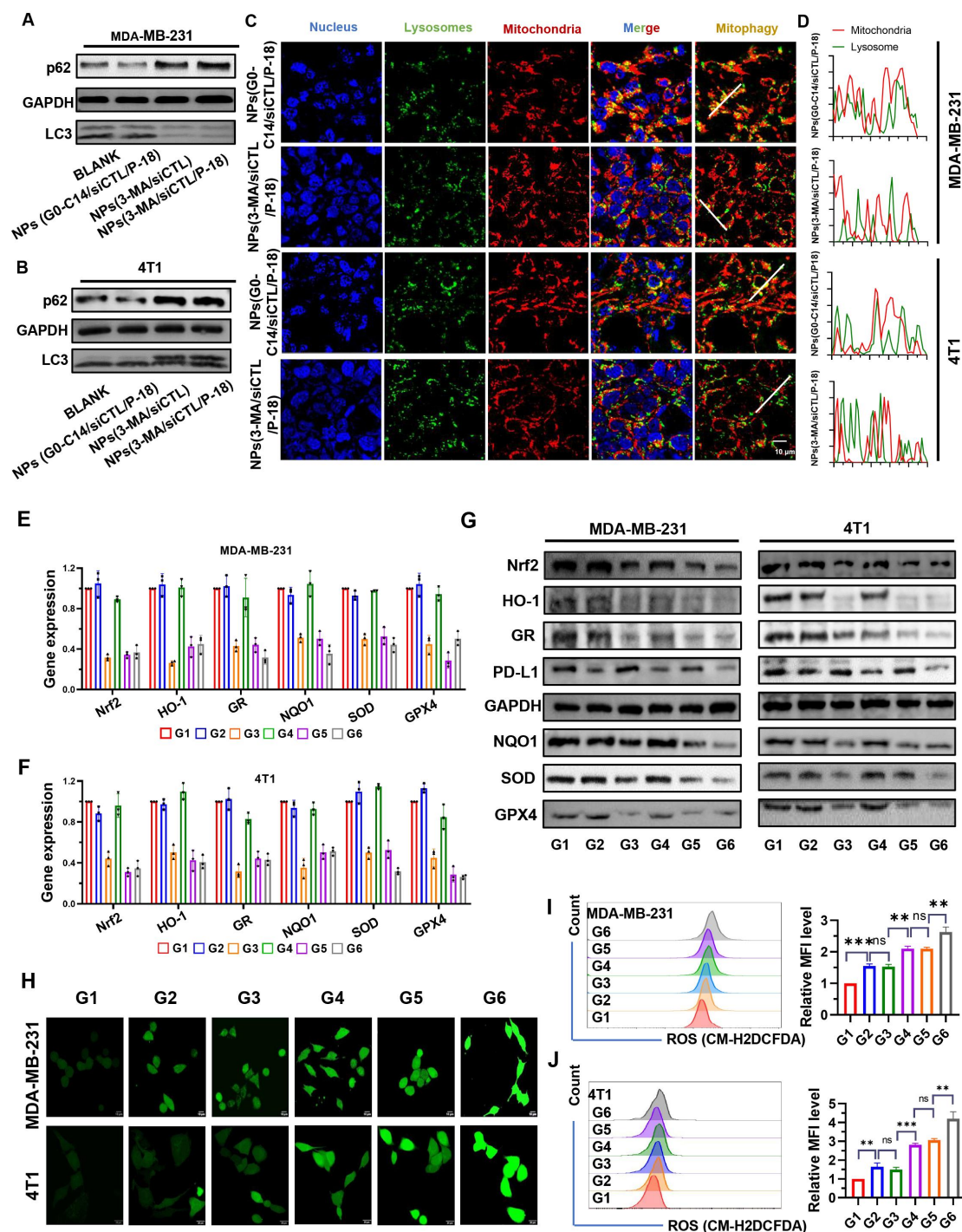
with combined SDT treatment inducing a 2.8-fold increase in 4T1 cells and a 3.4-fold increase in MDA-MB-231 cells (**Figure S5, Supporting Information**), thereby overwhelming cellular antioxidant defenses. These findings demonstrate that Nrf2 silencing disrupts mitochondrial redox balance through two interrelated mechanisms: (i) antioxidant Depletion-loss of key enzymes such as SOD2 and GPX1 impairs mtROS detoxification [26]; and (ii) metabolic destabilization-mtROS overload destabilizes ETC complexes, resulting in exacerbated electron leakage and propagation of oxidative stress, an effect further intensified by mitophagy inhibition [27]. This dual inhibition strategy establishes a self-amplifying cycle of mitochondrial damage, underscoring the necessity of targeting both pathways to achieve irreversible mitochondrial dysfunction and optimize SDT efficacy.

