

Supplementary Information for

Polymeric Nanocarrier *via* Metabolism Regulation Mediates Immunogenic Cell Death with Spatiotemporal Orchestration for Cancer Immunotherapy

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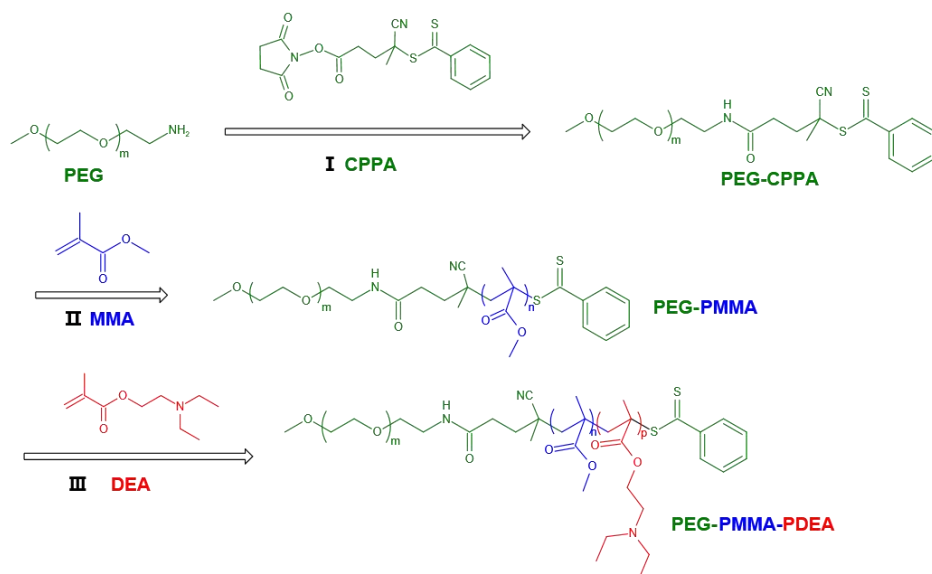
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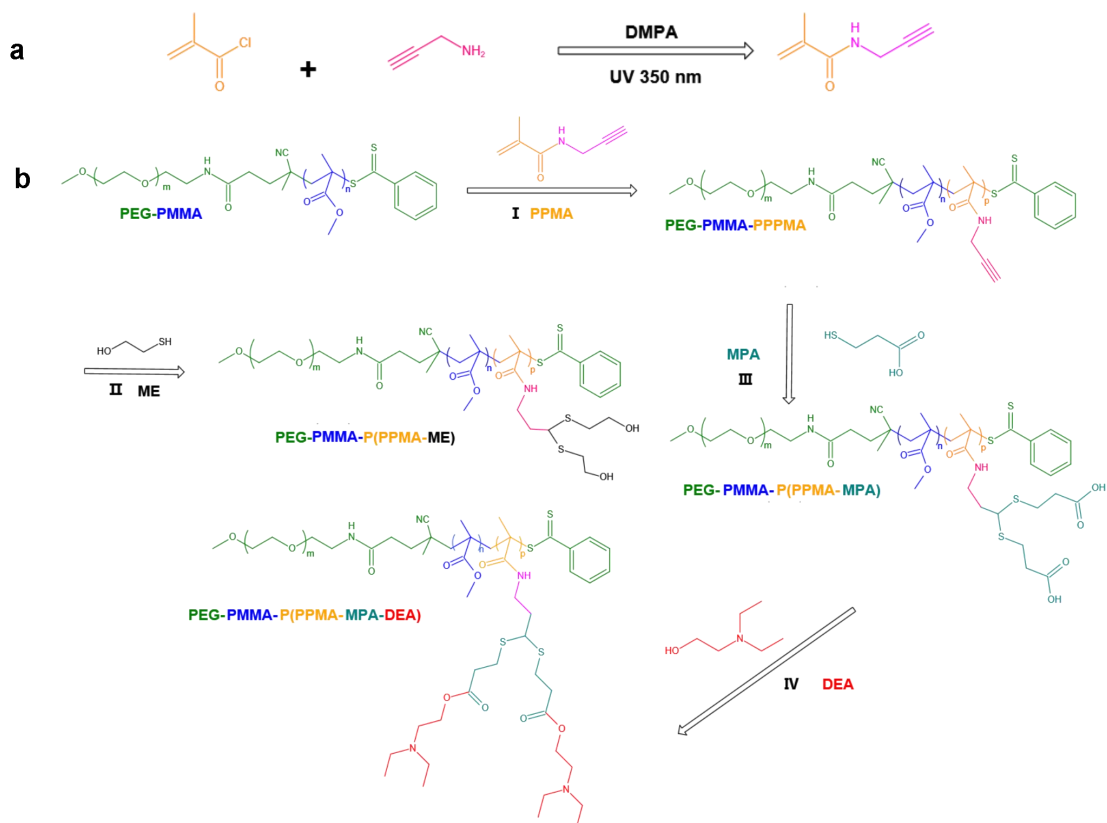
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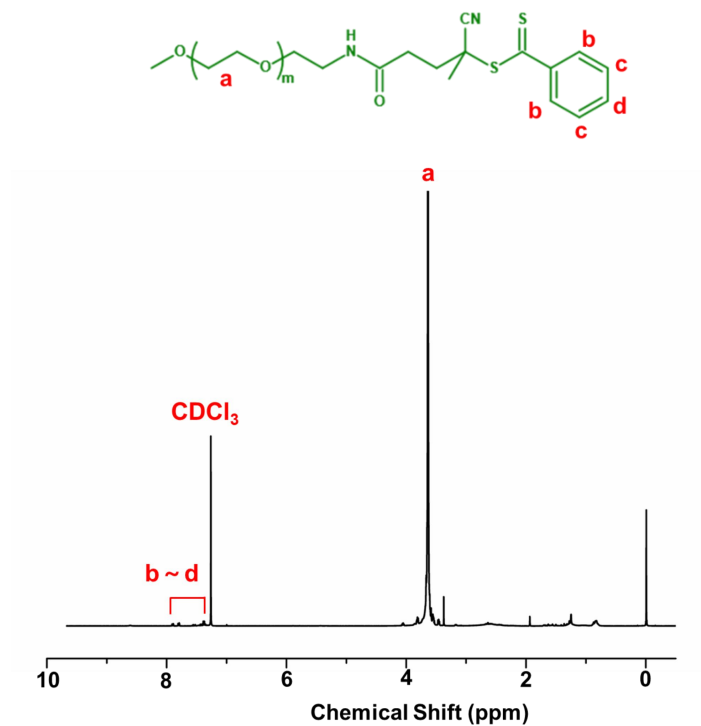
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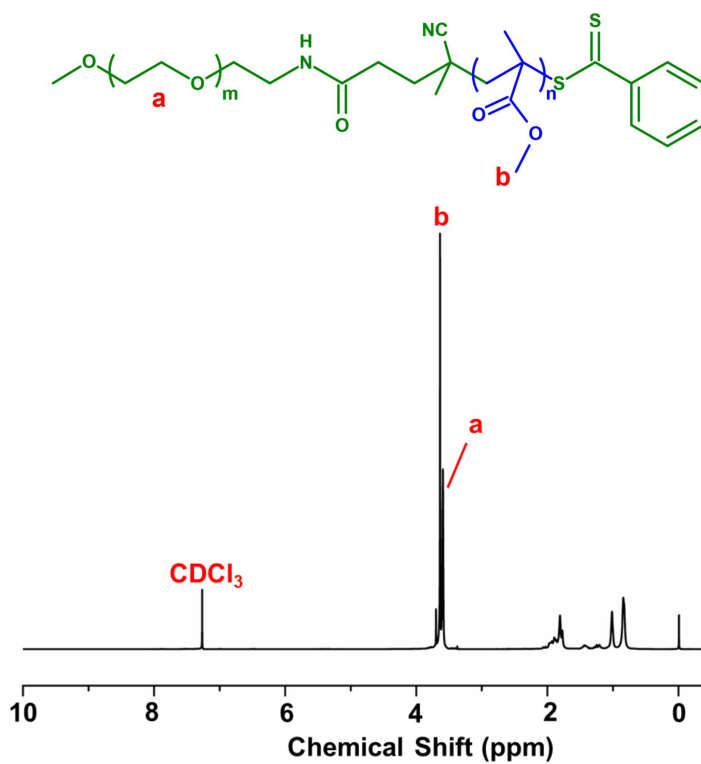
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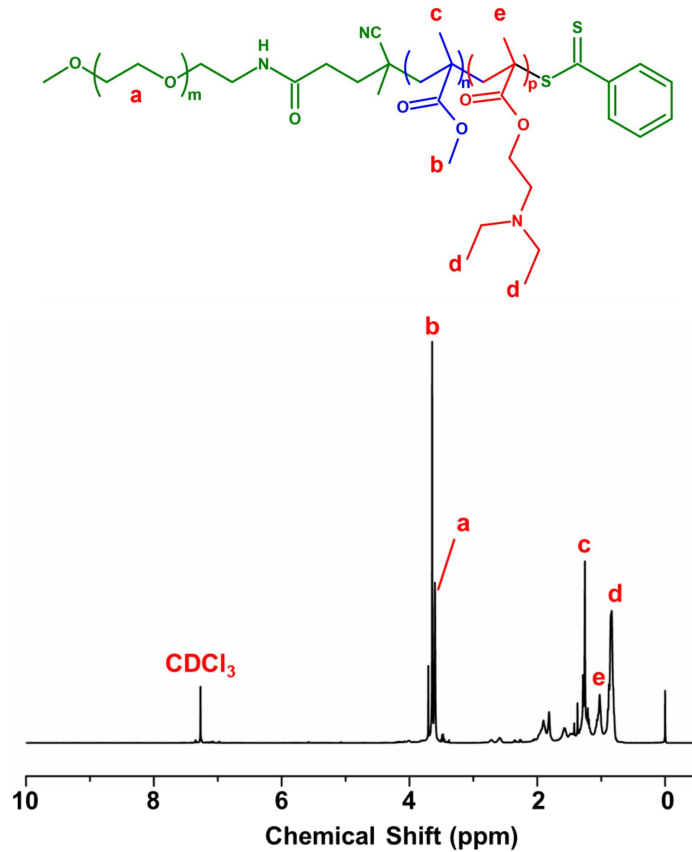
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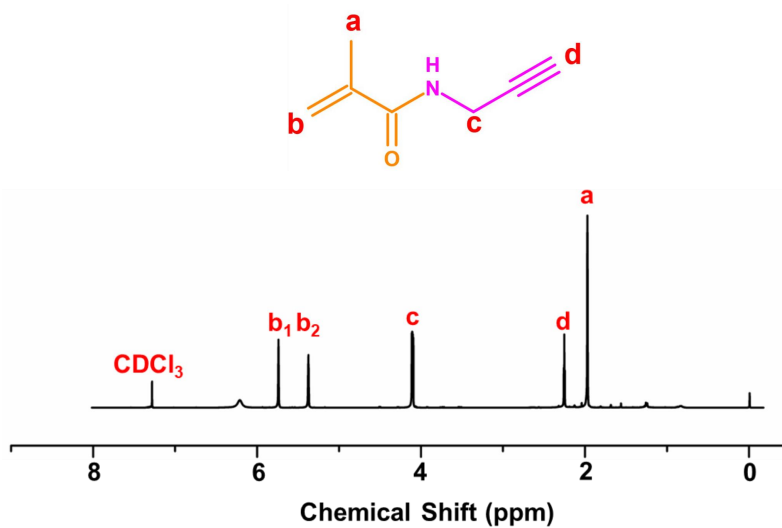
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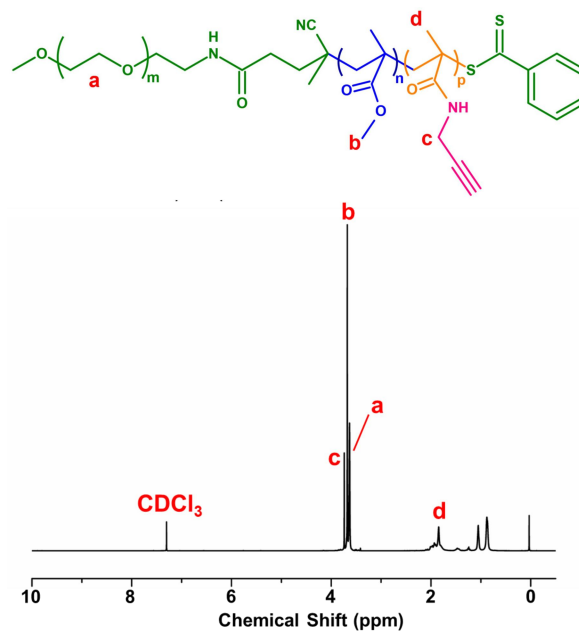
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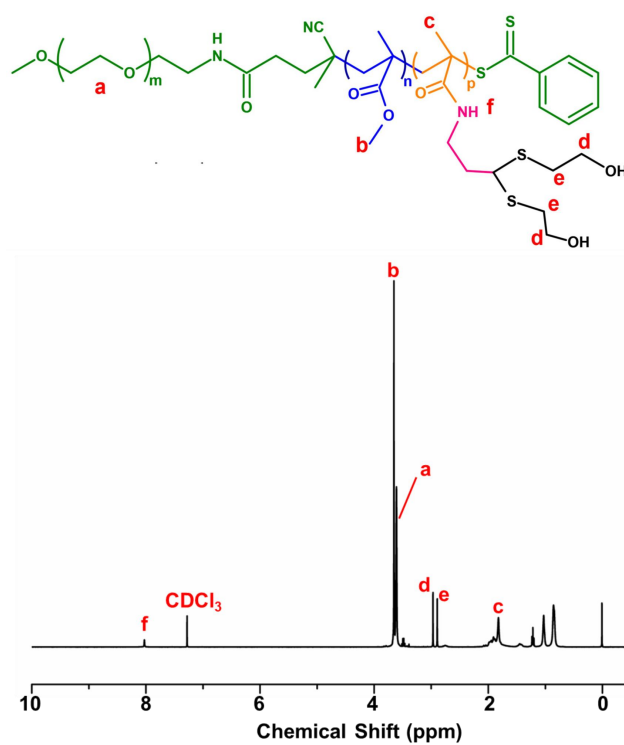
