

## Supporting information

# **Nanosensitizer-mediated augmentation of sonodynamic therapy efficacy and antitumor immunity**

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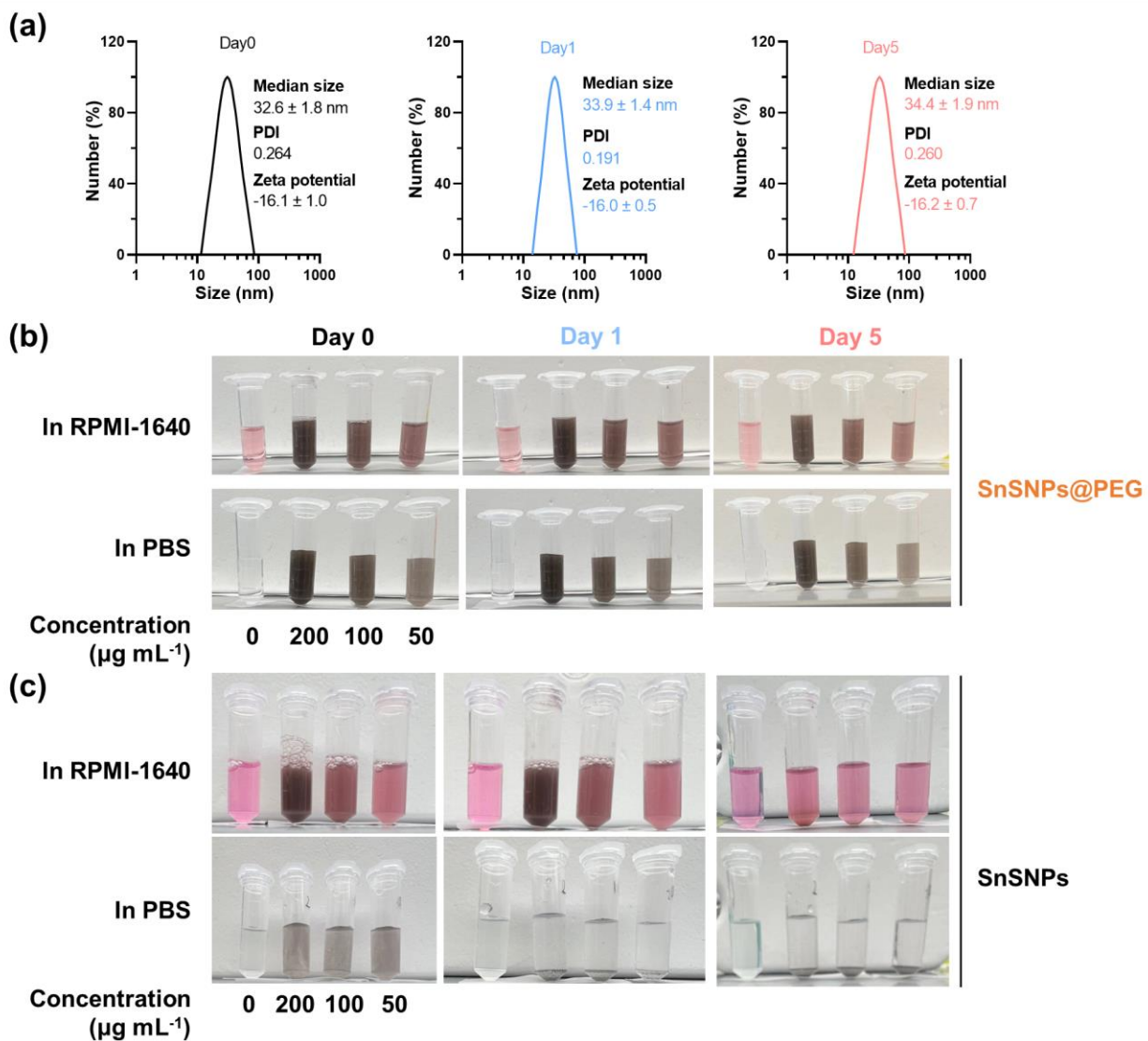
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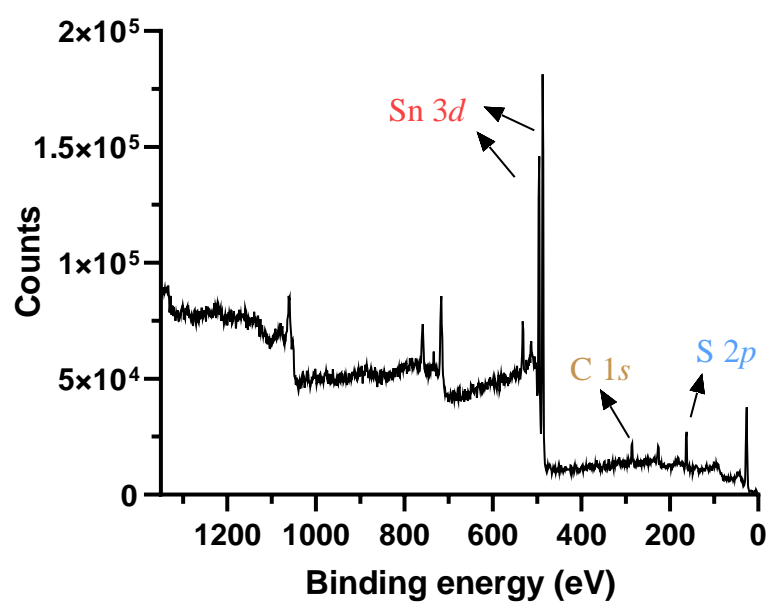
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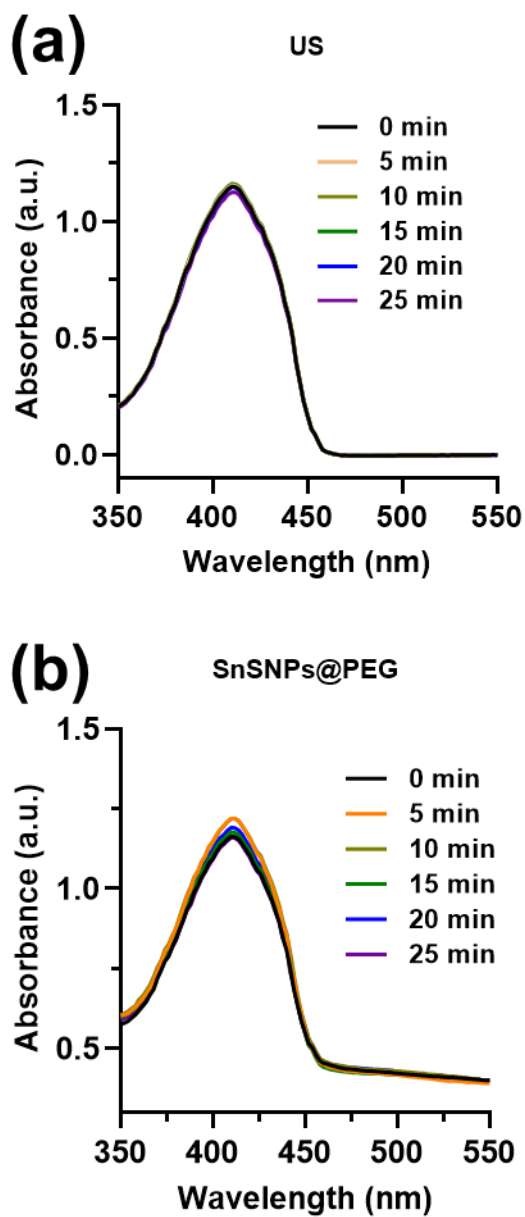
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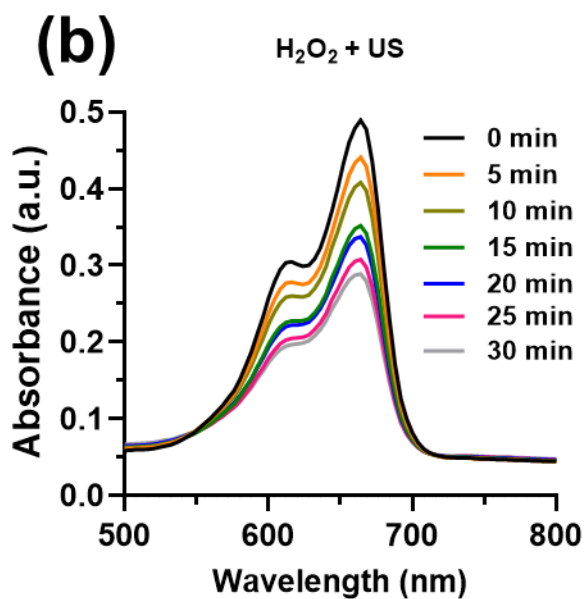
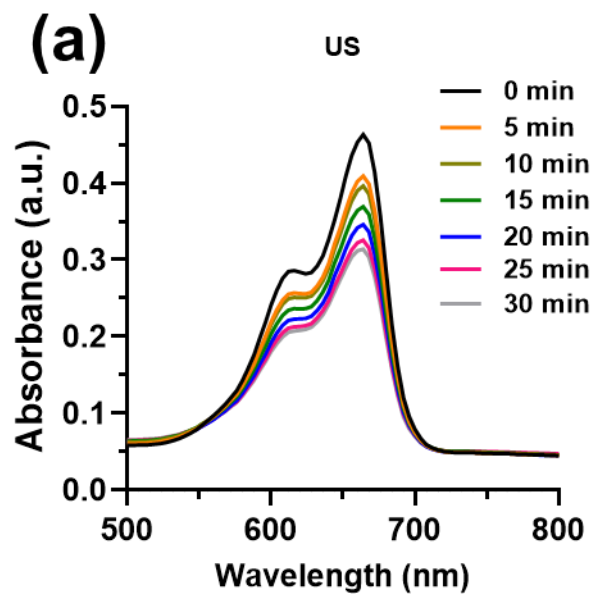
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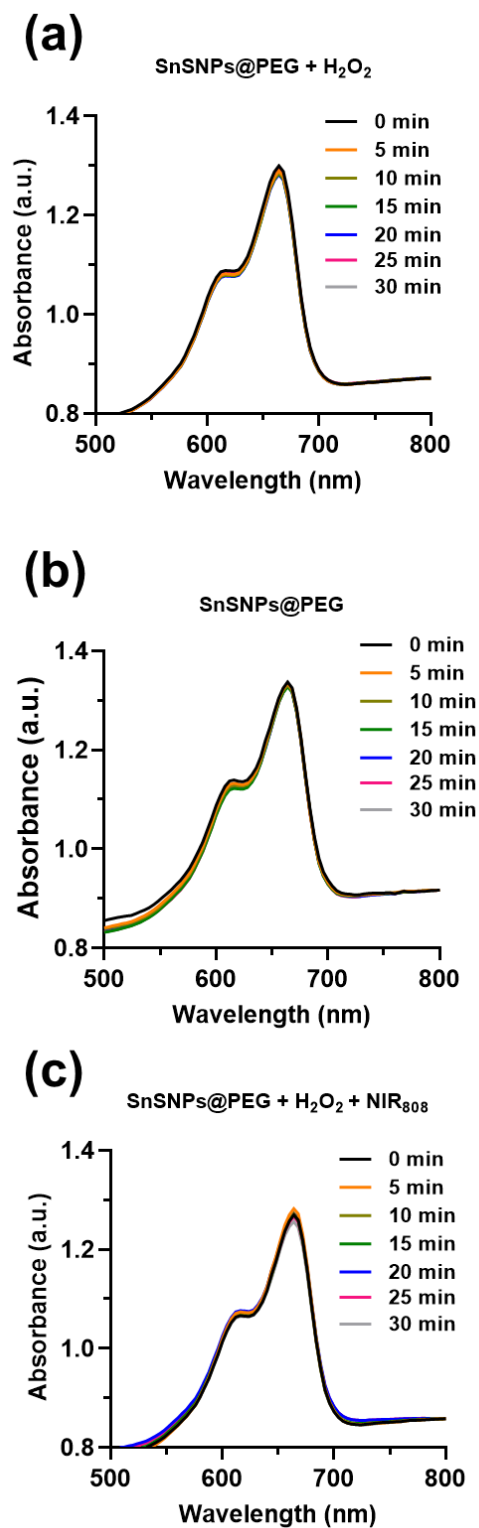
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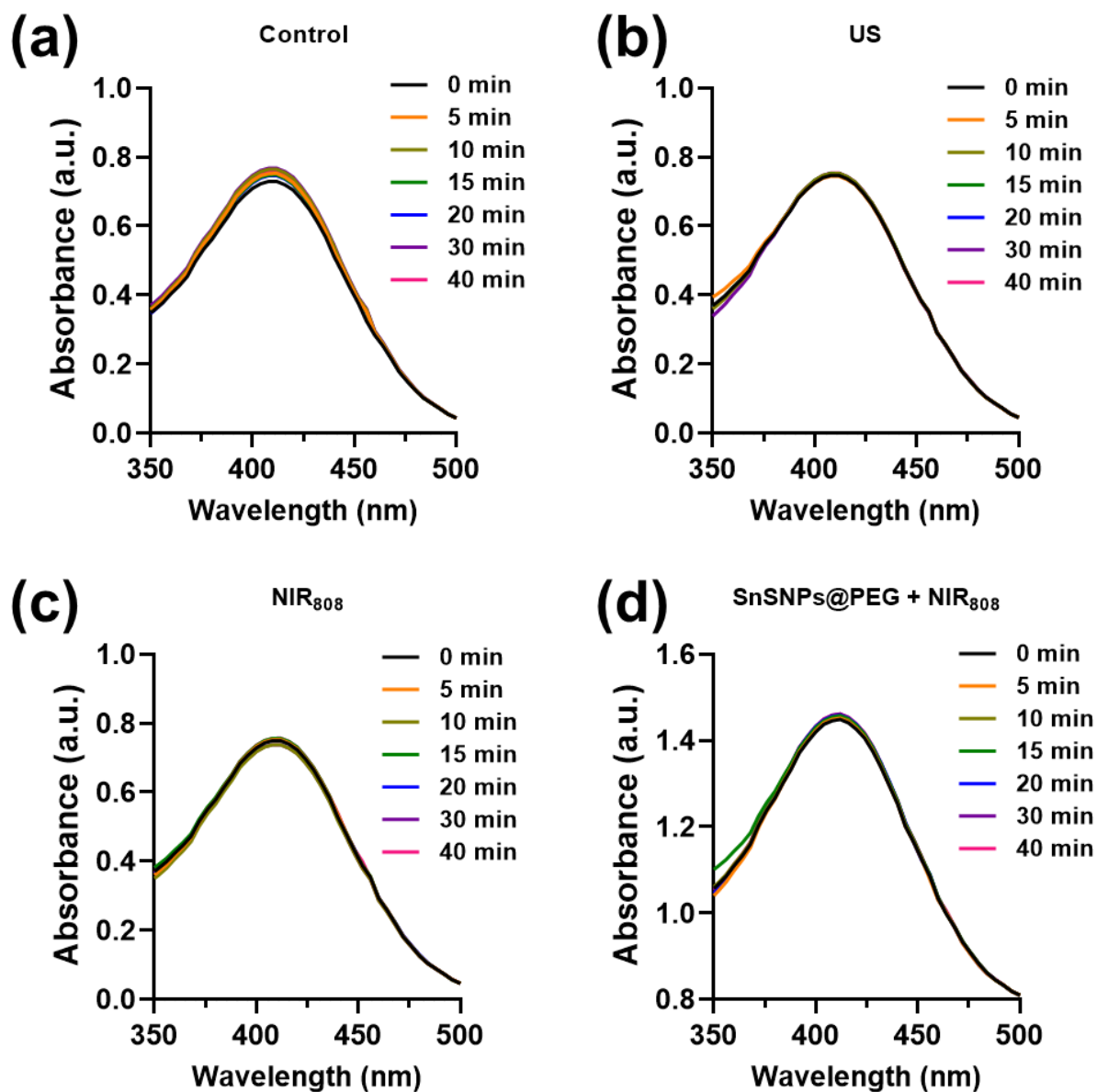