p62 is a well-established substrate for autophagy, and its expression level is inversely correlated with autophagic activity [28]. During autophagy, cytoplasmic LC3-I undergoes enzymatic processing and is subsequently conjugated to phosphatidylethanolamine (PE), forming the membrane-associated LC3-II, which is recruited to autophagosomal membranes [29]. Therefore, the LC3-II/I ratio serves as a widely used indicator of autophagic flux. Treatment with 3-MA loaded NP resulted in increased p62 accumulation and a decreased LC3-II/LC3-I ratio in both MDA-MB-231 (**Figure 3A**) and 4T1 cells (**Figure 3B**), indicating reduced autophagy levels. Moreover, compared to NPs(G0-C14/siCTL/P-18), treatment with NPs(3-MA/siCTL/P-18) inhibited the colocalization of mitochondria and lysosomes in both cell lines (**Figure 3C-3D**), confirming successful mitophagy inhibition by the 3-MA loaded formulation. 3-MA, a PI3K inhibitor, is widely used to inhibit autophagy by targeting class I PI3K [30]. These results confirm that the synthesized NP effectively replicates the autophagy-inhibitory function of the free drug 3-MA. Moreover, the expression of Nrf2-regulated downstream antioxidant genes-including heme oxygenase-1 (HO-1) [31, 32], glutathione reductase (GR) [33], NAD(P)H: quinone oxidoreductase (NQO1) [34, 35], superoxide dismutase (SOD), and glutathione peroxidase 4 (GPX4) [36, 37], was significantly downregulated at both

588   the mRNA (**Figure 3E-3F**) and protein levels (**Figure 3G**) following Nrf2 silencing.

589   In addition, PD-L1 expression was also reduced, suggesting that Nrf2 may contribute

590   to the regulation of immune checkpoint molecules. This is consistent with previous

591   reports indicating 3-MA can downregulate PD-L1 expression via an

592   NF-κB-dependent pathway [38, 39]. To further confirm the mechanism, BAY11-7082,

593   a NF-κB inhibitor, was used to mimic the effect of 3-MA and 3-MA-containing NP on

594   the NF-κB/PD-L1 signaling axis and p-p65 nuclear localization. These inhibitory

595   effects were reversed by MC-LR, a phosphatase inhibitor that indirectly promotes p65

596   phosphorylation by suppressing dephosphorylation, indicating that both 3-MA and

597   3-MA-loaded NP downregulate PD-L1 expression via the NF-κB pathway (**Figure S6,**

598   **Supporting Information**). To exclude the possibility that G0-C14 itself influences

599   mitophagy or PD-L1 expression, MDA-MB-231 and 4T1 cells were treated with

600   various concentrations of G0-C14. Western blot analysis showed no significant

601   changes in LC3, p62, or PD-L1 levels (**Figure S7, Supporting Information**),

602   confirming that G0-C14 functions solely as a cationic carrier to enhance siRNA

603   delivery, without directly affecting autophagic flux or immune checkpoint regulation.

604   Following confirmation of Nrf2 and mitophagy inhibition in BCa cells, intracellular

605   ROS levels were evaluated via confocal fluorescence imaging (**Figure 3H**) and flow

606   cytometry (**Figure 3I-3J**). Both methods demonstrated significantly increased ROS

