



Supporting Information

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NIR-Triggered Phototherapy and Immunotherapy via an Antigen-Capturing Nanoplatfrom for Metastatic Cancer Treatment

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Supplementary Figures

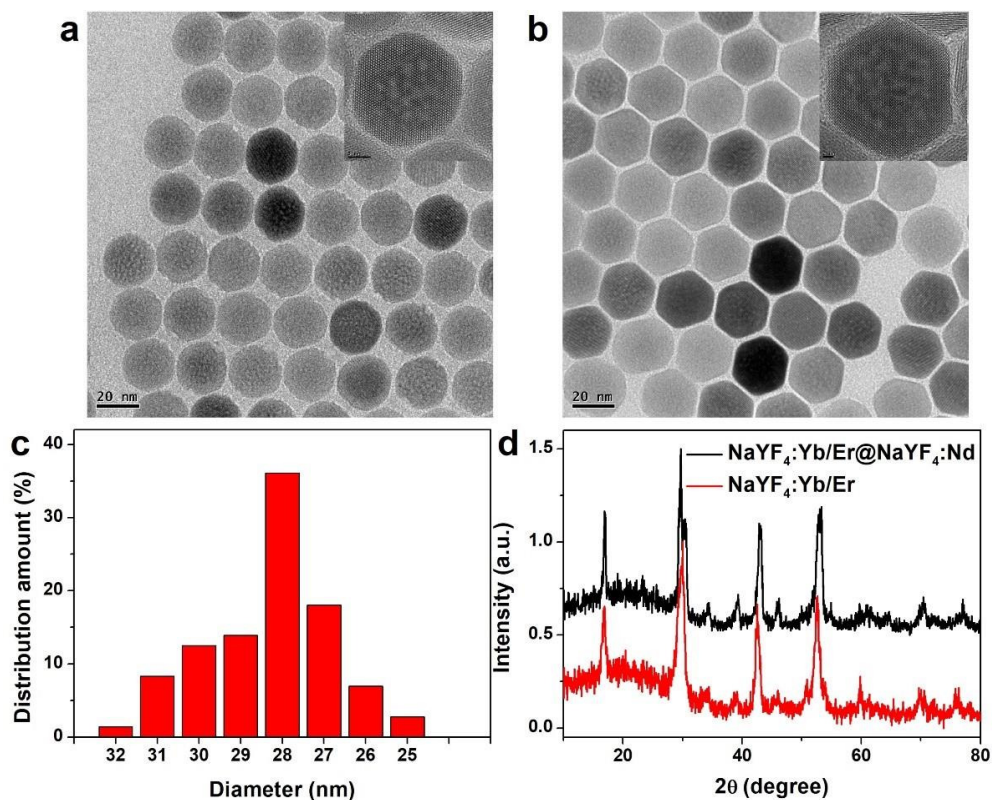


Figure S1. Transmission electron microscope (TEM) and High Resolution Transmission Electron Microscope (HRTEM) images of a) oleate-capped NaYF₄:Yb/Er UCNPs b) oleate-capped NaYF₄:Yb/Er@NaYF₄:Nd UCNPs. c) Size distribution of oleate-capped NaYF₄:Yb,Er/NaYF₄:Nd UCNPs as determined from 200 particles in the TEM image. d) Powder XRD pattern of NaYF₄:Yb/Er UCNPs and NaYF₄:Yb/Er@NaYF₄:Nd UCNPs.

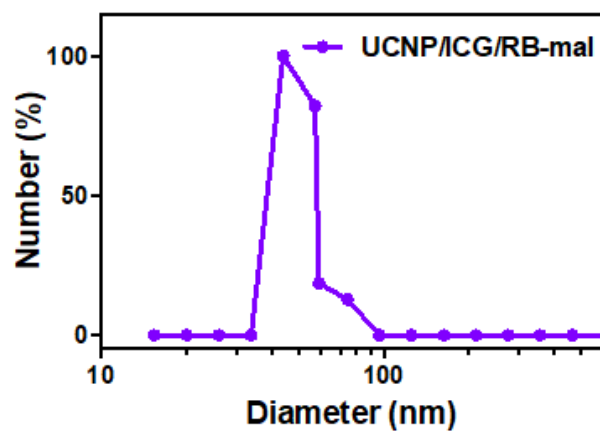


Figure S2. Hydrodynamic diameter distribution of UCNP/ICG/RB-mal. The hydrodynamic diameters of UCNP/ICG/RB-mal were determined to be 41 ± 1 nm.