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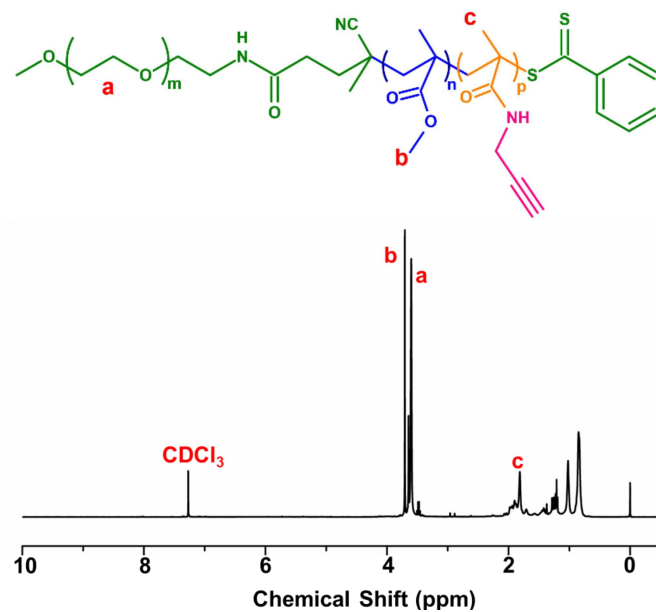
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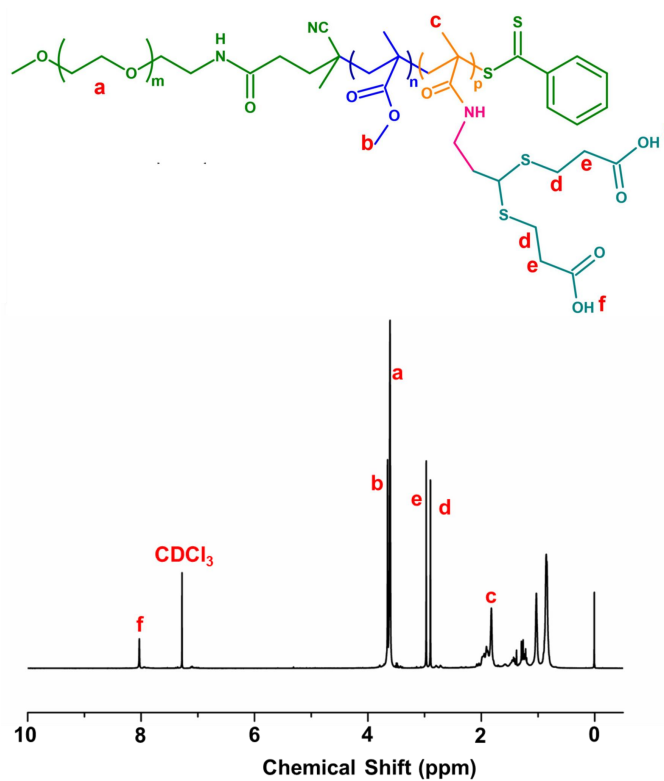
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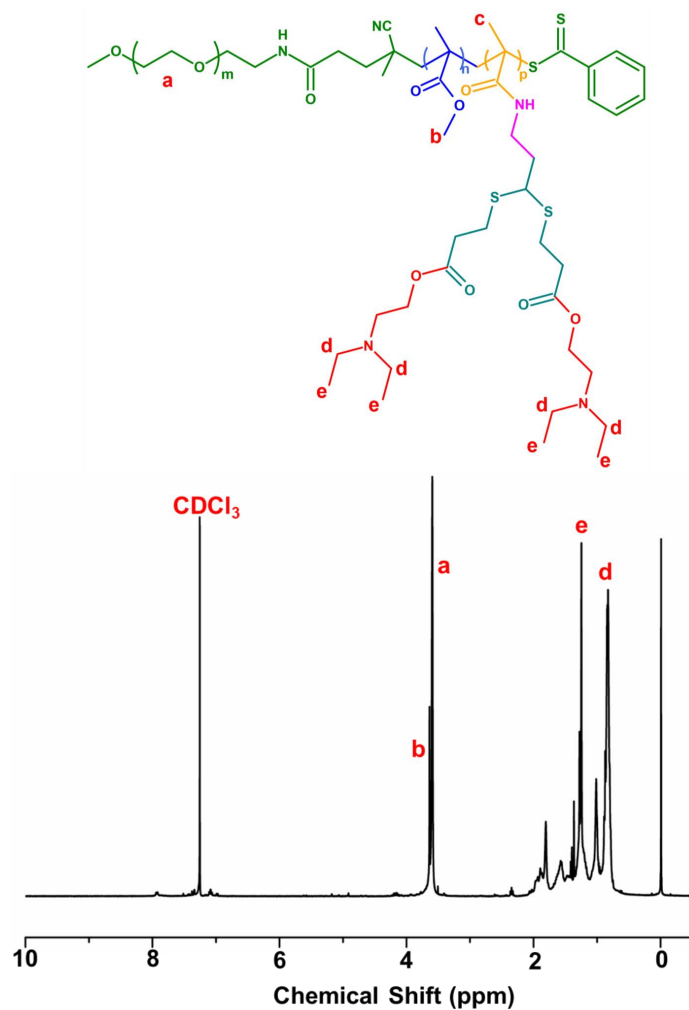
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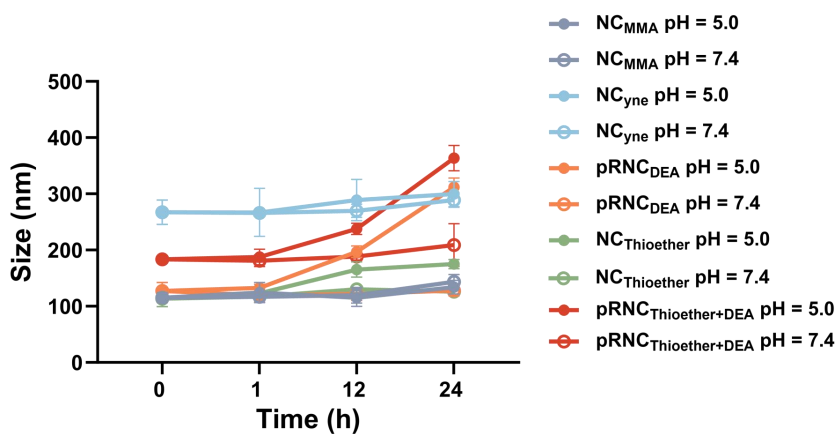
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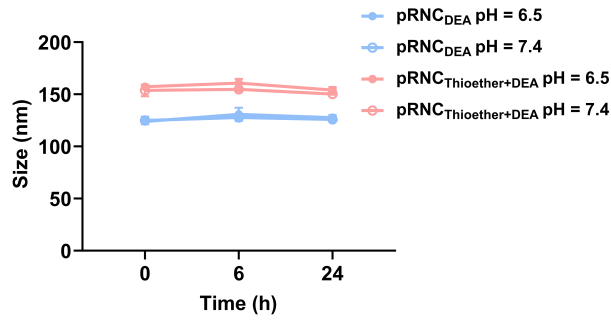
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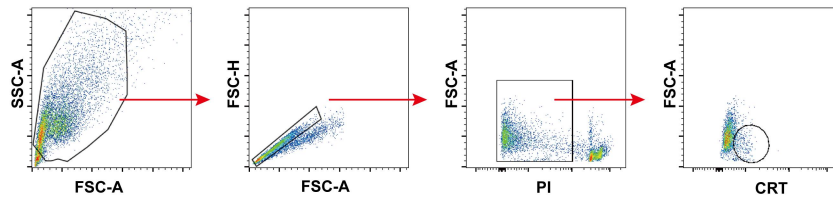
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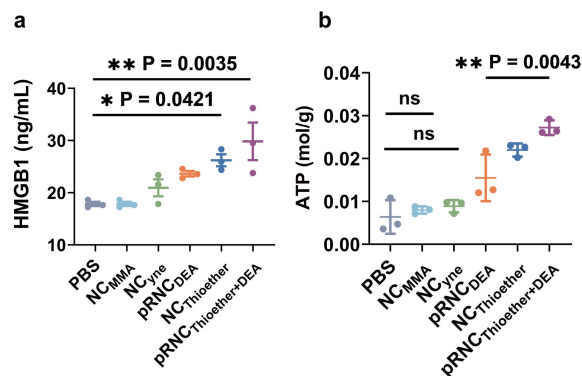
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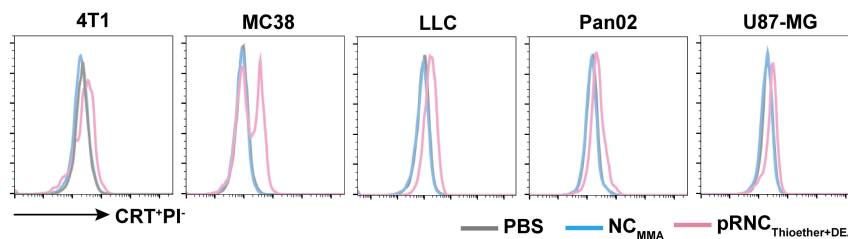
Supplementary Fig. 13. Hydrodynamic size of pRNC_{Thioether+DEA} and pRNC_{DEA} in acetate buffer solution (pH 6.5, 10 mM, 150 mM NaCl).



