Supplementary Information

A Sequentially Responsive and Structure-transformable Nanoparticle with Comprehensively Improved 'CAPIR cascade' for Enhanced Antitumor Effect

Chenfeng Xu^a, Yu Sun^b, YulinYu^a, Mei Hu^a, Conglian Yang^a, Zhiping Zhang^{a, c, d,}

AUTHOR ADDRESS

^a Tongji School of Pharmacy, Huazhong University of Science and Technology, Wuhan 430030, China

^b Ningbo First Hospital, Ningbo 315010, China.

^c National Engineering Research Center for Nanomedicine, Huazhong University of Science and Technology, Wuhan 430030, China

^d Hubei Engineering Research Centre for Novel Drug Delivery System, Huazhong University of Science and Technology, Wuhan 430030, China



Fig.S1 The HRMS spectrum of 2-(Nap)-FFKAGLDD-RGD.



Fig.S2 The HPLC spectrum of 2-(Nap)-FFKAGLDD-RGD, purify is 94.76%.



Fig.S3 The HRMS spectrum of 2-(Nap)-FFKWGLWD-RGD.



Fig.S4 The HPLC spectrum of 2-(Nap)-FFKWGLWD-RGD, purity is 93.37%.



Fig.S5 Synthesis route of 2-(Nap)-FFK_{TPA-DOX}AGLDD-RGD (compound 1).



Fig.S6 Synthesis route of 2-(Nap)-FFK_{TPA-DOX}WGLWD-RGD (compound 2).



Fig.S7 The HPLC spectra of DOX, 2-(Nap)-FFK_{TPA-DOX}AGLDD-RGD (compound 1) and 2-(Nap)-FFK_{TPA-DOX}WGLWD-RGD (compound 2).



Fig.S8 The HRMS spectrum of 2-(Nap)-FFK_{TPA-DOX}AGLDD-RGD (compound 1).



Fig.S9 The ¹H-NMR spectrum of 2-(Nap)-FFK_{TPA-DOX}AGLDD-RGD (compound 1, 400 MHz, *d6*-DMSO).



Fig.S10 The HRMS spectrum of 2-(Nap)-FFK $_{TPA-DOX}$ WGLWD-RGD (compound 2).



Fig.S11 The ¹H-NMR spectrum of 2-(Nap)-FFK_{TPA-DOX}WGLWD-RGD (compound 2, 400 MHz, *d6*-DMSO).



Fig.S12 The stability of RGD-sNPs and RGD-nNPs in PBS (pH7.4) containing 10% FBS at 37 °C for 48 h.



Fig.S13 The HRMS spectrum of intermediate generated from compound 1 that was treated for 12 h with MMP9, and the intermediate was determined as 2-(Nap) - $FFK_{TPA-DOX}AG$ according to the molecular weight, HRMS (ESI) m/z: ([M+H]⁺) calcd for C₇₆H₇₉N₇O₁₉,1394.4760; found 1394.5464.



Fig.S14 The chemical structure of 2-(Nap)-FFK_{TPA-DOX}AG.



Fig.S15 The CD spectra of RGD-nNPs with or without MMP9 treatment.



Fig.S16 Cellular uptake in HepG2 cells treated with RGD-nNPs. a) Confocal laser scanning microscope (CLSM) images of cellular uptake in HepG2 cells incubated with RGD-nNPs for 4 h or 24 h (white scale bar for 100 μ m and blue scale for 50 μ m), respectively. b) Quantitative analysis of DOX fluorescence intensity by flow cytometer and the amount of DOX internalized by HepG2 cells determined by HPLC (n=3).



Fig.S17 Bio-TEM of HepG2 cells incubated with pretreated RGS-sNPs (pretreated MMP9 for 12 h) for 2 h (red arrows represent nanofibers in vessels or lysosomes).



Fig.S18 Cellular uptake in 4T1 cells treated with RGD-sNPs or RGS-sNPs (pretreated MMP9 for 12 h) for 4 h and 24 h (white scale bar for 100 μ m and blue scale for 50 μ m), respectively.



Fig.S19 The expression levels of MMP9 in different tumors as analyzed by immunohistochemical section (scale bar for $200 \ \mu m$).



Fig.S20 The TEM images of Cy7.5-RGD-sNPs and Cy7.5-RGD-nNPs with or without MMP9 treatment.



Fig.S21 Bio-TEM of in situ formed nanofibers in tumor region (red arrows represent nanofibers formed in tumor).



Fig.S22 The average fluorescence intensity in tumor region at different time point (n=3).



Fig.S23 *In vivo* penetration of DOX extravasated from blood vessels after intravenous injection of DOX solution for 48 h at DOX dose of 2.5 mg/kg (scale bar for 100 μ m).



Fig.S24 *In vivo* penetration of DOX in tumors after intratumoral administration of DOX solution at DOX dose of 2.5 mg/kg for 48 h (scale bar for 100 μ m). Frozen sections of tumors were sliced at different depths below the injection position.



Fig.S25 The images of H&E for tumors resected from H22 tumor-bearing mice (scale bar for 200 μ m).



Fig.S26 The image of tumors (5/8) collected from 4T1 tumor-bearing mice after last treatment. The other three were stored at -80 °C after excision.



Fig.S27 The images of H&E for tumors resected from 4T1 tumor-bearing mice (scale bar for 100 μ m).



Fig.S28 The body weight change of H22 tumor-bearing mice (n=8).



Fig.S29 The body weight change of 4T1 tumor-bearing mice (n=8).



Fig.S30 The hematologic and biochemical parameters of plasma from H22 tumorbearing mice (n=8).

Table S1 The IC ₅	values against L	929 cell line after	er 48 h treatment ((n=6).
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Groups	IC ₅₀ values $(\mu g/mL)^{a}$	
	L929	
RGD-nNPs	504.25	
RGD-sNPs	493.44	

a) The half maximal inhibitory concentration.