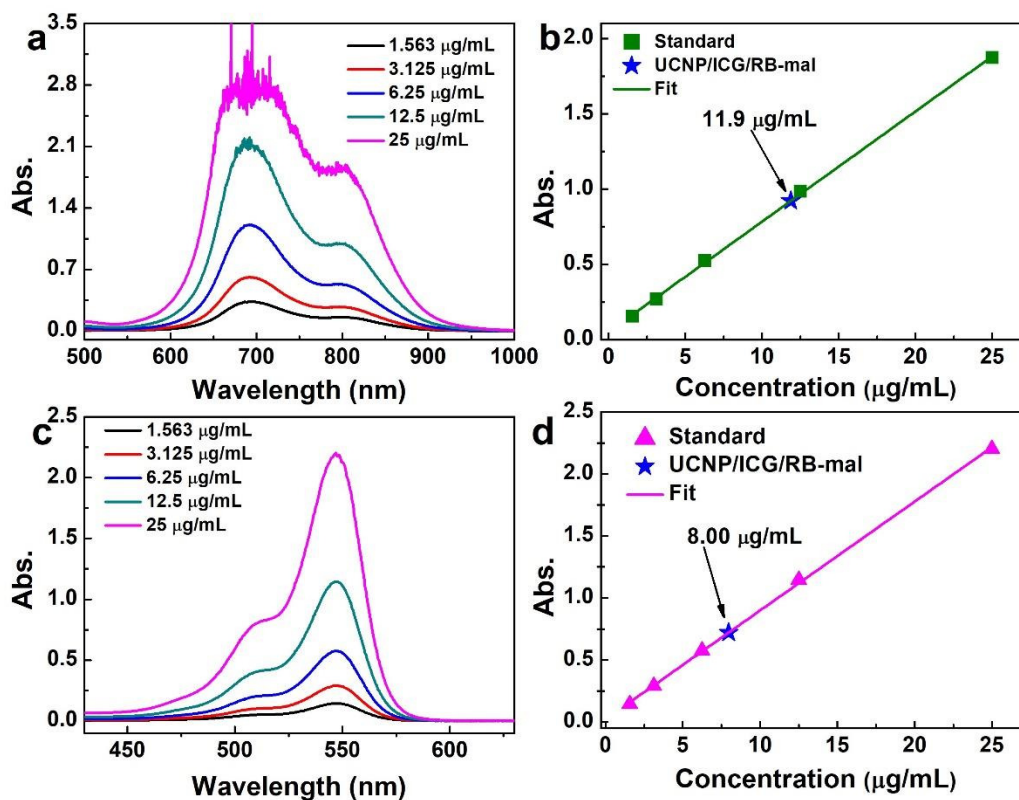


Figure S3. Quantitative analysis of indocyanine green (ICG) or rose bengal (RB) loaded in UCNP/ICG/RB-mal. a) Absorption spectra of ICG solution. b) The standard curve of absorption for ICG. The loading amount of ICG was determined to be 1.19% (w / w) in UCNP/ICG/RB-mal. c) Absorption spectra of RB solution. d) The standard curve of absorption for RB. The loading amount of RB was determined to be 0.8% (w / w) in UCNP/ICG/RB-mal.

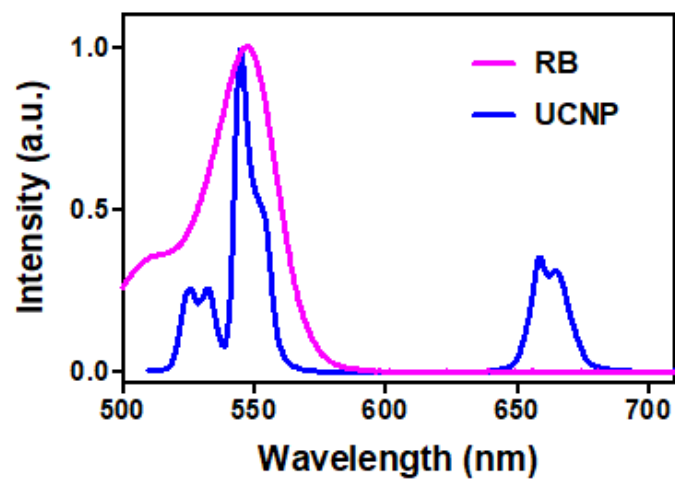


Figure S4. Absorption spectra of the RB solution and upconversion luminescence (UCL) spectra of oleate-capped NaYF₄:Yb/Er@NaYF₄:Nd nanoparticles, respectively.