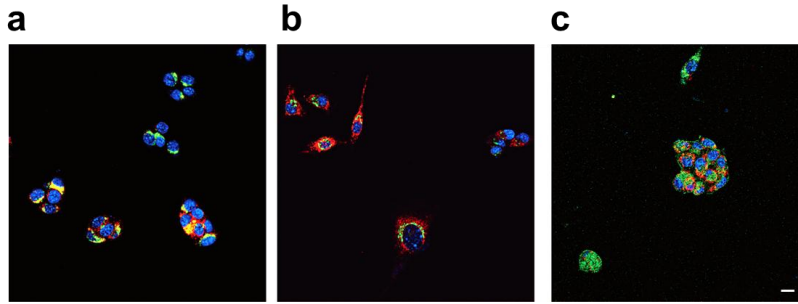
Supplementary Fig. 14. Scatter plots of the gating strategies for CRT⁺PI⁻ cells.



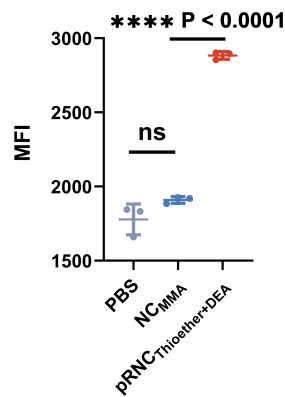
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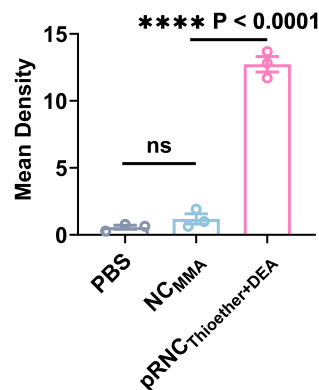
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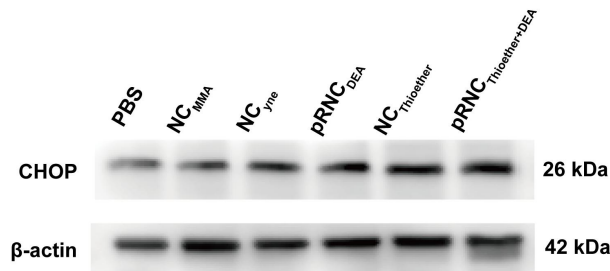


Supplementary Fig. 18. ROS flow cytometry quantification. Data are shown as mean ± SD (n = 3 independent experiments). The word “ns” represented non-significance, and ****p < 0.0001.