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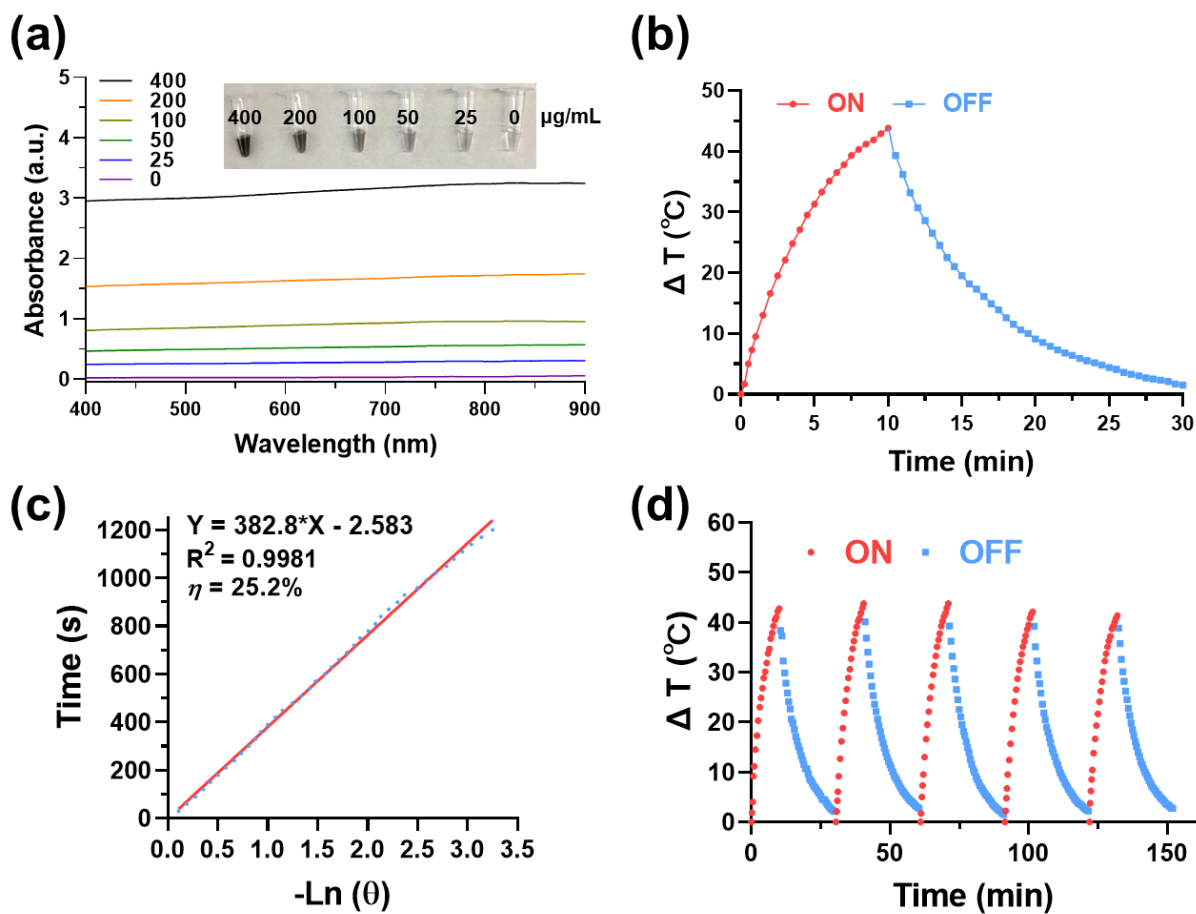