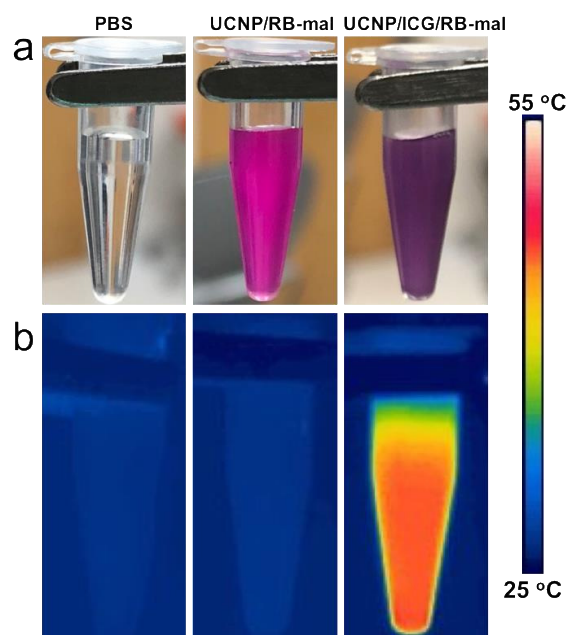


Figure S5. Infrared thermal images of PBS, UCNP/RB-mal, and UCNP/ICG/RB-mal solutions upon a 805-nm laser irradiation for 4 min. (0.75 W cm⁻²).

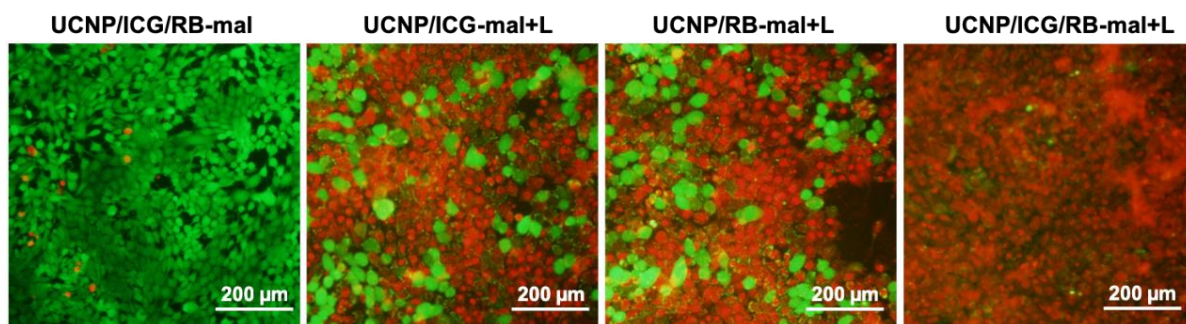


Figure S6. Fluorescence images of 4T1 cells after 4 h incubation with UCNP/ICG-mal, UCNP/RB-mal, and UCNP/ICG/RB-mal exposed to a 805-nm laser for 5 min. (0.75 W cm^{-2}) The cells were con-stained by calcein AM (green, live cells) and propidium iodide (red, dead cells) before imaging. Scale bar: 200 μm .