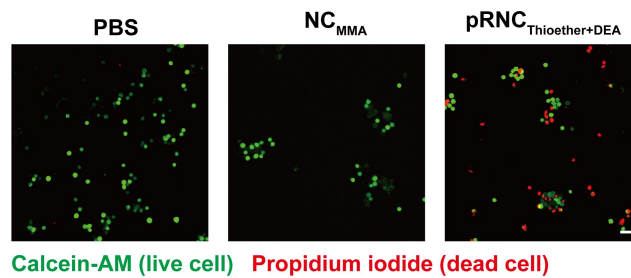


Supplementary Fig. 19. Quantification of intracellular mtROS level. Data are shown as

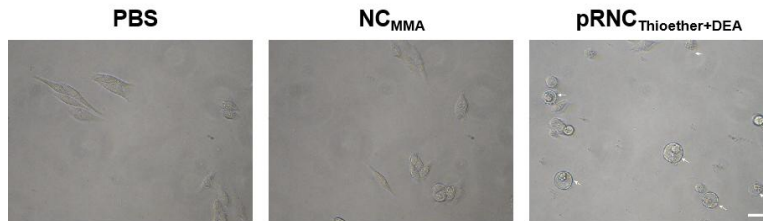
mean \pm SEM (n = 3 independent experiments). The word “ns” represented non-significance, and **** p < 0.0001.



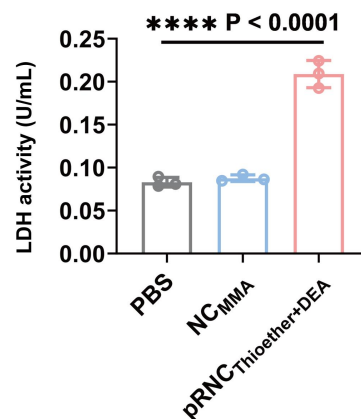
Supplementary Fig. 20. Western blot of CHOP expression after different treatments.



Supplementary Fig. 21. Imaging of Calcein-AM (green channel, living cells) and PI (red channel, dead cells) staining of cells after different treatments. Scale bar, 50 μ m.

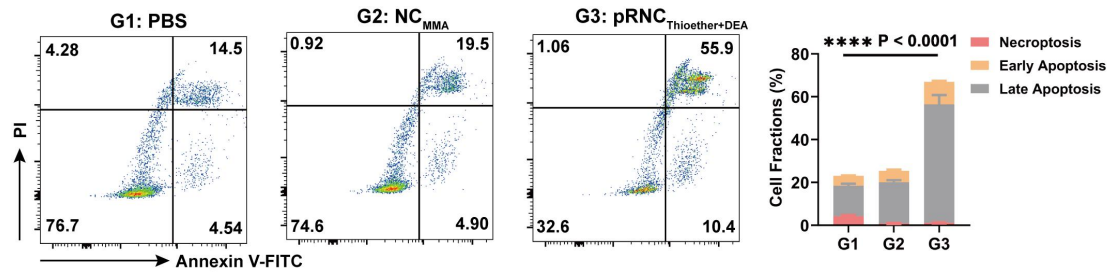


Supplementary Fig. 22. Morphological features of cells after various treatments. The white arrows represent pyroptotic cells. Scale bar, 30 μ m.

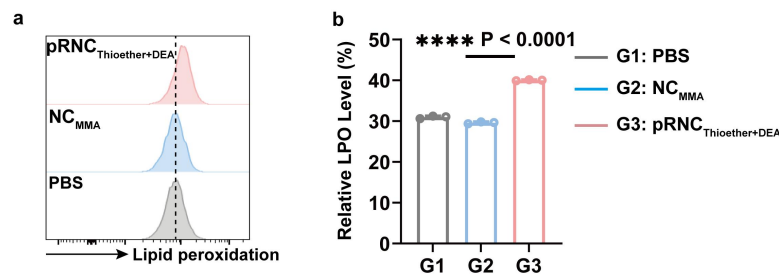


Supplementary Fig. 23. LDH activity in the supernatant after different treatments *via*

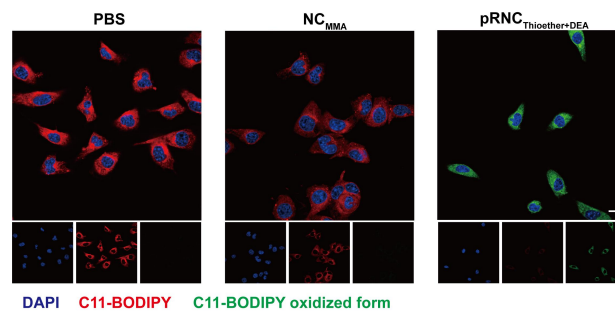
microplate plate test kit characterization (One unit means that pyruvate per micromole was produced in the reaction system when cell supernatant per milliliter was incubated with substrate at 37 °C for 15 min). Data are shown as mean ± SD (n = 3 independent experiments).



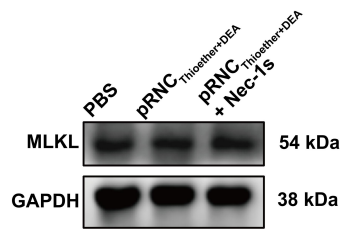
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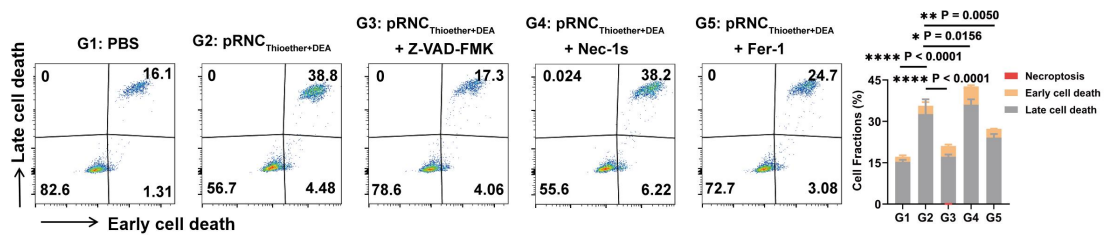
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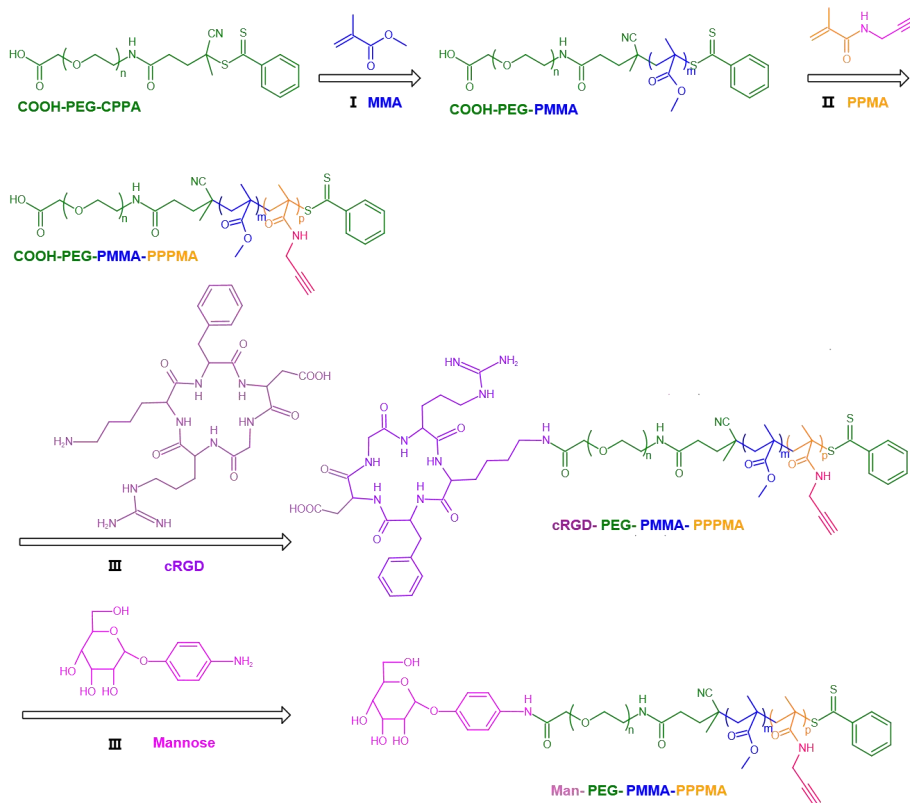
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Supplementary Fig. 27. Western blot of MLKL expression in B16F10 cells after treatments with PBS, pRNC_{Thioether+DEA} or pRNC_{Thioether+DEA} + Nec-1s. GAPDH was used as internal control.