607   accumulation in MDA-MB-231 and 4T1 cells after treatment with

608   NPs(siNrf2/3-MA/P-18) under US irradiation. In summary, the reduction-responsive

609   RNAi NP facilitates cascade amplification and accumulation of ROS within tumor

610   cells by simultaneously silencing Nrf2 silencing and inhibiting mitochondrial
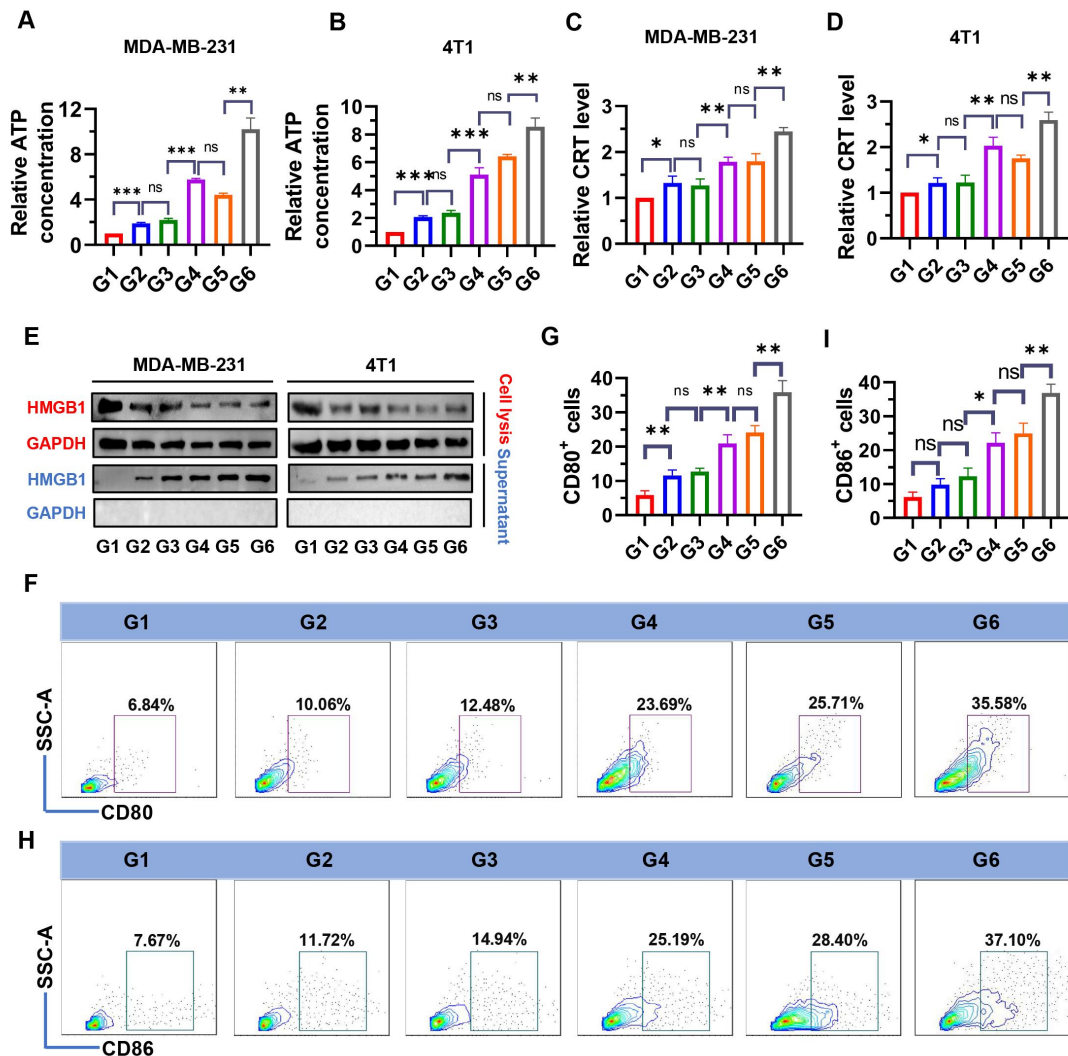
611   autophagy.

Figure 3. Evaluation of autophagy inhibition, antioxidant gene regulation, PD-L1 expression, and ROS accumulation following treatment with NPs(3-MA/siNrf2/P-18). (A, B) Western blot analysis of autophagy-related proteins p62 and LC3 in MDA-MB-231 and 4T1 cells after treatment with NPs(3-MA/siNrf2/P-18), indicating mitophagy inhibition. (C, D) Confocal microscopy images showing colocalization of mitochondria and lysosomes in MDA-MB-231 and 4T1 cells, respectively,

demonstrating altered mitophagy dynamics following treatment. (E, F) Quantitative RT-PCR analysis of antioxidant-related gene expression (Nrf2, HO-1, GR, NQO1, SOD, and GPX4) in MDA-MB-231 and 4T1 cells after nanoparticle treatment. (G) Western blot analysis of PD-L1 and antioxidant protein levels (Nrf2, HO-1, GR, NQO1, SOD, and GPX4) in both cell lines post-treatment, indicating immunomodulatory and redox-disruptive effects. (H) Confocal fluorescence imaging of intracellular ROS generation in MDA-MB-231 and 4T1 cells following US-activated treatment with NPs(3-MA/siNrf2/P-18). (I, J) Flow cytometry analysis and quantification of ROS levels in MDA-MB-231 and 4T1 cells, respectively, after various treatments with or without ultrasound irradiation. Treatment groups: G1: PBS; G2: NPs(3-MA/siCTL/P-18); G3: NPs(G0-C14/siNrf2/P-18); G4: NPs(3-MA/siCTL/P-18) + US; G5: NPs(G0-C14/siNrf2/P-18) + US; G6: NPs(3-MA/siNrf2/P-18) + US. Data are presented as mean ± SD (n = 3). Error bars represent standard deviation. Statistical analysis was performed using one-way ANOVA for multiple comparisons. Significance levels are indicated as *P < 0.05, **P < 0.01 and ***P < 0.001, ns, no significance.