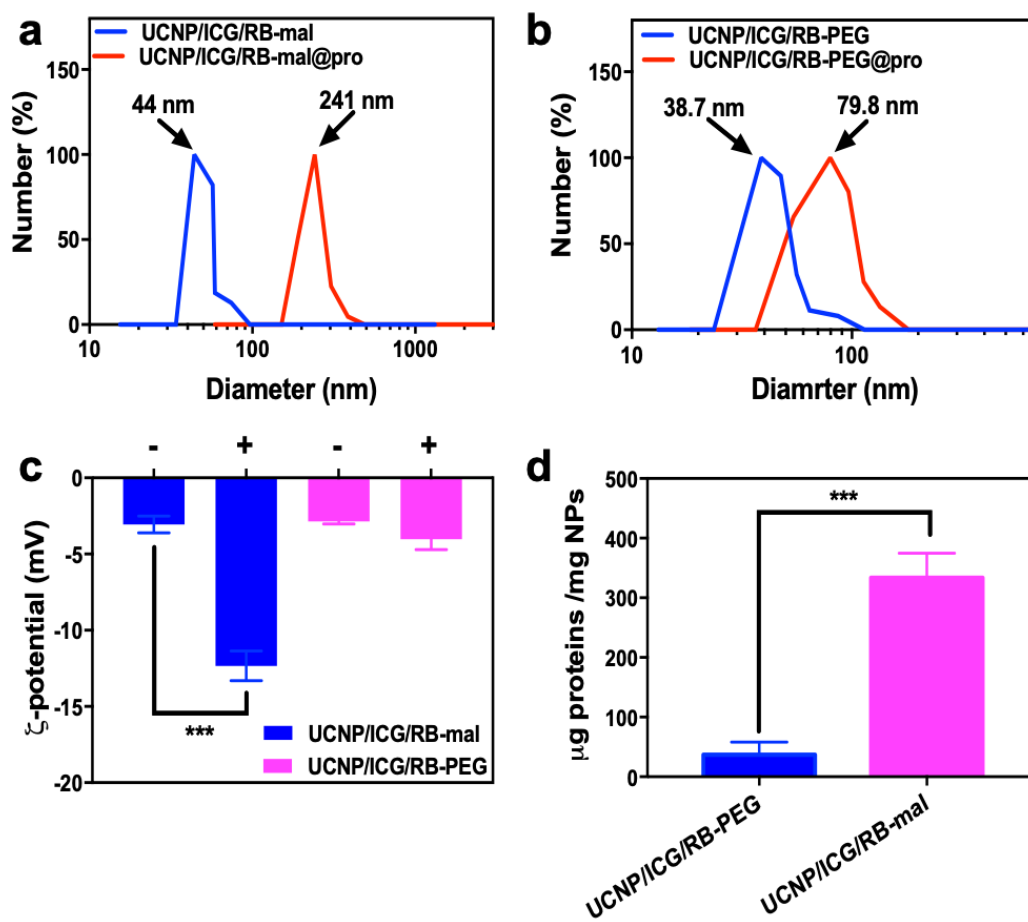


Figure S7. The size of a) UCNP/ICG/RB-mal and b) UCNP/ICG/RB-PEG before and after antigen capture. c) The zeta potential of UCNP/ICG/RB-mal or UCNP/ICG/RB-PEG before and after antigen capture. (***) $P < 0.001$ d) Quantification of protein captured by nanoparticles. (***) $P < 0.001$). Data are expressed as means \pm SD ($n = 4$).

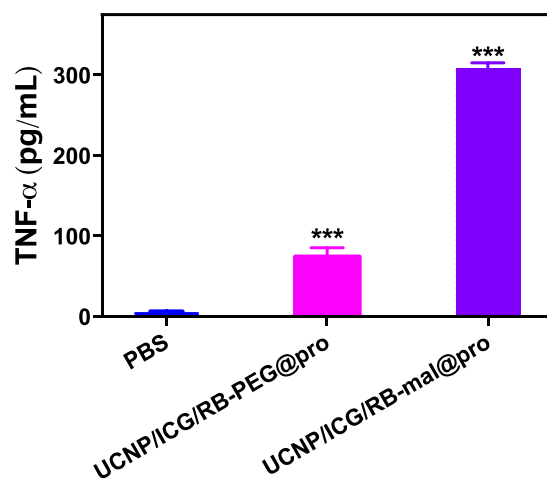


Figure S8. Secretion of TNF- α in BMDC suspensions measured by ELISA (^{***}P < 0.001 vs PBS). Data are expressed as means \pm SD (n = 4).

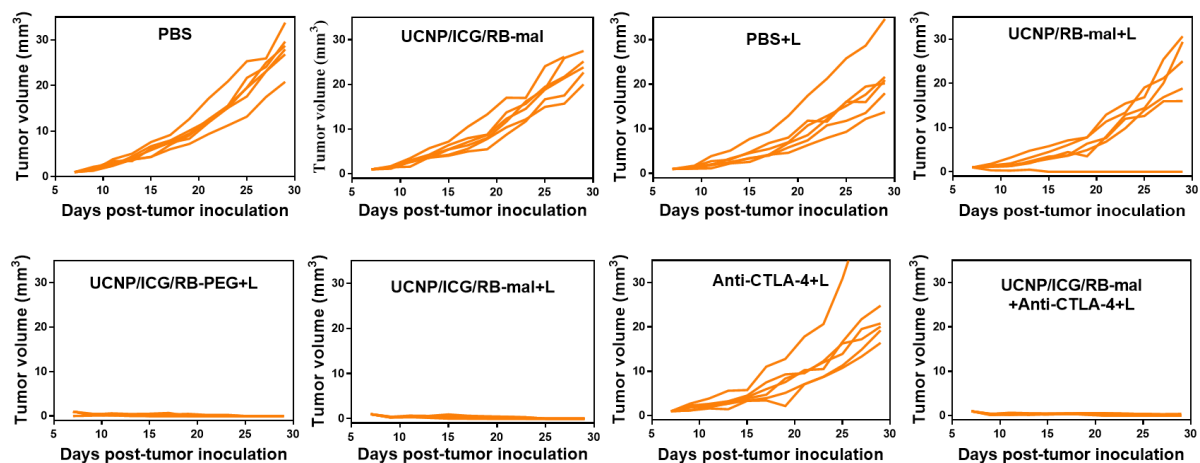


Figure S9. Growth curves of tumors in individual mice after various treatments.