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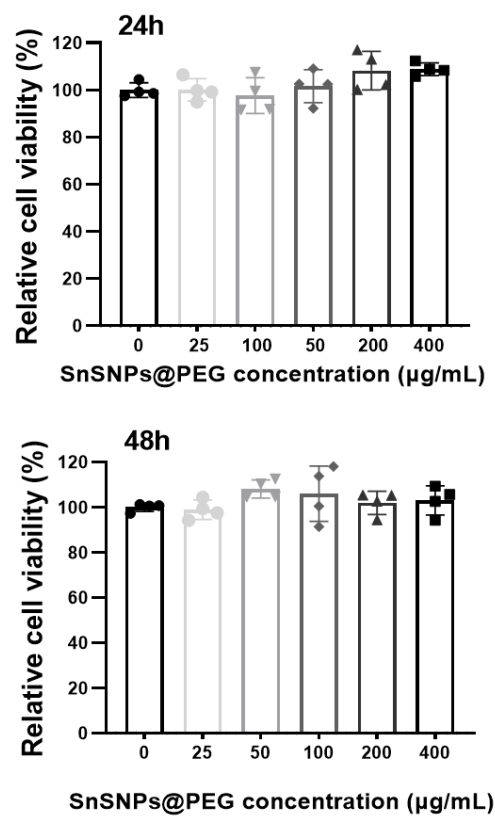




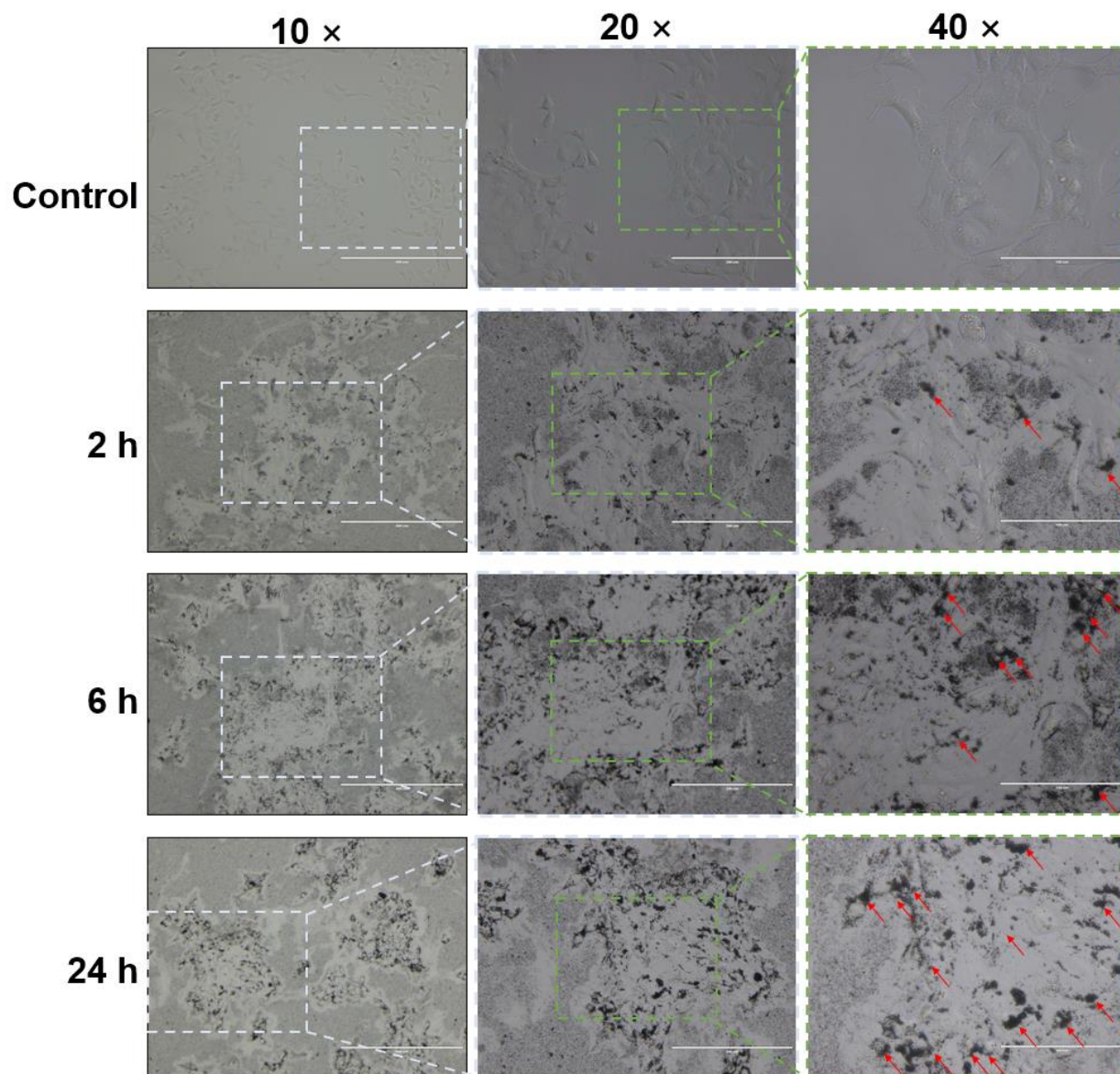
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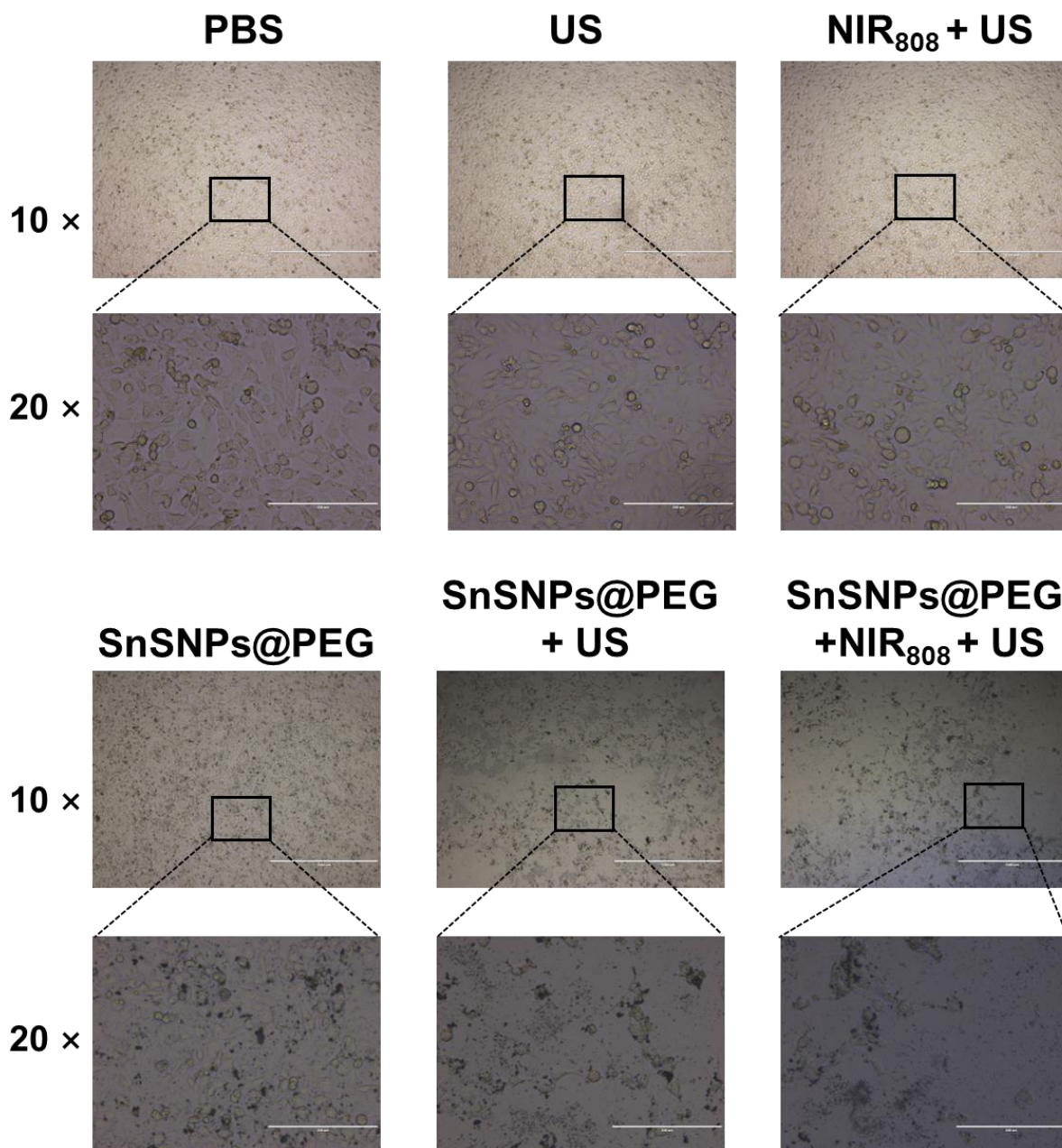
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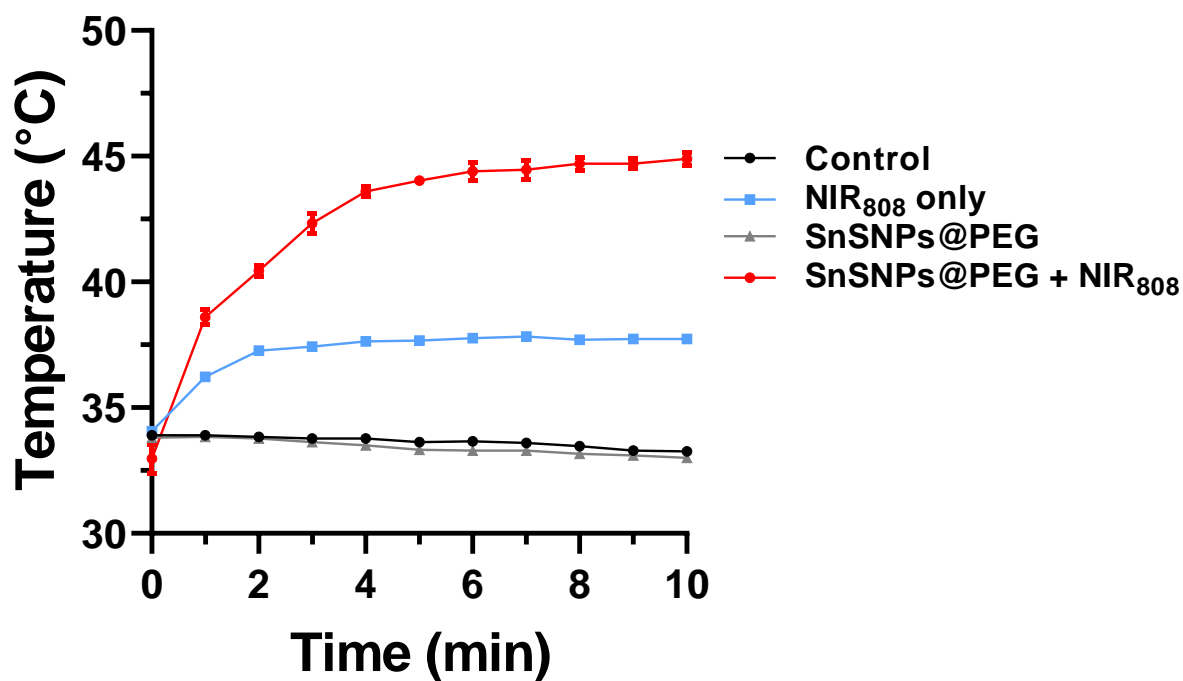
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**Figure S9.** Bright-field microscope images showing time-dependent cellular uptake of SnSNPs@PEG by 4T1 cells. The substantial uptake was highlighted by red arrows. No cytotoxic effect was observed on 4T1 cells given that the substantial cellular uptake of SnSNPs@PEG was observed. Scale bar: 10 $\times$  = 400  $\mu$ m, 20 $\times$  = 200  $\mu$ m, 40 $\times$  = 100  $\mu$ m.