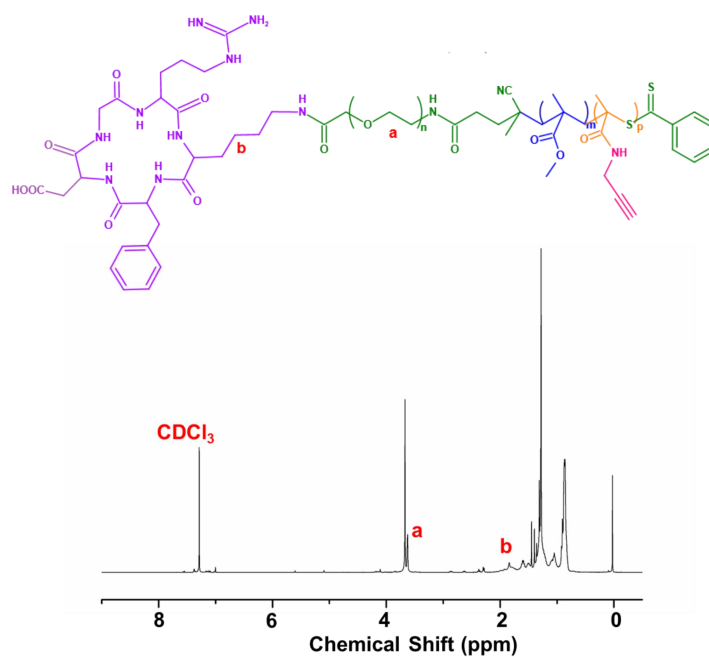


Supplementary Fig. 28. Representative flow cytometric images and the semi-quantitative analysis to show the cell death populations in B16F10 cells after treatment with PBS, pRNC_{Thioether+DEA} with or without cell death inhibitors. Data are shown as mean \pm SD (n = 3 independent experiments), *P < 0.05, **P < 0.01, ***P < 0.001, and ****P < 0.0001.

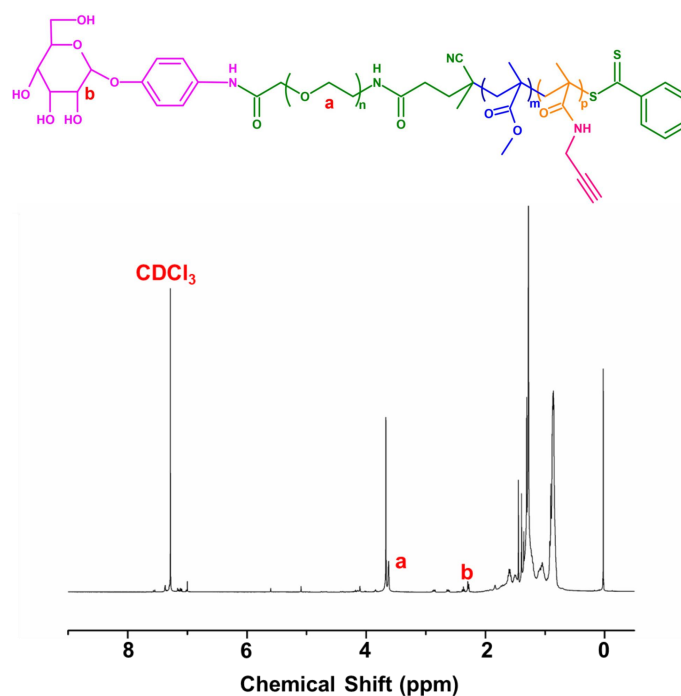


Supplementary Fig. 29. Synthesis of Man-PEG-PMMA-PPPMA and cRGD-PEG-PMMA-

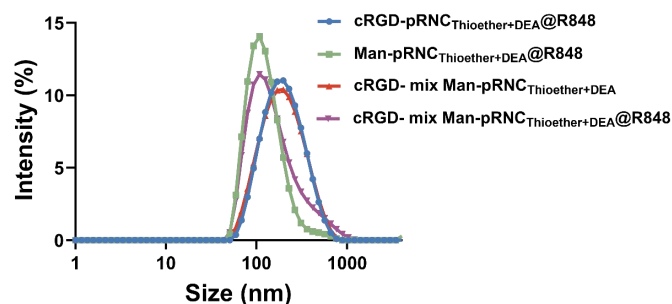
PPPMA. Conditions: (I) AIBN, 1, 4-dioxane, 70 °C, 48 h. (II) AIBN, 1, 4-dioxane, 70 °C, 48 h. (III) EDC·HCl, NHS, DMF, 30 °C, 24 h.



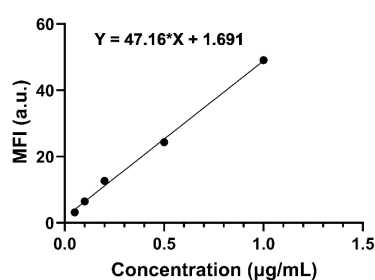
Supplementary Fig. 30. ¹H NMR spectrum (400 MHz, CDCl₃) of cRGD-PEG-PMMA-PPPMA.



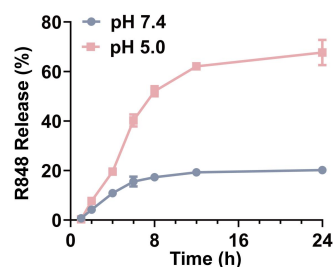
Supplementary Fig. 31. ¹H NMR spectrum (400 MHz, CDCl₃) of Man-PEG-PMMA-PPPMA.



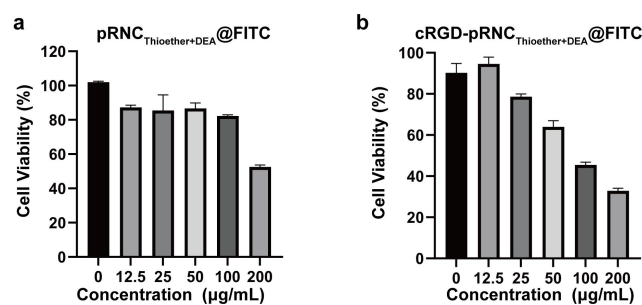
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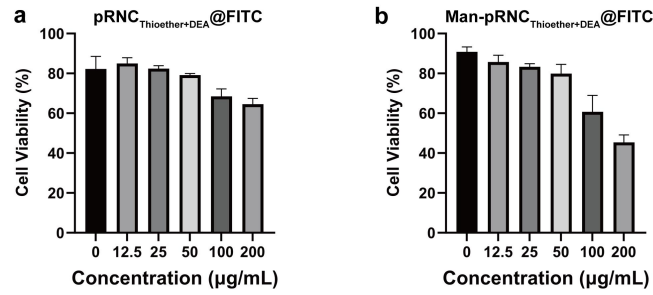
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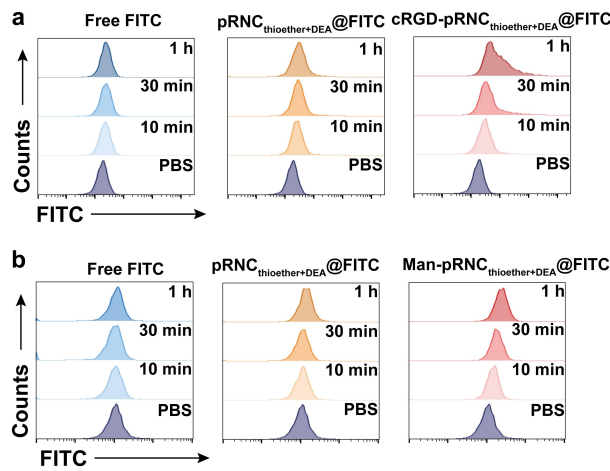
Supplementary Fig. 34. *In vitro* R848 release from cRGD- mix Man-pRNC_{Thioether+DEA} within 24 h at different pH values (pH 7.4 or 5.0) (n = 3 independent experiments).



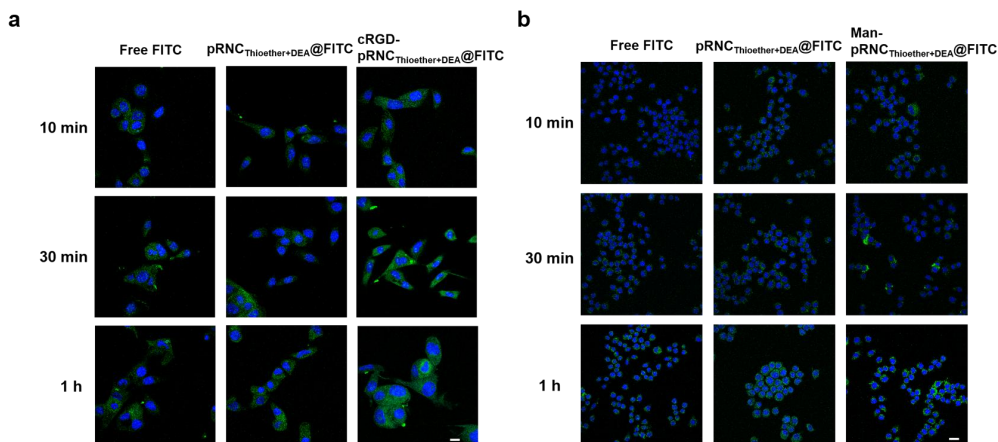
Supplementary Fig. 35. Dosage dependent cytotoxicity of a) pRNC_{Thioether+DEA}@FITC and b) cRGD-pRNC_{Thioether+DEA}@FITC in B16F10 cells by MTT assays. Data are shown as mean \pm SD (n = 4 independent experiments).