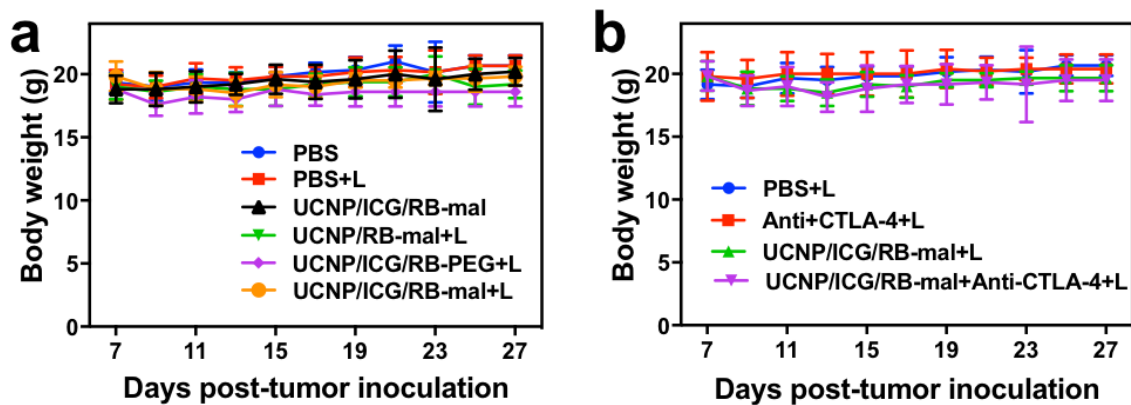


Figure S10. Average body weight of orthotopic 4T1 tumor-bearing mice of different groups. Data are expressed as means \pm SD (n = 6).

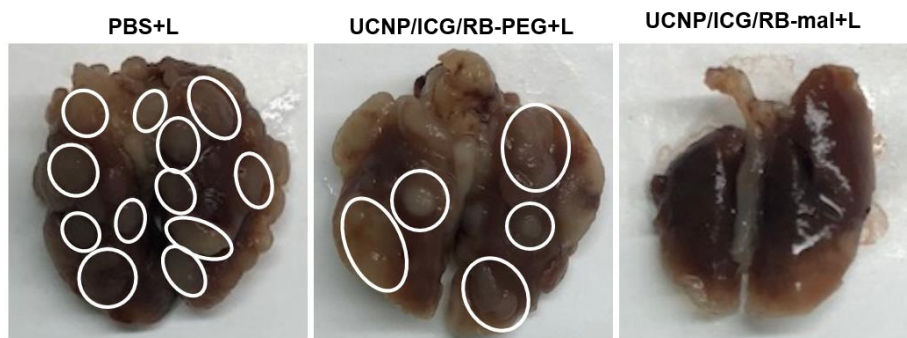


Figure S11. Photographs of metastatic nodules in lungs of mice in different groups 25 days after treatment.