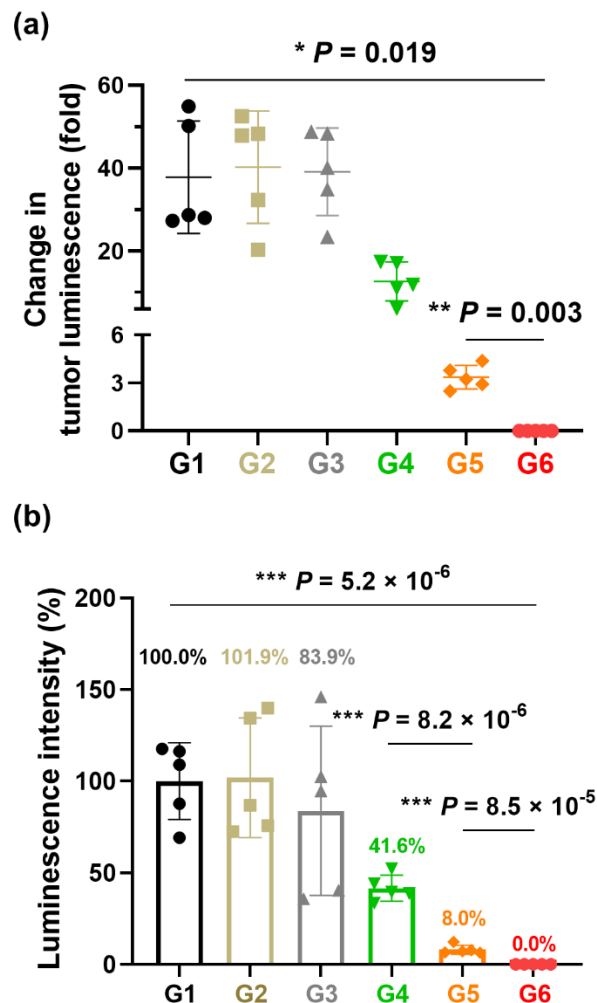


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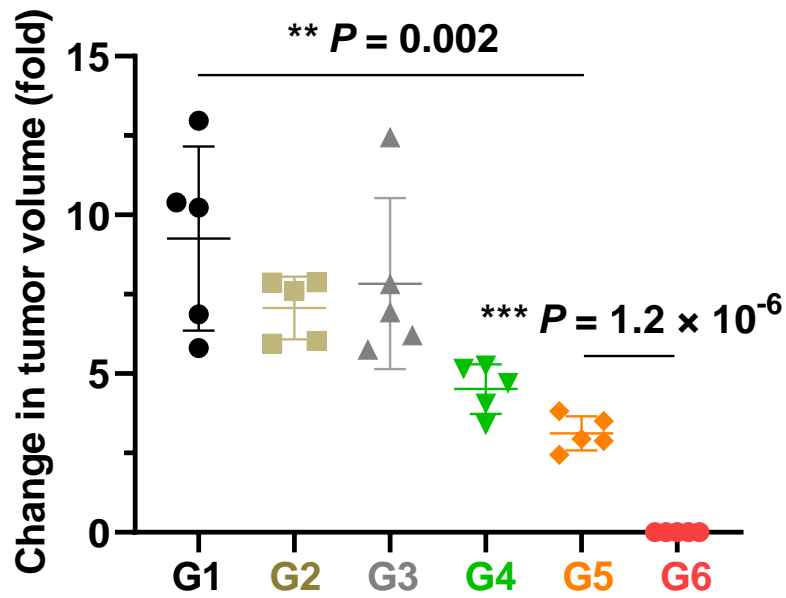


