SUPPLEMENTARY INFORMATION

Precise delivery of a multifunctional nanosystem for MR-guided cancer therapy and monitoring of tumor response by functional diffusion-weighted MRI

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Results



Figure S1. The chemical structures of the PLGA-PEI-mPEG-cRGD copolymers. The ¹H NMR and ¹³C NMR spectra of PLGA, PLGA-PEI, PLGA-PEI-mPEG, and PLGA-PEI-mPEG-cRGD are shown. The chemical shifts of the characteristic H and C are shown in the tables.



Figure S2. The FT-IR spectra of the PLGA, PLGA-PEI, PLGA-PEI-mPEG, and PLGA-cRGD copolymers. The black arrows indicate the amine group (1559 cm⁻¹) and the amide group (1280 cm⁻¹).



Figure S3. TEM images of cRGD-PLGA-SPIO@DOX nanoparticles under different pH conditions (pH = 7.4, 6.0 and 5.3).



Figure S4. Change in particle size of the cRGD-PLGA-SPIO@DOX nanoparticles in DMEM+FBS and PBS solution for 10 days.



Figure S5. IC₅₀ values for *in vitro* cytotoxicity against NIH-3T3 cancer cells. The averages and standard deviations from three experiments are shown.



Figure S6. Quantitative analysis of cellular uptake of DOX and cRGD-PLGA-SPIO@DOX nanoparticles in L02 cells for 0.5, 1, 2, 4 and 6 h.



Figure S7. (A)The competition cellular uptake assay of cRGD-PLGA-SPIO@DOX nanoparticles. A549 cells were pretreated with 0.25, 0.5, or 1 mg/mL cRGD peptide and then treated with 10 μ M cRGD-PLGA-SPIO@DOX nanoparticles for 6 h. (B) A549 cell viabilities in the competition uptake assays. A549 cells were pretreated with 0.25, 0.5, or 1.0 mg/mL cRGD peptide for 1 h, followed by incubation with 10, 5, 2.5, 1.25, or 0.625 μ M cRGD-PLGA-SPIO@DOX nanoparticles for 72 h. The data are presented as the average ± standard deviation (n=3). Significant differences between the DOX and cRGD-PLGA-SPIO@DOX groups are indicated at the *P*< 0.05 (*) or P< 0.01 (**) level.



Figure S8. (A) Hematoxylin-eosin (H&E) and Prussian blue staining assays to verify Fe accumulation in the tumor, liver, spleen and lung. (B) Fe quantification in the tumor, liver, spleen and lung by ICP-MS in the control, SPIO, cRGD-SPIO, and cRGD-PLGA-SPIO@DOX groups.



Figure S9. H&E-stained slice images of major organs (×20) from the DOX (2 mg/kg), cRGD-PLGA-SPIO@DOX (1 mg/kg), and cRGD-PLGA-SPIO@DOX (2 mg/kg) groups.



Figure S10. MR T₂WI anatomical images of representative Control, DOX and cRGD-PLGA-SPIO@DOX-treated tumors before and at different time points after treatment.