Supplementary Fig. 36. Dosage dependent cytotoxicity of a) pRNC_{Thioether+DEA}@FITC and b) Man-pRNC_{Thioether+DEA}@FITC in RAW264.7 cells by MTT assays. Data are shown as mean ± SD (n = 4 independent experiments).

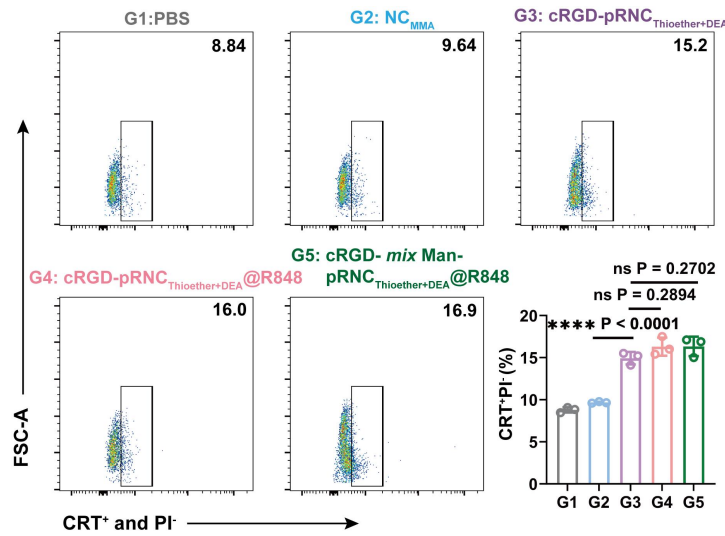


Supplementary Fig. 37. Cellular internalization of targeted nanoformulations *via* flow cytometry characterization. Representative flow cytometric plots showing the cellular uptake of a) pRNC_{Thioether+DEA}, cRGD-pRNC_{Thioether+DEA} and free FITC at different time points in B16F10 cells and b) pRNC_{Thioether+DEA}, Man-pRNC_{Thioether+DEA} and free FITC in RAW 264.7 cells.

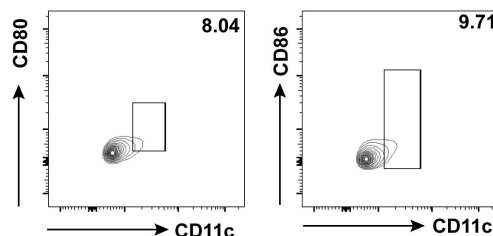


Supplementary Fig. 38. Cellular internalization of different nanoformulations *via* CLSM characterization. Representative images of a) cRGD-pRNC_{Thioether+DEA}@FITC uptake in

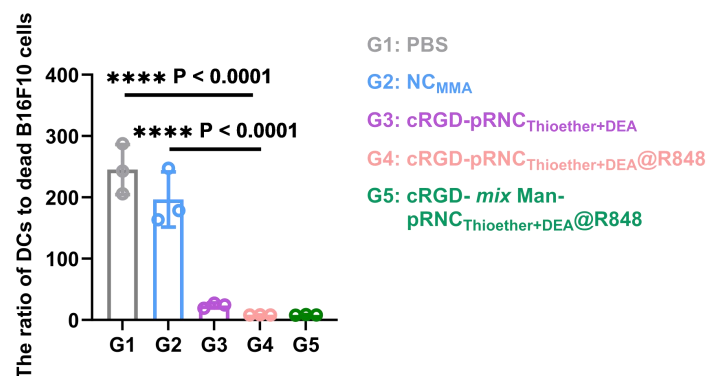
B16F10 cells, and b) Man-pRNC_{Thioether+DEA}@FITC in RAW264.7 cells. Scale bar = 20 μ m.



Supplementary Fig. 39. Representative flow cytometric images and quantification analysis of CRT exposure in B16F10 cells after different treatments *via* flow cytometry characterization. Data are presented as mean \pm SD (n = 3 independent experiments). Statistical significance was calculated through one-way ANOVA for multiple comparisons using a Tukey post-hoc test.

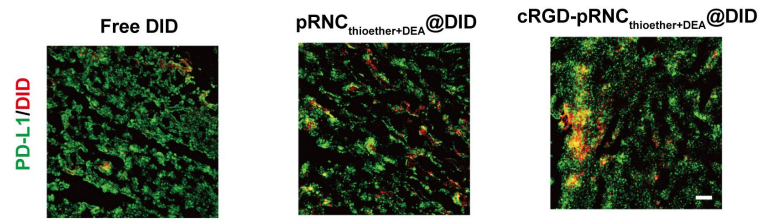


Supplementary Fig. 40. Negative control of DC maturation investigation using viable B16F10 cells directly to incubate with DCs.

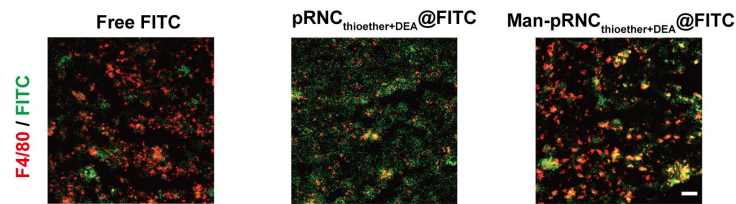


Supplementary Fig. 41. The ratio of DCs to dead B16F10 cells after different treatments. Data are presented as mean \pm SD (n = 3 independent experiments). Statistical significance was

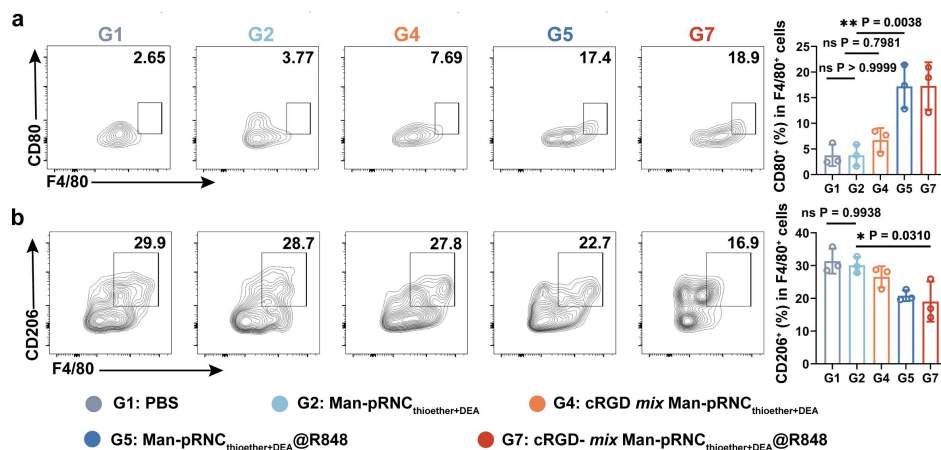
calculated through one-way ANOVA for multiple comparisons using a Tukey post-hoc test.



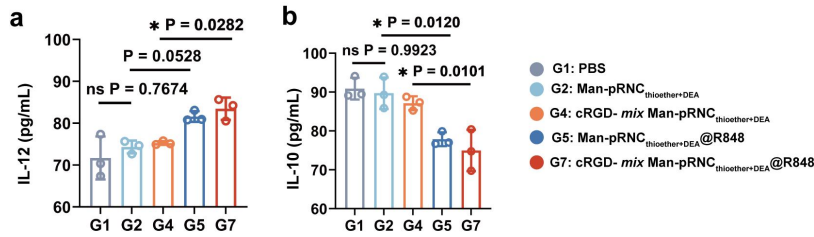
Supplementary Fig. 42. Representative immunofluorescence images showing tumor targeting of cRGD-pRNC_{Thioether+DEA}@DID at 24 h treatment. Scale bar = 50 μ m.