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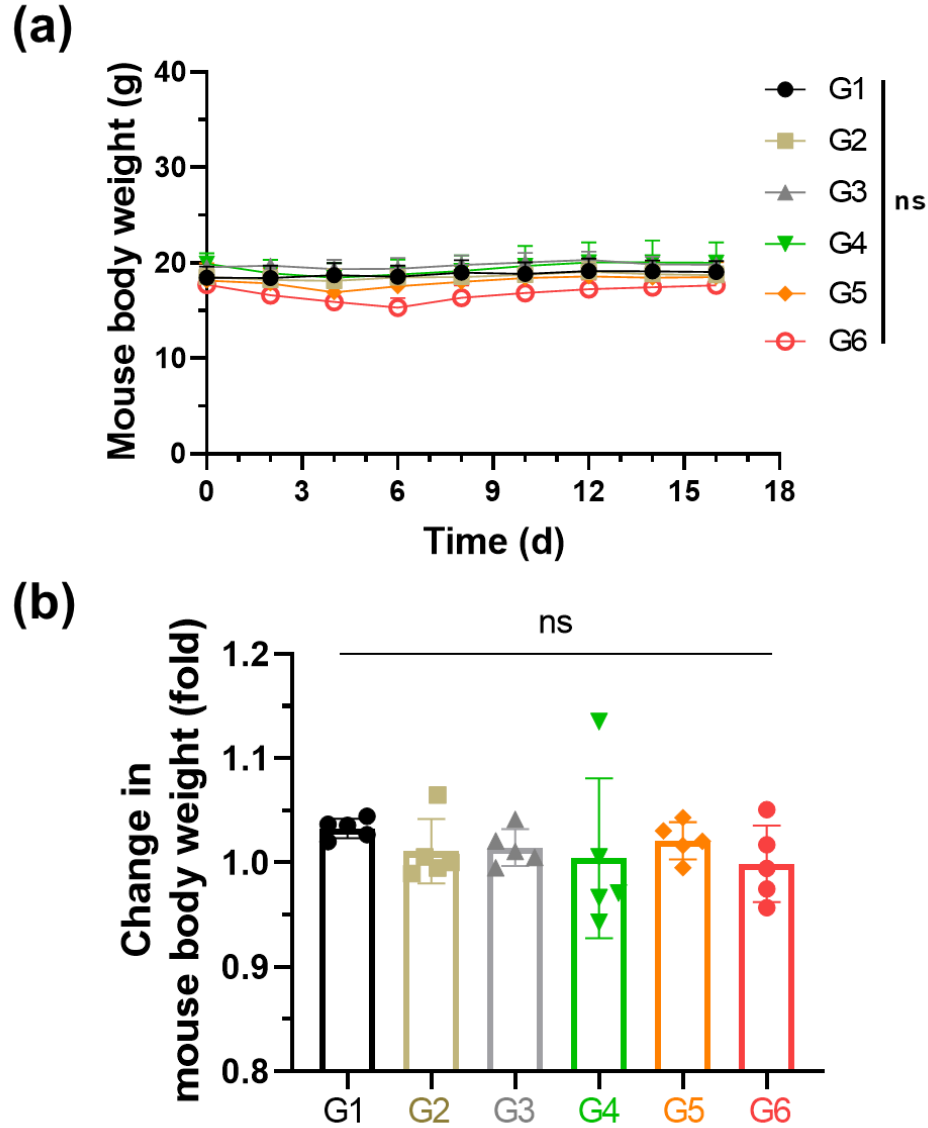


**Figure S12.** Comparison of luminescence intensity of 4T1 tumor areas after various treatments. (a) Fold-change of tumor luminescence intensity. The intensity change was calculated as luminescence intensity on Day 16/ luminescence intensity on Day 0 (before treatment). (b) Percentage of tumor luminescence intensity. The percentage was calculated as follows: luminescence intensity at Day 16 of each tumor-bearing mice in group/ average luminescence intensity of mice in G1 at Day 16. Statistical analysis between the two groups was performed using two-sided student's *t*-test.  $** P < 0.01$ ,  $*** P < 0.001$ . Groups are as follows: control (G1), SnSNPs@PEG (G2), NIR + US (G3), SnSNPs@PEG + US (G4), SnSNPs@PEG + NIR + US (G5), SnSNPs@PEG + NIR + US  $\times 2$  (G6), (n = 5).

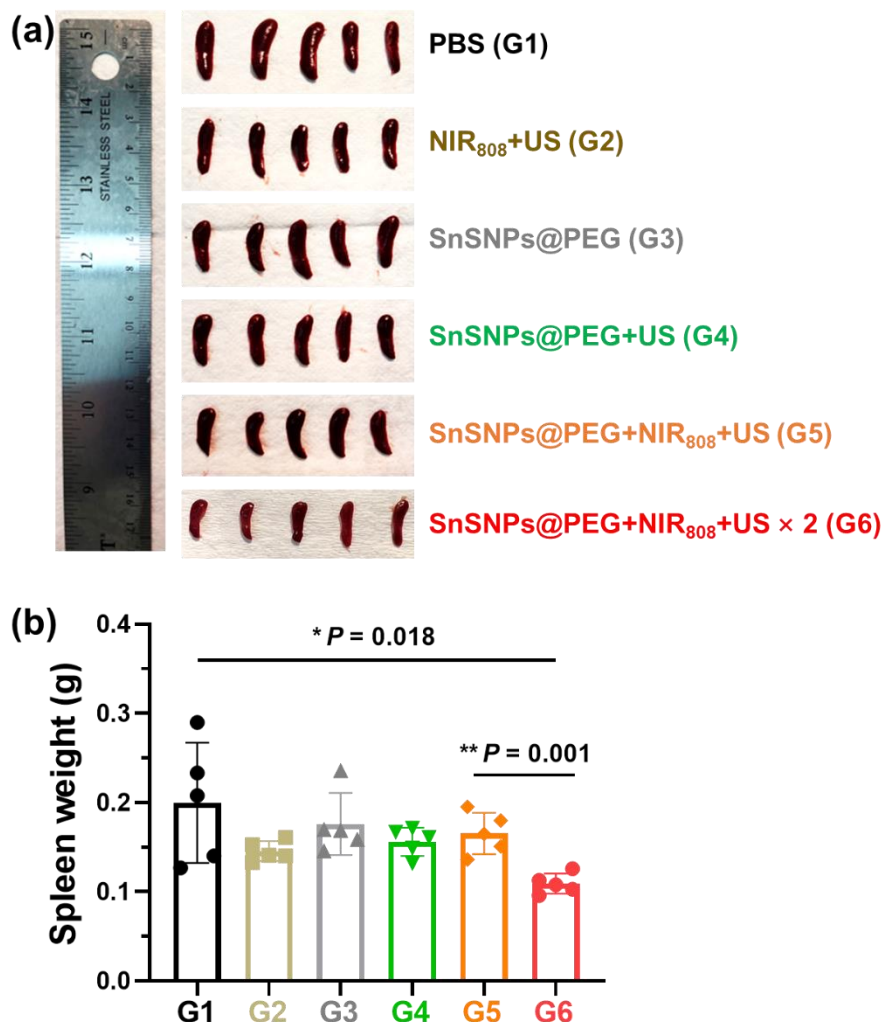


**Figure S13.** Change in the 4T1 tumor volume after various treatments. The volume change was calculated as follows: tumor volume at Day 16/ tumor volume at Day 0 (before treatment). Data are presented as mean  $\pm$  SD (n = 5). Statistical analysis between two groups was performed using two-sided student's *t*-test. \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ . Groups are as follows: control (G1), SnSNPs@PEG (G2), NIR + US (G3), SnSNPs@PEG + US (G4), SnSNPs@PEG + NIR + US (G5), SnSNPs@PEG + NIR + US  $\times$  2 (G6), (n = 5).