Increased ROS are an important cause of ICD induction [40]. To examine whether NPs(3-MA/siNrf2/P-18) could trigger intensive ICD, the release of DAMPs, including ATP, CRT, and HMGB1 were evaluated in dying BCa cells. As shown in **Figure 4**, treatment with NPs(3-MA/siNrf2/P-18) under US irradiation significantly promoted ATP released from MDA-MB-231 (**Figure 4A**) and 4T1 cells (**Figure 4B**), as well as CRT exposure in MDA-MB-231 (**Figure 4C**) and 4T1 cells (**Figure 4D**). In addition, HMGB1 protein levels in the culture supernatant were significantly increased following NPs(3-MA/siNrf2/P-18) treatment with US irradiation (**Figure 4E**), indicating enhanced passive cell death. These results demonstrate that SDT induced by NPs(3-MA/siNrf2/P-18) effectively promotes apoptosis and ICD, leading to substantial release of DAMPs from BCa cells. DAMPs released during ICD are recognized by pattern recognition receptors (PRRs) on the surface of DCs, initiating a

series of cytological responses that ultimately activate both innate and adaptive immune responses [14]. To determine whether NPs(3-MA/siNrf2/P-18)-induced ICD could successfully activate DCs *in vitro*, a co-culture system was established using mouse-derived DCs and 4T1 cells in a Boyden chamber device (**Figure S8, Supporting Information**). Following treatment with NPs(3-MA/siNrf2/P-18) and US irradiation, a significant increase in the percentage of $CD80^+$ and $CD86^+$ DCs was observed-by approximately fivefold-compared to controls (**Figure 4F-4I**), indicating robust DCs activation. To rule out the possibility that US alone contributed to ICD induction, an additional control group with US-only treatment was included. 4T1 and MDA-MB-231 cells were exposed to the same US conditions described in the study, and ICD markers were assessed. The results confirmed that US treatment alone did not induce significant changes in ICD indicators (**Figure S9, Supporting Information**).
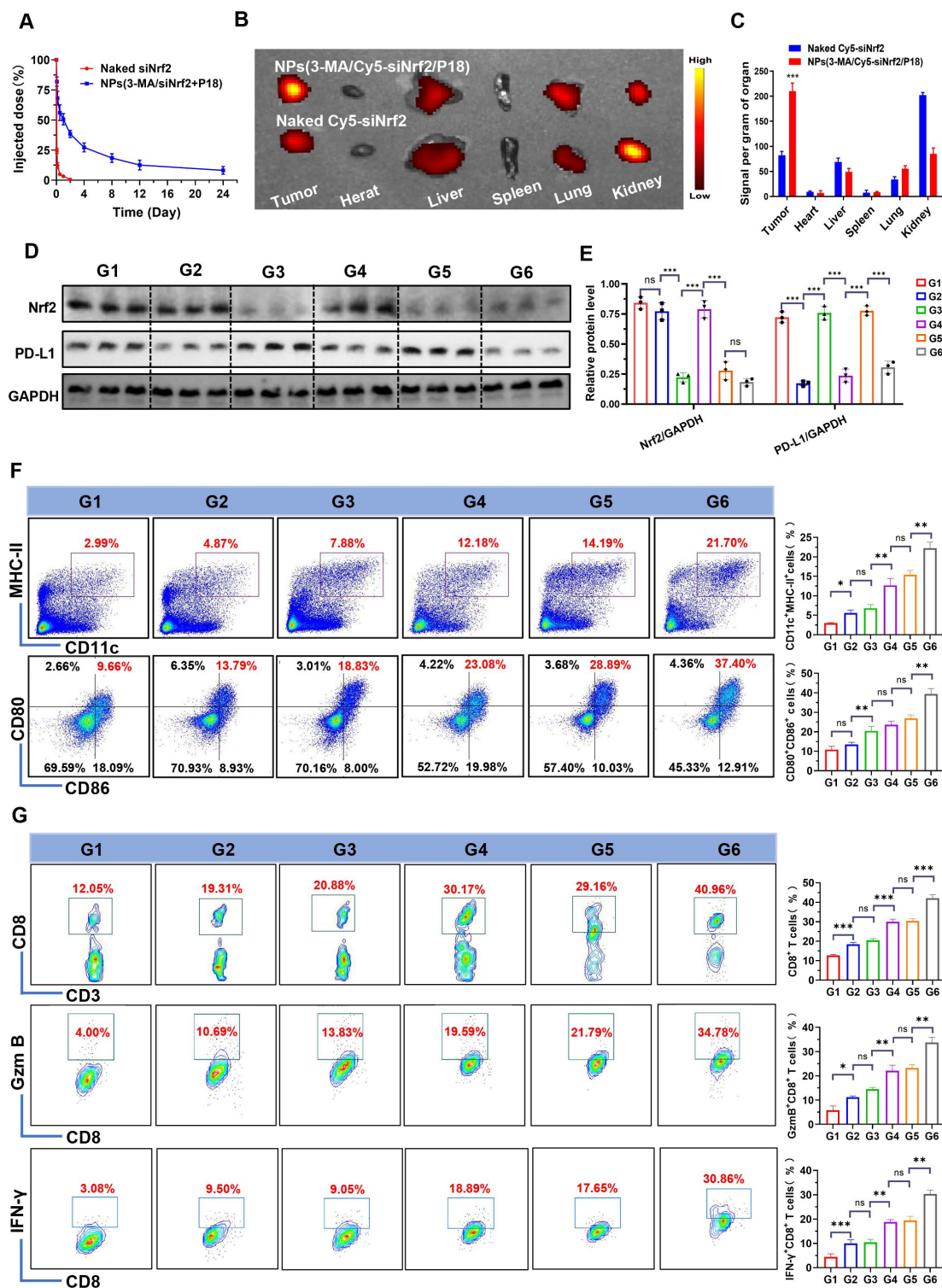
Figure 4. Evaluation of ICD markers and DCs activation following NPs(3-MA/siNrf2/P-18) treatment under ultrasound (US) irradiation. (A, B) Quantification of extracellular ATP levels in the supernatant of MDA-MB-231 and 4T1 cells, respectively, determined by ELISA after treatment with NPs(3-MA/siNrf2/P-18) with US activation. (C, D) Flow cytometry analysis of CRT surface exposure on MDA-MB-231 and 4T1 cells, respectively, indicating ICD induction. (E) HMGB1 protein levels in cell supernatants, measured by ELISA, from MDA-MB-231 and 4T1 cells post-treatment under US irradiation. (F, G) Percentage of CD80+ DCs and corresponding quantitative analysis following co-culture with pretreated 4T1 tumor cells, indicating enhanced DC maturation. (H, I) Percentage of CD80+ DCs and corresponding statistical analysis following co-culture under the