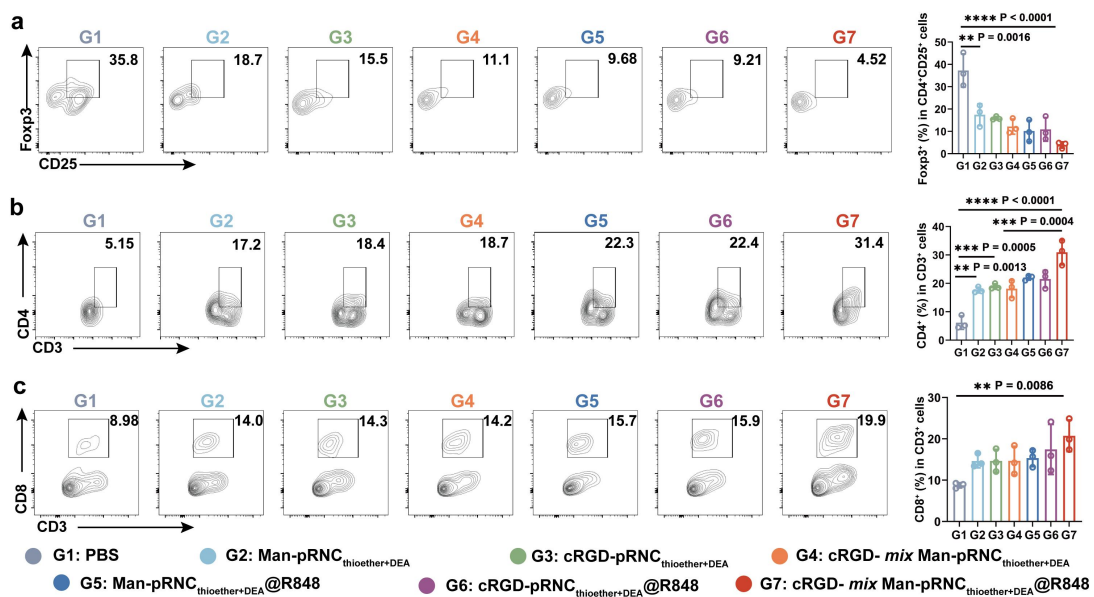
Supplementary Fig. 43. Representative immunofluorescence images showing TAMs targeting of Man-pRNC_{Thioether+DEA}@FITC *in vivo* at 24 h treatment. Scale bar = 50 μ m.



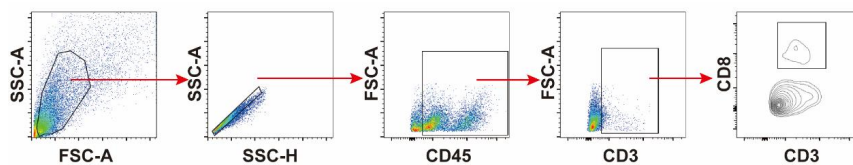
Supplementary Fig. 44. Flow cytometric analysis to show TAM polarization. a) Representative flow cytometry plots (left) and quantitative analysis (right) of CD80⁺F4/80⁺ TAM. b) Representative flow cytometry images (left) and quantitative data (right) of CD206⁺F4/80⁺ TAM. Data are presented as mean \pm SD (n = 3 mice per group). Statistical significance was calculated through one-way ANOVA for multiple comparisons using a Tukey post-hoc test.



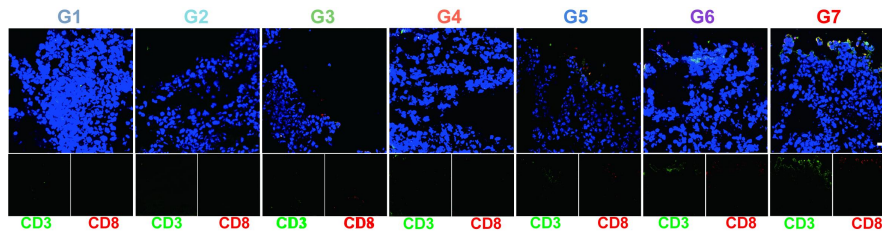
Supplementary Fig. 45. IL-12 and IL-10 levels in mice after different treatments. a) IL-12 and b) IL-10 levels measured by ELISA. Data are presented as mean \pm SD (n = 3 mice per group). Statistical significance was calculated through one-way ANOVA for multiple comparisons using a Tukey post-hoc test.



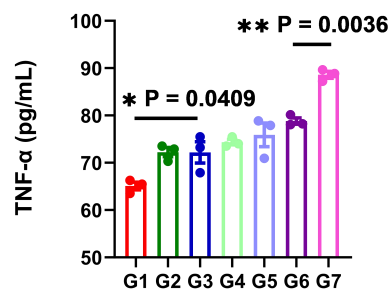
Supplementary Fig. 46. Immune cell ratios analysis in tumor tissues for mice after different treatments. a) Representative flow cytometric analysis gating on CD25⁺ cells and quantification of Foxp3⁺ Tregs in tumors. b) Representative flow cytometric analysis and quantification of CD4⁺ gating on CD3⁺ T cells. c) Representative flow cytometric analysis and quantification of CD8⁺ gating on CD3⁺ T cells. Data are presented as mean \pm SD (n = 3 mice per group). Statistical significance was calculated through one-way ANOVA for multiple comparisons using a Tukey post-hoc test.



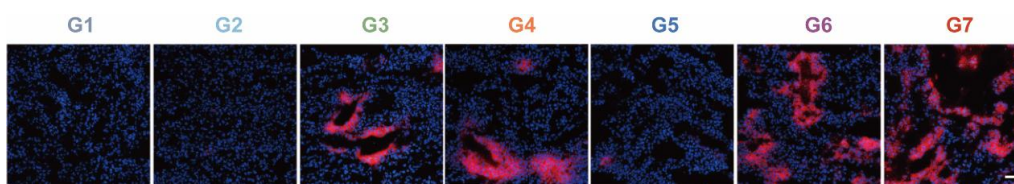
Supplementary Fig. 47. Scatter plots show the gating strategies for CD3⁺CD8⁺ T cells.



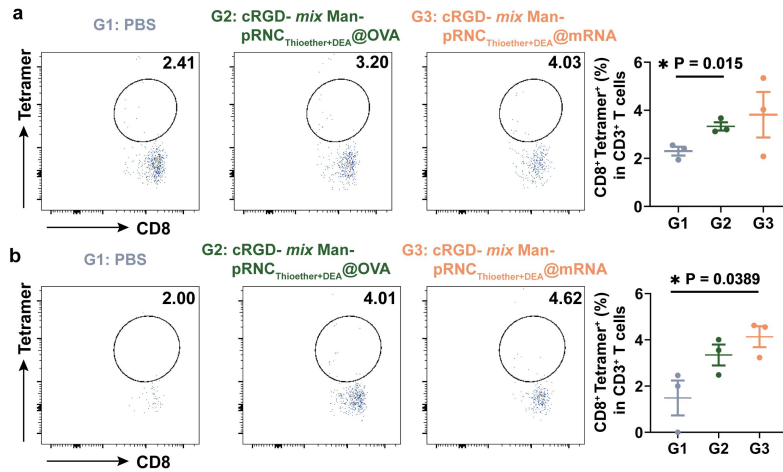
Supplementary Fig. 48. Representative immunofluorescence images showing CD3⁺CD8⁺ T lymphocytes infiltration. Nuclei were stained with DAPI. Scale bar = 50 μ m. (G1: PBS, G2: Man-pRNC_{Thioether+DEA}, G3: cRGD-PRNC_{Thioether+DEA}, G4: cRGD- *mix* Man-pRNC_{Thioether+DEA}, G5: Man-pRNC_{Thioether+DEA@R848}, G6: cRGD-PRNC_{Thioether+DEA@R848}, G7: cRGD- *mix* Man- pRNC_{Thioether+DEA@R848})



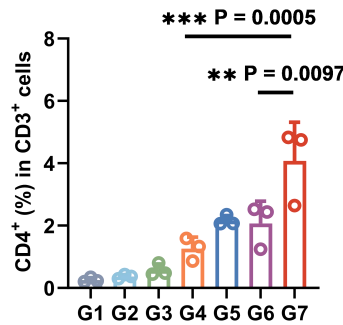
Supplementary Fig. 49. TNF- α level in mouse serum characterization by ELISA kit. (G1: PBS, G2: Man-pRNC_{Thioether+DEA}, G3: cRGD-PRNC_{Thioether+DEA}, G4: cRGD- *mix* Man-pRNC_{Thioether+DEA}, G5: Man-pRNC_{Thioether+DEA@R848}, G6: cRGD-PRNC_{Thioether+DEA@R848}, G7: cRGD- *mix* Man- pRNC_{Thioether+DEA@R848}). Data are presented as mean \pm SEM (n = 3 mice per group). Statistical significance was calculated through one-way ANOVA for multiple comparisons using a Tukey post-hoc test.