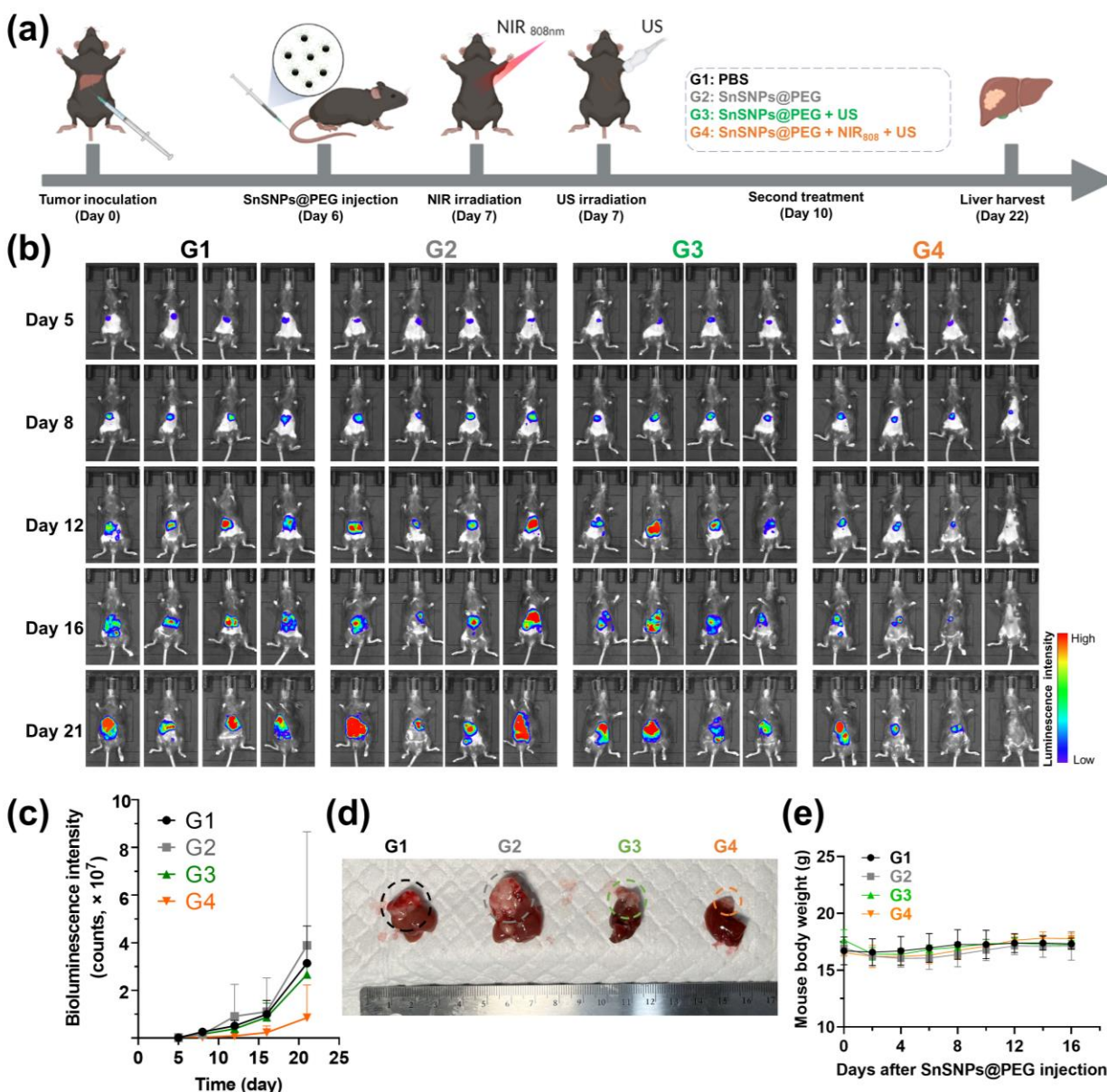




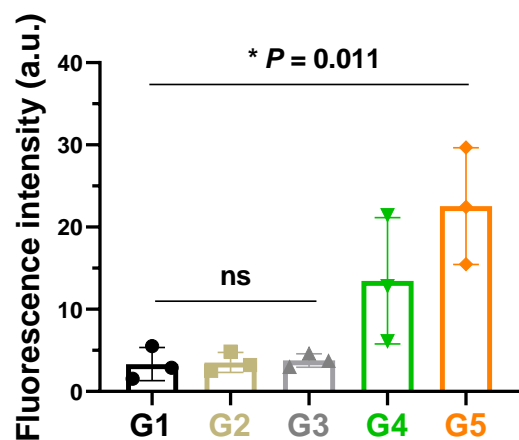
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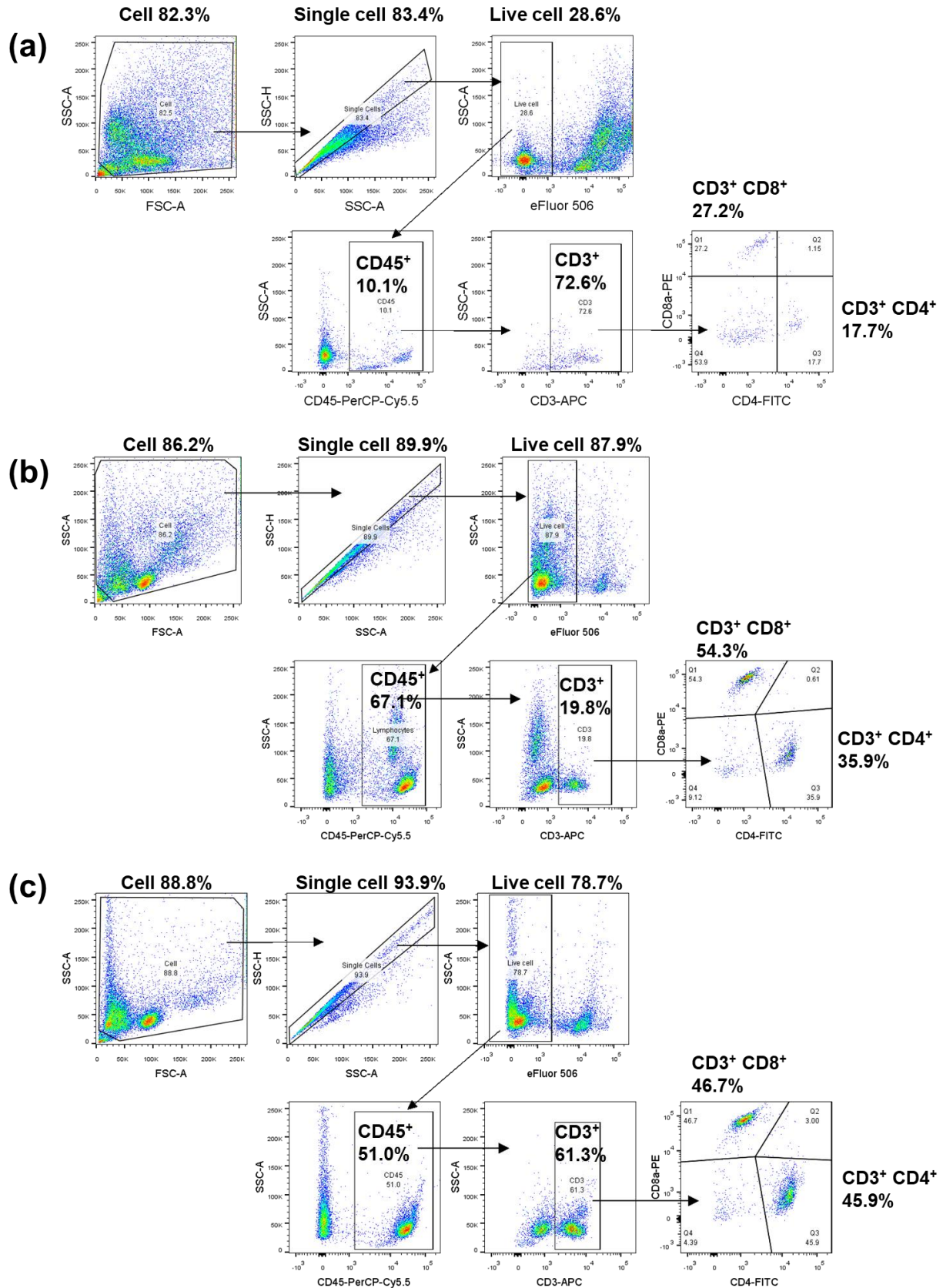
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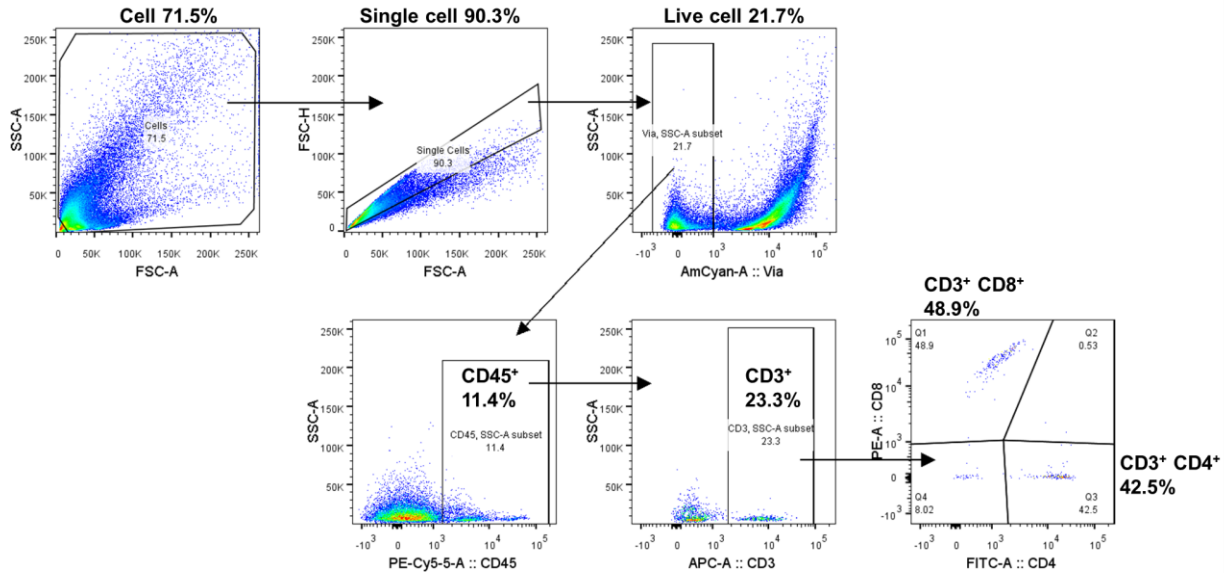
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**Figure S17.** Semi-quantitative analysis of green fluorescence intensity showing the ROS level in tumors of 4T1 tumor-bearing mice after various treatments. Data are presented as mean  $\pm$  SD ( $n = 3$ ). Statistical analysis between two groups was performed using student's  $t$ -test. \*  $P < 0.05$ . Groups are as follows: control (G1), NIR + US (G2), SnSNPs@PEG (G3), SnSNPs@PEG + US (G4), SnSNPs@PEG + NIR + US (G5), ( $n = 3$ ).