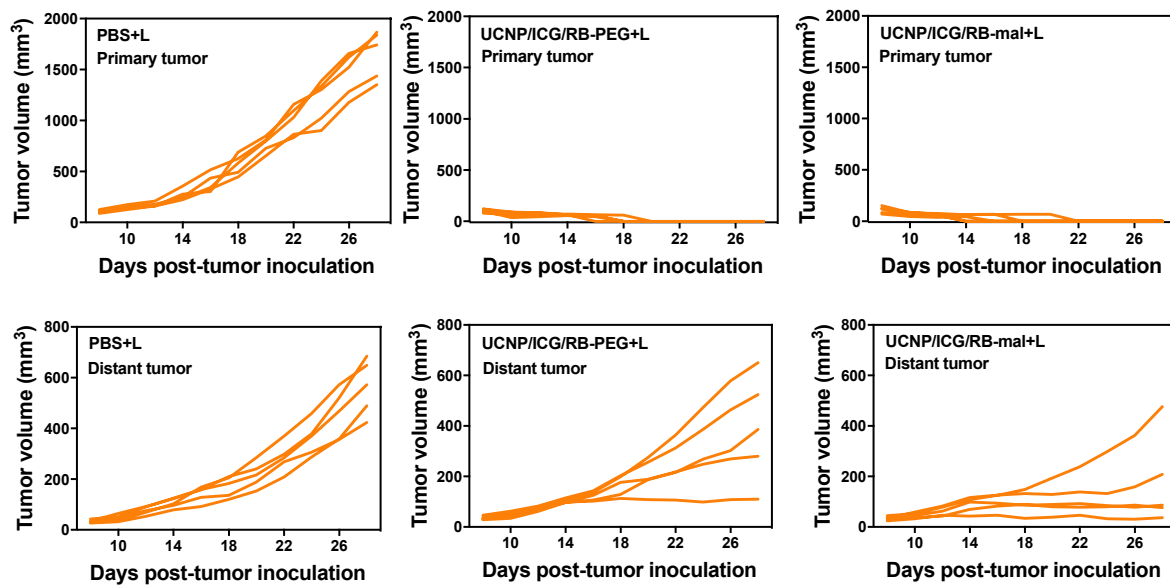


Figure S12. Growth curves of primary and distant tumors in individual mice after various treatments.

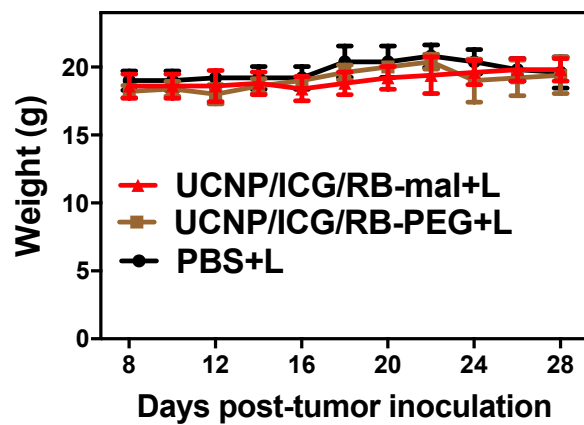


Figure S13. Average body weight of bilateral 4T1 tumor-bearing mice of different groups. Data are expressed as means \pm SD (n = 5).

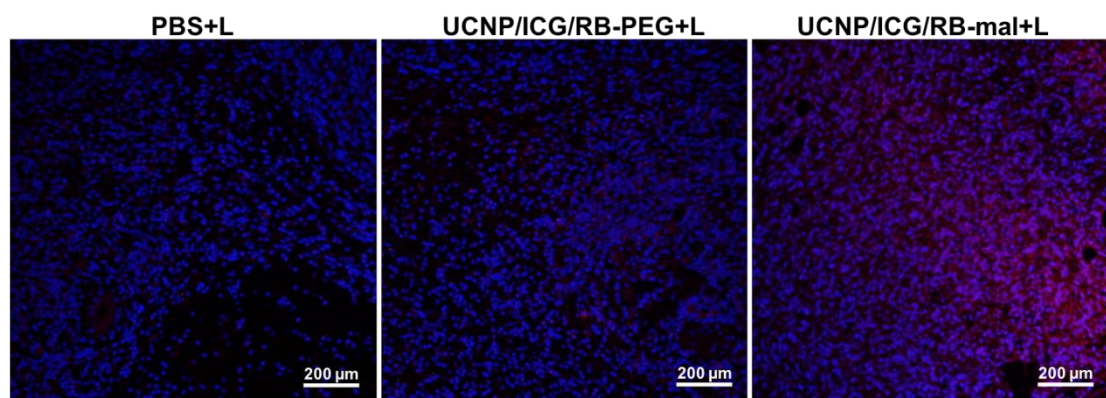


Figure S14. Immunohistochemistry staining of CD11c⁺ dendritic cells (red) in 4T1 primary tumors 1 day after the treatments. Scale bar = 200 μM.