same conditions, further confirming DC activation. Treatment groups: G1: PBS; G2: NPs(3-MA/siCTL/P-18); G3: NPs(G0-C14/siNrf2/P-18); G4: NPs(3-MA/siCTL/P-18) + US; G5: NPs(G0-C14/siNrf2/P-18) + US; G6: NPs(3-MA/siNrf2/P-18) + US. Data are presented as mean ± SD (n = 3). Error bars represent standard deviation. Statistical analysis was performed using one-way ANOVA for multiple comparisons. Significance levels are indicated as *P < 0.05, **P < 0.01 and ***P < 0.001, ns, no significance.

Antitumor effect of NPs(3-MA/siNrf2/P-18) *in vivo*

Following the validation of NPs(3-MA/siNrf2/P-18)-mediated SDT in silencing Nrf2, downregulating PD-L1 expression, inhibiting protective mitophagy, and inducing intensive ICD to activate DCs, the *in vivo* antitumor efficacy was next evaluated. To determine whether these characteristics could enhance the anti-tumor immune response, NPs(3-MA/siNrf2/P-18) were intravenously administered into 4T1 orthotopic tumor-bearing mice. Due to the protective outer PEG chains [22], NPs(3-MA/siNrf2/P-18) exhibited prolonged blood circulation (**Figure 5A**) and efficient tumor accumulation (**Figure 5B-5C and Figure S10, Supporting Information**). This biodistribution profile resulted in significant silencing of Nrf2 and downregulation of PD-L1 in tumor tissue (**Figure 5D-5E**). The *in vitro* stability observed earlier correlated with the favorable *in vivo* pharmacokinetics, as the PEG shell minimized opsonization and clearance by the reticuloendothelial system. The prolonged circulation time allowed for enhanced tumor accumulation via enhanced permeability and retention effect, yielding a 2.8-fold increase in tumor-targeted delivery compared to free siRNA. To investigate the impact on immune cell activation, CD45$^+$ cells were isolated from tumor tissues following NPs(3-MA/siNrf2/P-18) administration under US irradiation. Flow cytometry analysis showed a ~10-fold increase in CD11c$^+$ MHC-II$^+$ DCs and a ~4-fold increase in CD80$^+$ CD86$^+$ DCs (**Figure 5F**), indicating enhanced DCs maturation and antigen presentation capacity within the TME. As a result, significantly higher levels of tumor-infiltrating CD8$^+$ T cells, Granzyme B$^+$ CD8$^+$ T cells, and IFN-γ$^+$ CD8$^+$ T cells were detected in tumor