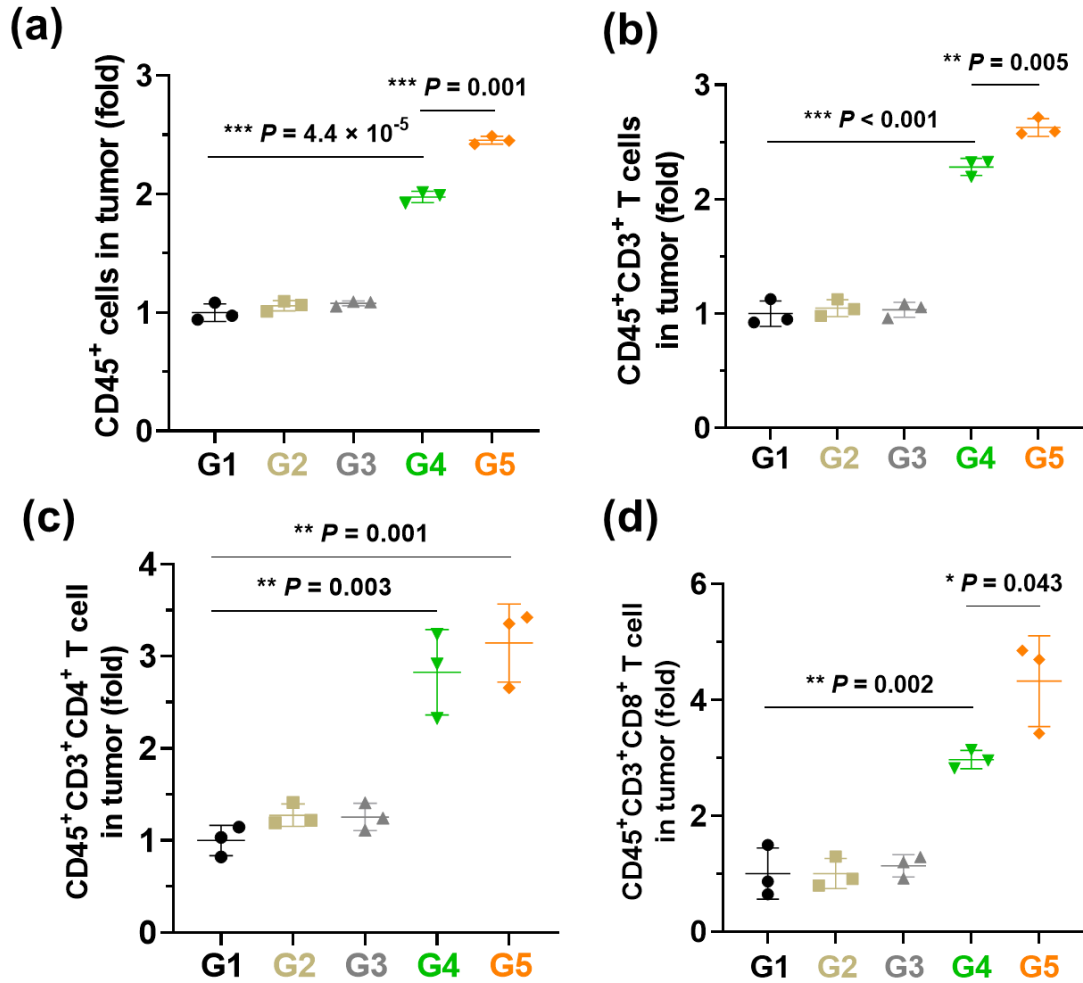


**Figure S18.** Gating strategies to identify CD45<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup> and CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> T lymphocytes in the tumor, spleen and lymph nodes of 4T1 tumor-bearing mice following the SnSNPs@PEG-mediated treatment strategy. (a) tumor, (b) spleen and (c) lymph node. Single live cells were first gated, followed by a selection of CD45<sup>+</sup>CD3<sup>+</sup> T lymphocytes. Subsequently, the populations of CD45<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup> and CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> T lymphocytes were distinguished and analyzed.

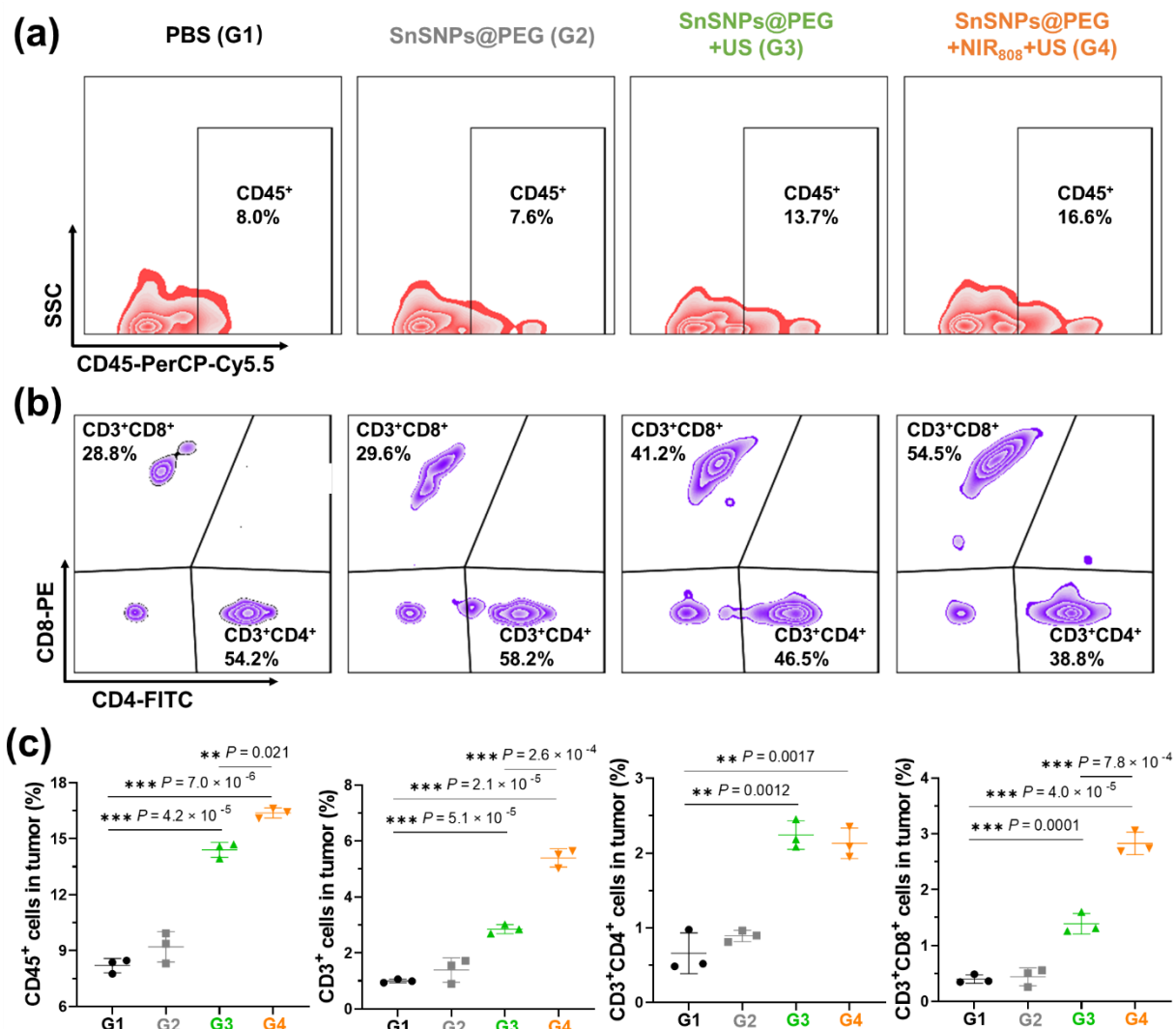


**Figure S19.** Gating strategy to identify CD45<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup> and CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> T lymphocytes in the RIL-175-HCC following the SnSNPs@PEG-mediated treatment strategy. Single live cells were first gated, followed by a selection of CD45<sup>+</sup>CD3<sup>+</sup> T lymphocytes. Subsequently, the populations of CD45<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup> and CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> T lymphocytes were distinguished and analyzed.