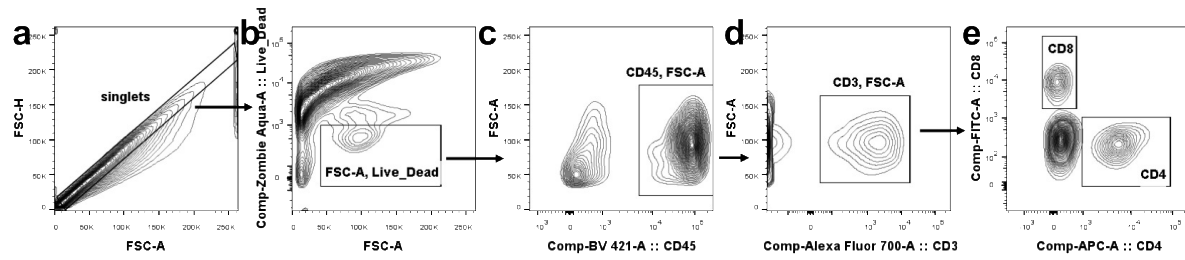


Figure S15. Gating strategy determine frequencies of T cells from distant tumors. The gating strategy was performed based on exclusion of doublets by FSC-A and FSC-H a), exclusion of dead cells b), selection of CD45⁺ c), selection of CD3⁺ d), and further staining using CD8 e) and CD4 e) with appropriate fluorescent dyes to select CD8⁺ or CD4⁺ T cells, respectively.

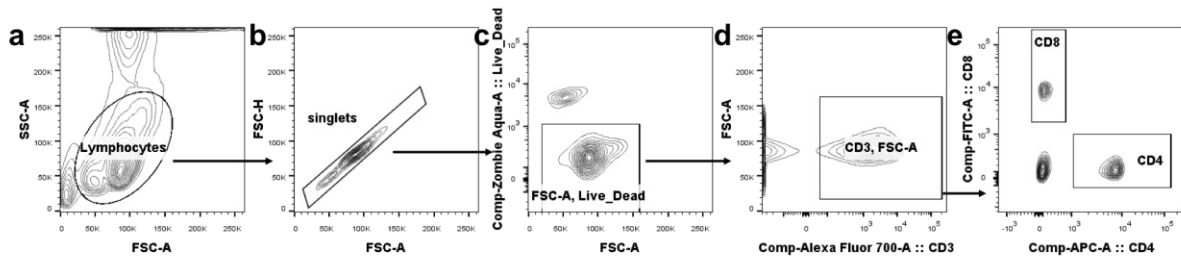


Figure S16. Gating strategy determine frequencies of T cells from spleens. The gating strategy was performed based on the right justification of the first gate a), exclusion of doublets by FSC-A and FSC-H b), exclusion of dead cells c), selection of CD3⁺ d), and further staining using CD8 e) and CD4 e) with appropriate fluorescent dyes to select CD8⁺ or CD4⁺ T cells, respectively.