tissues (**Figure 5G**), reflecting robust activation of the adaptive immune response. Furthermore, due to PD-l1 downregulation by the 3-MA-loaded nanoparticle, immune checkpoint blockade was achieved, alleviating T cell exhaustion and contributing to the elevated percentages of CD8$^+$ Granzyme B$^+$ and CD8$^+$ IFN-γ^+ T cells. The gating strategy used for flow cytometry analysis in this experiment is shown in **Figure S11** (**Supporting Information**).
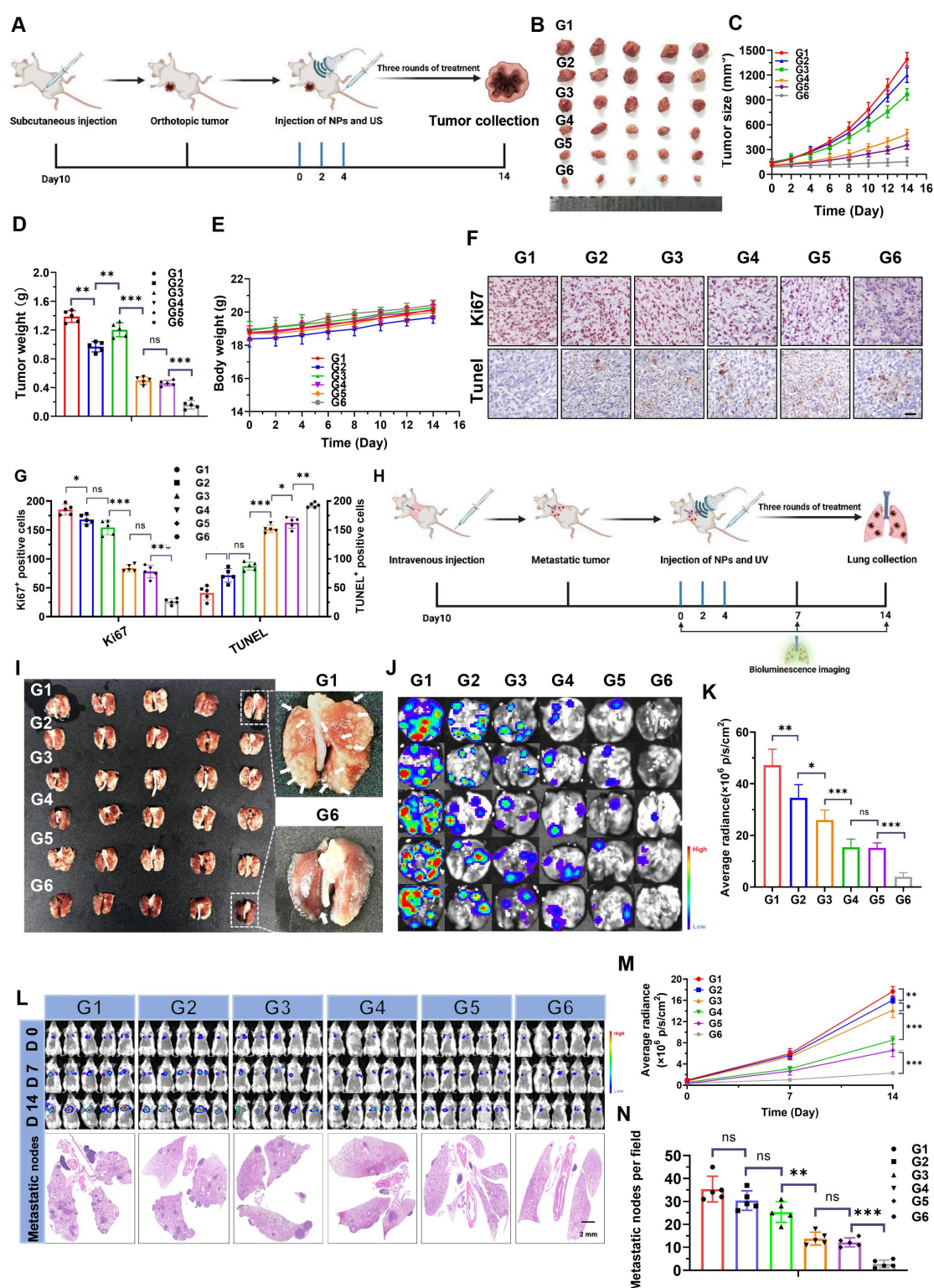
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Figure 5. *In vivo* biodistribution, gene silencing, and immune activation induced by NPs(3-MA/Cy5-siNrf2/P-18) in 4T1 orthotopic tumor-bearing mice. (A) Blood circulation profiles of naked Cy5-siNrf2 and NPs(3-MA/Cy5-siNrf2/P-18) following intravenous administration in healthy mice, indicating enhanced stability and circulation time of the nanoparticle formulation. (B) In vivo fluorescence imaging of