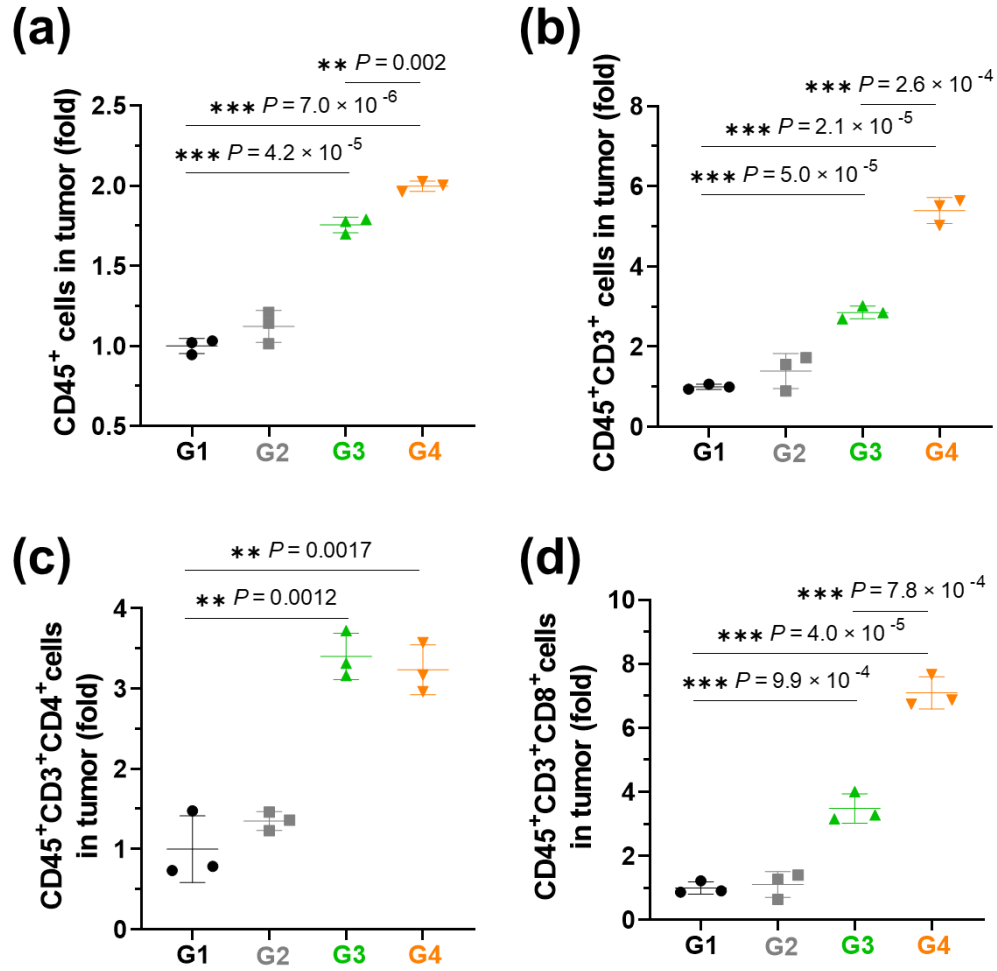


**Figure S20.** Comparison of immune cell levels in 4T1 tumors in different groups after SnSNPs@PEG-mediated treatment strategy. (a) CD45<sup>+</sup> immune cells, (b) CD45<sup>+</sup>CD3<sup>+</sup> T cells, (c) CD45<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup> T cells, and (d) CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> T cells in tumors after various treatments. Data are presented as mean  $\pm$  SD (n = 3). Statistical analysis between two groups was performed using two-sided student's *t*-test. \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ . Groups are as follows: control (G1), NIR + US (G2), SnSNPs@PEG (G3), SnSNPs@PEG + US (G4), SnSNPs@PEG + NIR + US (G5), (n = 3).

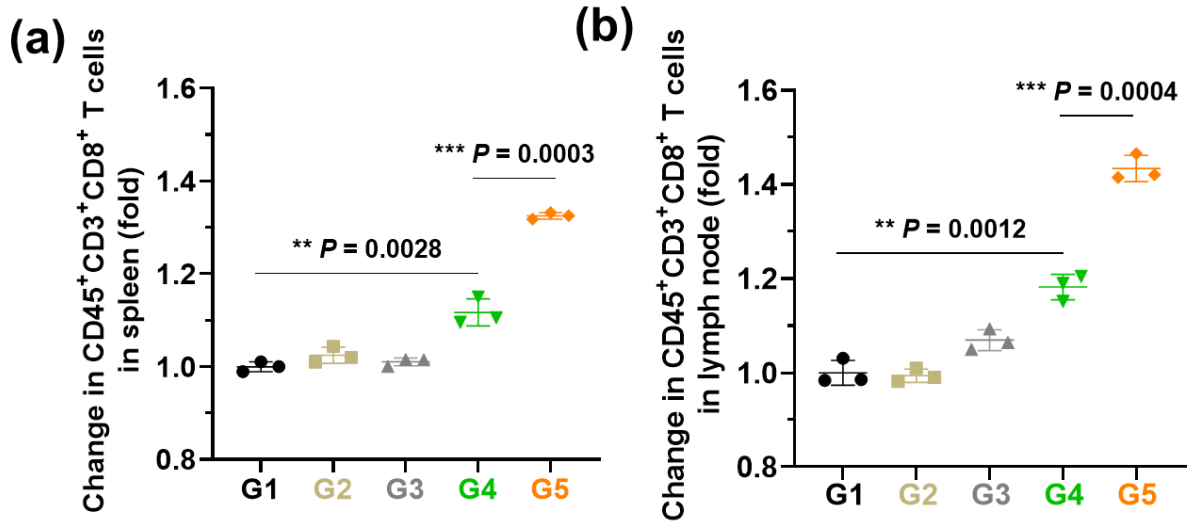


**Figure S21.** Flow cytometry analysis of (a) CD45<sup>+</sup> cells, (b)(c) CD45<sup>+</sup>CD3<sup>+</sup> lymphocytes, CD45<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup> T lymphocytes and CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> T lymphocytes in RIL-175-HCC after different treatments (n = 3). Statistical analysis between two groups was performed using two-sided student's *t*-test. Data are presented as mean  $\pm$  SD (n = 3) \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ . Groups are as follows: control (G1), SnSNPs@PEG (G2), SnSNPs@PEG + US (G3), SnSNPs@PEG + NIR + US (G4), (n = 3).





**Figure S22.** Comparison of the levels of (a) CD45<sup>+</sup> immune cells, (b) CD45<sup>+</sup>CD3<sup>+</sup> T cells, (c) CD45<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup> T cells, and (d) CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> T cells in RIL-175-HCC-bearing mice after various treatments (n = 3). Statistical analysis between two groups was performed using two-sided student's *t*-test. Data are presented as mean ± SD (n = 3). \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ . Groups are as follows: control (G1), SnSNPs@PEG (G2), SnSNPs@PEG + US (G3), SnSNPs@PEG + NIR + US (G4), (n = 3).



**Figure S23.** Comparison of CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> T cell levels in spleens and lymph nodes of 4T1 tumor-bearing mice in different groups after SnSNPs@PEG-mediated treatment strategy. (a) spleens and (b) lymph nodes of tumor-bearing mice after various treatments. Statistical analysis between two groups was performed using two-sided student's *t*-test. Data are presented as mean  $\pm$  SD (*n* = 3) \* *P* < 0.05, \*\* *P* < 0.01, \*\*\* *P* < 0.001. Groups are as follows: control (G1), NIR + US (G2), SnSNPs@PEG (G3), SnSNPs@PEG + US (G4), SnSNPs@PEG + NIR + US (G5), (*n* = 3).

**Table S1.** Comparison of the bandgap of SnSNPs with other nano-sonosensitizers.

<i>Sonosensitizer</i>	<i>Structure</i>	<i>Reported bandgap</i>	<i>Reference</i>
<i>SnS</i>	Nanoparticle	1.18 eV	This work
<i>Bi@BiO<sub>2-x</sub>@Bi<sub>2</sub>S<sub>3</sub></i>	Nanoparticle	1.43 eV	1
<i><math>\alpha</math>-Fe<sub>2</sub>O<sub>3</sub>@Pt</i>	Heterostructure particle	1.83 eV	2
<i>WO<sub>x</sub></i>	Nanobelt	2.11 eV	3
<i>Ti(Oi-Pr)<sub>4</sub>@Ag</i>	Metal–organic framework	2.11 eV	4
<i>Sn</i>	Nanosheet	2.3 eV	5
<i>BiVO<sub>4</sub></i>	Nanorod	2.5 eV	6
<i>Vanadium carbide</i>	Carbon dot	2.57 eV	7
<i>Sodium molybdenum bronze</i>	Nanoparticle	2.7 eV	8
<i>TiH<sub>1.924</sub></i>	Nanodot	2.7 eV	9
<i>TiO<sub>2</sub></i>	Nanoparticle	3.2 eV	10

**Table S2.** Antibody used for the flow cytometry analysis of T cells.

<b>Antibody</b>	<b>Brand and Catalog #</b>	<b>Concentration for use</b>
TruStain FcX™ (anti-mouse CD16/32)	Miltenyi Biotec; 130-092-575	1.0 µg/mL
PerCP/Cyanine5.5 anti-mouse CD45	BioLegend; 109828	2.0 µg/mL
APC anti-mouse CD3	BioLegend; 100236	2.5 µg/mL
FITC anti-mouse CD4	BioLegend; 100406	1.5 µg/mL
PE anti-mouse CD8	BioLegend; 100708	1.5 µg/mL

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