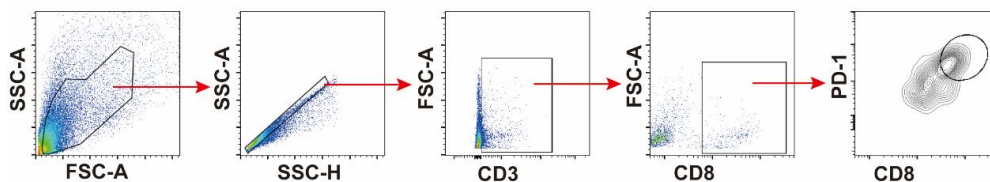
Supplementary Fig. 50. Representative immunofluorescence images of tumors with HMGB1 release. Red represented HMGB1 stained with Alexa Fluor[®]647 anti HMGB1, and blue represented nuclei stained with DAPI. Scale bar = 50 μ m. (G1: PBS, G2: Man-pRNC_{Thioether+DEA}, G3: cRGD-PRNC_{Thioether+DEA}, G4: cRGD- *mix* Man-pRNC_{Thioether+DEA}, G5: Man-pRNC_{Thioether+DEA@R848}, G6: cRGD-PRNC_{Thioether+DEA@R848}, G7: cRGD- *mix* Man-pRNC_{Thioether+DEA@R848}).



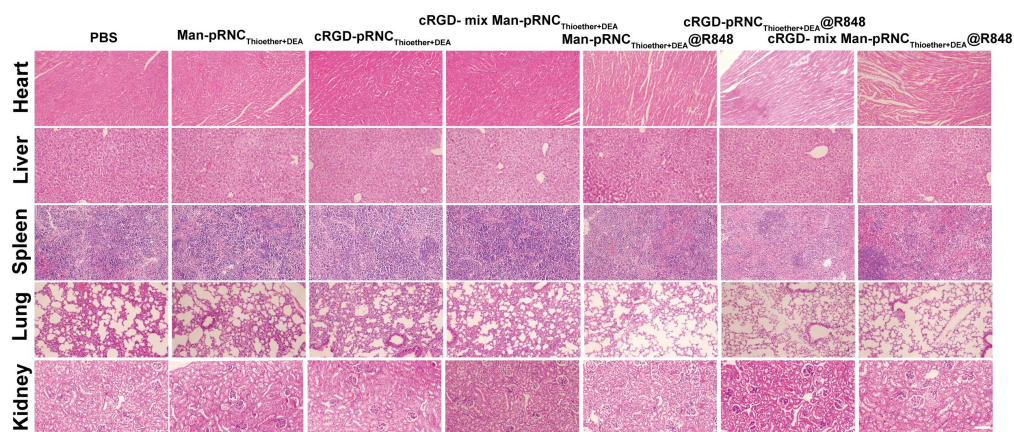
Supplementary Fig. 51. Representative flow dot plots and statistics of CD8⁺Tetramer⁺ T cells in peripheral blood mononuclear cells of mice a) day-7 and b) day-14 after immunization. Data are presented as mean ± SEM (n = 3 mice per group). Statistical significance was calculated through two-tailed student's *t* test.



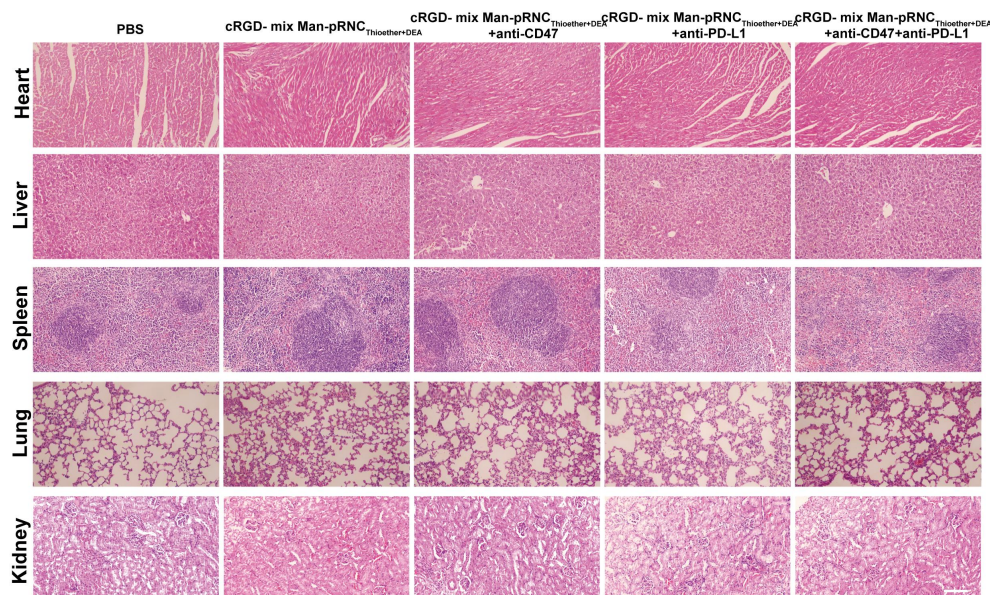
Supplementary Fig. 52. Quantitative ratio of CD4⁺ in CD3⁺ T cells. (G1: PBS, G2: Man-pRNC_{Thioether+DEA}, G3: cRGD-PRNC_{Thioether+DEA}, G4: cRGD- *mix* Man-pRNC_{Thioether+DEA}, G5: Man-pRNC_{Thioether+DEA@R848}, G6: cRGD-PRNC_{Thioether+DEA@R848}, G7: cRGD- *mix* Man-pRNC_{Thioether+DEA@R848}). Data are presented as mean ± SD (n = 3 mice per group). Statistical significance was calculated through one-way ANOVA for multiple comparisons using a Tukey post-hoc test.



Supplementary Fig. 53. Scatter plots of the gating strategies for CD8⁺PD-1⁺ T cells.



Supplementary Fig. 54. H&E staining of normal tissues harvested from mice in different groups. Scale bar = 100 μm .



Supplementary Fig. 55. H&E staining of normal tissues harvested from mice in different groups. Scale bar = 100 μm .

Supplementary Table 1. Characteristics of block copolymers.

Copolymers	M_n (kg/mol)		GPC	
	Design	$^1\text{H NMR}^a$	M_n^b (kg/mol)	M_w/M_n^b
PEG-PMMA	5.0-12.0	5.0-10.0	15.3	1.30
PEG-PMMA-PPMA	5.0-12.0-3.0	5.0-10.0-6.3	18.9	1.39
PEG-PMMA-PDEA	5.0-12.0-4.6	5.0-10.0-3.8	19.2	1.35
PEG-PMMA-P(PPMA-ME)	5.0-12.0-9.3	5.0-10.0-10.4	30.7	1.14
PEG-PMMA-P(PPMA-MPA-DEA)	5.0-12.0-16.5	5.0-7.3-18.4	36.1	1.41

^a Calculated from $^1\text{H NMR}$ by comparing the intensities of signals at δ 3.65, 3.60, 2.97, 1.82,

1.24 and 0.82, respectively.

^b Analyzed by GPC using THF as an eluent at a flow rate of 1.0 mL/min (standards: PMMA, 30 °C).

Supplementary Table 2. Preparation of blank polymersomes.

Polymersomes	Size (nm) ^a	PDI ^a
NC _{MMA}	118.2 ± 1.5	0.24 ± 0.006
NC _{yne}	138.7 ± 8.2	0.25 ± 0.032
pRNC _{DEA}	126.5 ± 3.6	0.22 ± 0.013
NC _{Thioether}	132.8 ± 5.6	0.22 ± 0.025
pRNC _{Thioether+DEA}	140.7 ± 3.5	0.29 ± 0.010

^a Determined by Malvern Panalytical Zetasizer Pro at 25 °C in PBS (10 mM, pH 7.4).

Supplementary Table 3. Drug loading content and efficiency characterization.

Polymersomes	Thero. DLC (wt.%)	Size ^a (nm)	PDI ^a	DLC ^b (wt.%)	DLE ^b (%)
cRGD-NP _{Thioether+DEA@R848}	5%	168.0 ± 4.8	0.22 ± 0.02	4.79	95.6
	10%	236.4 ± 2.5	0.12 ± 0.10	8.28	81.3
	20%	246.5 ± 5.9	0.18 ± 0.02	12.70	58.2
Man-NP _{Thioether+DEA@R848}	5%	143.3 ± 5.5	0.16 ± 0.01	4.85	96.8
	10%	143.5 ± 1.0	0.17 ± 0.01	8.46	83.2
	20%	171.4 ± 4.0	0.06 ± 0.02	13.29	61.3
cRGD- <i>mix</i> Man- pRNC _{Thioether+DEA@R848}	5%	169.9 ± 22.7	0.21 ± 0.06	4.65	92.6
	10%	183.3 ± 4.6	0.16 ± 0.04	8.15	79.9
	20%	176.2 ± 3.9	0.28 ± 0.03	12.55	57.4

^a Determined using Malvern Panalytical Zetasizer Pro at 25 °C in PBS (10 mM, pH 7.4).

^b Determined by fluorescence spectrophotometer measurement.