4T1 orthotopic tumor-bearing mice at 24 h post-injection, showing tumor-targeted accumulation of NPs(3-MA/Cy5-siNrf2/P-18). (C) *Ex vivo* fluorescence images of major organs and tumors harvested from the mice in (B), confirming preferential tumor accumulation and biodistribution. (D, E) Western blot analysis and corresponding quantification of Nrf2 and PD-L1 protein expression in tumor tissues after treatment with the indicated formulations, demonstrating gene silencing and immune checkpoint regulation. (F) Flow cytometry analysis of matured DCs (CD11c$^+$ CD80$^+$CD86$^+$) within the tumor microenvironment, indicating enhanced antigen-presenting activity. (G) Flow cytometry quantification of tumor-infiltrating CD8$^+$ T cells, Granzyme B$^+$ CD8$^+$ T cells, and IFN-γ$^+$ CD8$^+$ T cells, assessing cytotoxic T cell activation in response to treatment.. Treatment groups: G1: PBS; G2: NPs(3-MA/siCTL/P-18); G3: NPs(G0-C14/siNrf2/P-18); G4: NPs(3-MA/siCTL/P-18) + US; G5: NPs(G0-C14/siNrf2/P-18) + US; G6: NPs(3-MA/siNrf2/P-18) + US. Data are presented as mean ± SD (n = 5). Error bars represent standard deviation. Statistical analysis was performed using one-way ANOVA for multiple comparisons. Significance levels are indicated as *$P < 0.05$, **$P < 0.01$ and ***$P < 0.001$, ns, no significance.

Building on the encouraging results described above, the therapeutic efficacy of NPs(3-MA/siNrf2/P-18) *in vivo* was further evaluated in 4T1 orthotopic tumor-bearing mice under US irradiation (**Figure 6A**). As anticipated, intravenous administration of NPs(3-MA/siNrf2/P-18) combined with US irradiation significantly inhibited tumor growth without affecting overall body weight, indicating low systemic toxicity (**Figure 6B-6E**). Over the two-week treatment period, tumor volume in the PBS control group increased by approximately 13-fold, whereas tumor volume in the NPs(3-MA/siNrf2/P-18) + US group increased by only ~1.5-fold (**Figure S12, Supporting Information**). To dissect the individual contributions of mitophagy inhibition and Nrf2 silencing, NPs(3-MA/siCTL/P-18) and NPs(G0-C14/siNrf2/P-18) were administered with US irradiation. Each treatment led to ROS accumulation and moderate tumor growth inhibition over 14 days, underscoring the necessity of