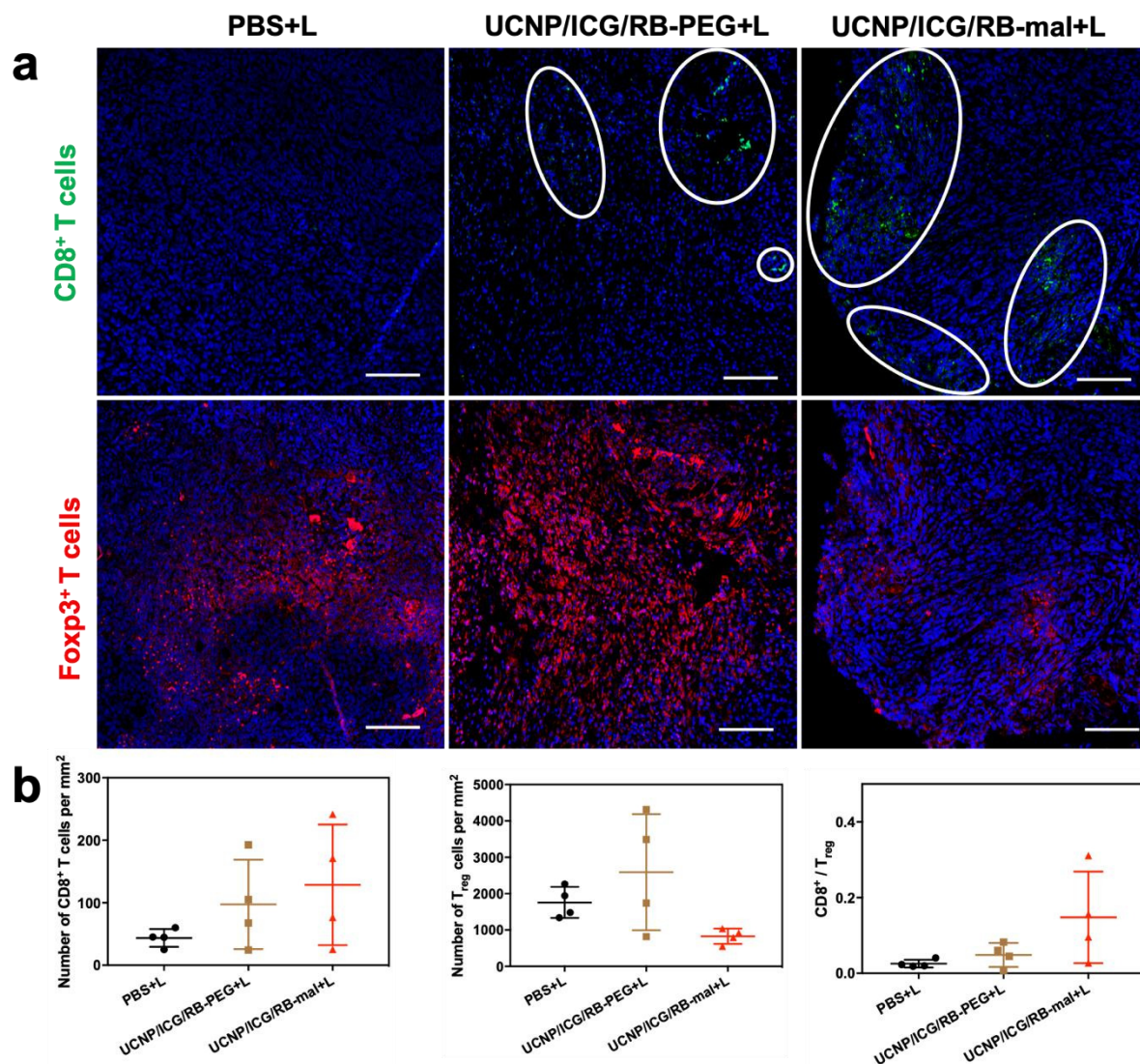


Figure S17. T cells immunofluorescence assay. a) Representative images of tumors after staining of CD8⁺ T cells (Green) and FoXp3⁺ T cells (T_{reg}) (Red) in 4T1 untreated second tumors 7 days after the treatments of the primary tumors. Scale bar = 200 μ M. b) The densities of CD8⁺ T cells and T_{reg} cells in the tumors. Data are expressed as means \pm SD (n = 4).

Table S1. The capture of TDPAs by UCNP/ICG/RB-mal.

	Gene	Description	iBAQ
1	Hist2h2ab	Histone H2A type 2-B, PE=1 SV=3	145001.046
2	Hist2h2ac	Histone H2A type 2-C, PE=1 SV=3	25180.155
3	Hist1h4a,	Histone H4, PE=1 SV=2	668804.651
4	Hist1h2bf	Histone H2B type 1-F/J/L, PE=1 SV=2	1220846.62
5	Hmgb1	High mobility group protein, PE=1 SV=4	875.588785
6	Hspd1	60 kDa heat shock protein, PE=1 SV=1	82812.8744
7	Hsph1	Heat shock protein 105 kDa, PE=1 SV=2	5658.95921
8	Hsp90ab1	Heat shock protein HSP 90-beta, PE=1 SV=3	21224.0829
9	Hspa1b	Heat shock 70 kDa protein 1B, PE=1 SV=3	6718.14642
10	Hspa8	Heat shock cognate 71 kDa protein, PE=1 SV=1	109807.119
11	Hspa4l	Heat shock 70 kDa protein 4L, PE=1 SV=2	349.673031
12	Hsp90aa1	Heat shock protein HSP 90-alpha, PE=1 SV=4	15045.1419
13	Hsp90ab1	Heat shock protein HSP 90-beta, PE=1 SV=3	21224.0829
14	Hspd1	60 kDa heat shock protein, PE=1 SV=1	82812.8744
15	Trap1	Heat shock protein 75 kDa, PE=1 SV=1	43148.83
16	Hspa4	Heat shock 70 kDa protein 4, PE=1 SV=1	12998.5256
17	Hspd1	60 kDa heat shock protein, PE=1 SV=1	2155.54439
18	Mthfd1	C-1-tetrahydrofolate synthase, PE=1 SV=4	36050.4115
19	Mthfd11	Monofunctional C1-tetrahydrofolate synthase, PE=1 SV=2	2155.54439
20	Mthfd1	C-1-tetrahydrofolate synthase, PE=1 SV=4	6686.73136
21	Actn4	Alpha-actinin-4, PE=1 SV=1	26667.0449