combinational therapy to achieve maximal therapeutic efficacy. This synergistic effect was further proven by histological analysis: TUNEL staining showed increased apoptosis, and Ki67 staining revealed reduced proliferation in tumor tissues treated with NPs(3-MA/siNrf2/P-18) (**Figure 6F-6G**). To evaluate the anti-metastatic potential of the NP, a luciferase-expressing 4T1 (Luc-4T1) lung metastasis model was established (**Figure 6H**) [8]. Similar to the inhibition observed in orthotopic tumors, NPs(3-MA/siNrf2/P-18) treatment significantly suppressed lung metastases compared to all other treatment groups. This was evidenced by a lower number of metastatic nodules (**Figure 6I**), reduced bioluminescence signal from lung tissues (**Figure 6J-6K**), and diminished whole-body bioluminescence intensity (**Figure 6L, Figure 6M** and **Figure S13, Supporting Information**). Hematoxylin and eosin (H&E) staining further confirmed a marked reduction in metastatic nodules in lung sections (**Figures 6L** and **Figure 6N**). Notably, no apparent histological abnormalities were observed in major organs of mice treated with the NPs(3-MA/siNrf2/P-18) under US irradiation (**Figure S14, Supporting Information**), and no obvious fluctuations were detected in hematological parameters (**Figure S15, Supporting Information**) or in liver and kidney function tests (**Figure S16, Supporting Information**), confirming the favorable *in vivo* biosafety profile of NPs(3-MA/siNrf2/P-18) under US irradiation. In summary, the reduction-responsive RNAi NP demonstrated excellent anti-tumor and anti-metastatic efficacy *in vivo*, along with good biosafety and biocompatibility, supporting its potential for translational cancer therapy.

764

Figure 6. *In vivo* antitumor and antimetastatic efficacy of NPs(3-MA/siNrf2/P-18) under US irradiation. (A) Schematic illustration of the experimental timeline showing tumor inoculation, treatment schedule, and formulation details in 4T1 orthotopic tumor-bearing mice. Each treatment contained 5 mg/kg P-18, 6 mg/kg 3-MA, and/or 1

769 nM siNrf2 per mouse. (B) Representative images of excised primary tumors at the

770 study endpoint. (C) Tumor growth curves for each treatment group. (D) Tumor

771 weights measured at the endpoint, reflecting therapeutic efficacy. (E) Body weight

772 changes of mice over the course of treatment, indicating systemic tolerability. (F)

773 Representative IHC staining images of Ki67and TUNEL in tumor sections. (G)

774 Quantification of Ki67$^+$ and TUNEL$^+$ cells per high-power field. (H) Schematic of the

775 experimental setup for the lung metastasis model using luciferase-expressing 4T1

776 cells and treatment allocation. (I) Photographs of excised lungs showing visible

777 metastatic nodules. (J) Representative bioluminescence imaging of lung metastases.

778 (K) Quantitative analysis of lung bioluminescence intensity, indicating metastatic

779 burden. (L) Whole-body bioluminescence imaging at days 0, 7, and 14 post-treatment,

780 along with H&E staining of lung tissue sections to assess metastatic infiltration. (M)

781 Quantification of whole-body bioluminescence signal intensity over time. (N)

782 Quantification of metastatic nodules in H&E-stained lung sections. Treatment groups:

783 G1: PBS; G2: NPs(3-MA/siCTL/P-18); G3: NPs(G0-C14/siNrf2/P-18); G4:

784 NPs(3-MA/siCTL/P-18) + US; G5: NPs(G0-C14/siNrf2/P-18) + US; G6:

785 NPs(3-MA/siNrf2/P-18) + US. Data are presented as mean ± SD (n = 5). Error bars

786 represent standard deviation. Statistical analysis was performed using one-way

787 ANOVA for multiple comparisons. Significance levels are indicated as *P < 0.05, **P

788 < 0.01 and ***P < 0.001, ns, no significant,

789 **Conclusion**

790 This study presents an innovative approach that concurrently targets the mitophagy

791 and Nrf2 pathways to enhance SDT by amplifying intracellular ROS generation

792 within tumor cells. This strategy significantly curtails both progression and metastasis

793 in breast cancer. The ROS-induced oxidative stress promotes apoptosis and triggers

794 extensive ICD, leading to the release of DAMPs that promote DCs maturation and

795 antigen presentation to CD8$^+$ T cells. In addition, 3-MA downregulates PD-L1

796 expression via an NF-κB-dependent pathway, thereby mitigating T cell exhaustion

797 and bolstering CD8$^+$ T cell-mediated antitumor immunity. The reduction-responsive

798 RNAi NP developed in this study proves to be a potent tool for augmenting

799 therapeutic outcomes in breast cancer. Beyond SDT, the ROS amplification capability

800 of this NP may also enhance other ROS-dependent modalities, including

801 chemodynamic therapy, photodynamic therapy, radiotherapy, and chemotherapy.

802 Furthermore, it synergizes effectively with immune checkpoint blockade. Collectively,

803 the SDT-mediated, reduction-responsive RNAi NP represents a novel, versatile, and

804 effective strategy for advancing multimodal cancer treatment.

805

806 **Conflict of interest**

807 The authors declare no conflict of interest.

808

820

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