Supporting Information

Improving the Carrier Stability and Drug Loading of Unimolecular Micelle-based Nanotherapeutic for Acidactived Drug Delivery and Enhanced Antitumor Therapy

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Figure S1. The synthetic route of rod-like DMO@DOX polymeric prodrug.



Figure S2. The ¹H NMR spectrum of DEX-Br macroinitiator.



Figure S3. The ¹H NMR spectrum of MGMA monomer.



Figure S4. GPC traces of DEX-PMGMA₂₄ and DEX-P(MGMA)₂₄-*b*-P(OEGMA)₃ (DMO-1), DEX-P(MGMA)₂₄-*b*-P(OEGMA)₈ (DMO-2) and DEX-P(MGMA)₂₄-*b*-P(OEGMA)₂₁ (DMO-3).



Figure S5. Typical FTIR spectra of DEX-Br, DEX-PMGMA, DEX-PMGMA-*b*-POEGMA (DMO), DMO-hydrazide and DMO@DOX, respectively.



Figure S6. UV-vis spectra of free DOX, DMO-1@DOX, DMO-2@DOX and DMO-3@DOX prodrugs in water or DMF solution.



Figure S7. The fluorescence spectra of free DOX, DMO-1@DOX, DMO-2@DOX and DMO-3@DOX prodrugs in water or DMF solution.



Figure S8. Fluorescence images of HeLa cells incubated with (A) DMO-2@DOX UMs and (B) DMO-3@DOX UMs for 24 h, respectively (final equivalent DOX concentration: 15 μ g/mL). The fluorescence of Calcein AM (live cells) and PI (dead cells) were labelled as green and red, respectively. Scale bars: 200 μ m.



Figure S9. (A) CLSM images showing the distribution of free DOX and DMO-2@DOX UMs in HeLa cells (final equivalent DOX concentration: 20 μ g/mL). Cell nuclei and cytoplasm were stained with DAPI (Blue) and AF-633 (Red), respectively. Scale bars: 50 μ m. (B) Flow cytometry analysis and (C) the mean fluorescence intensity of DMO-2@DOX treated HeLa cells at 2, 4 and 6 h (data are presented as means \pm SD, n =3).



Figure S10 (A) CLSM images showing the distribution of free DOX and DMO-3@DOX UMs in HeLa cells (final equivalent DOX concentration: 20 μ g/mL). Cell nuclei and cytoplasm were stained with DAPI (Blue) and AF-633 (Red), respectively. Scale bars: 50 μ m. (B) Flow cytometry analysis and (C) the mean fluorescence intensity of DMO-3@DOX treated HeLa cells at 2, 4 and 6 h (data are presented as means ± SD, n =3).



Figure S11 Flow cytometry dot plots for HeLa cells treated with (A) DMO-@DOX, (B) DMO-2@DOX and (C)DMO-3@DOX UMs under different incubation time.



Figure S12 CLSM images showing the time-dependent accumulation process of DMO-2@DOX UMs using Lyso Tracker as the tracer (final equivalent DOX concentration: 20 μg/mL). Scale bars: 50 μm.



Figure S13 CLSM images showing the time-dependent accumulation process of DMO-2@DOX UMs using Lyso Tracker as the tracer (final equivalent DOX concentration: $20 \ \mu g/mL$). Scale bars: $50 \ \mu m$.

	M n, NMR ^b	M n, GPC c	$M_{\rm w}/M_{\rm n}{}^d$	D^e	D^{f}	$LC^{g}(wt\%)$
Polymer ^{<i>a</i>}	(g mol ⁻¹)	(g mol ⁻¹)		(Water)	(DMF)	
DEX-P(MGMA)24	252900	38900	1.46		—	_
DEX-P(MGMA)24-b-P(OEGMA)3	339200	42900	1.56	34.83	45.89	80.4
DEX-P(MGMA)24-b-P(OEGMA)8	496900	48500	1.28	41.35	55.96	72.2
DEX-P(MGMA)24-b-P(OEGMA)21	912200	95500	1.24	68.01	74.23	40.4

Table S1. Structural information of the rod-like DMO@DOX prodrug and their precursors

^a DP_n of polymerized alternating units (MGMA or OEGMA) and ^b $M_{n,NMR}$ was measured by ¹H NMR results. ^c $M_{n,GPC}$ and ^d M_w / M_n were measured by GPC. ^eHydrodynamic diameter of prodrug micelles were determined by DLS. ^gThe loading content (LC) were determined by Fluorescence (FL) spectrophotometer, respectively.

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Sample	IC ₅₀ (HeLa)	IC ₅₀ (MCF-7)	IC ₅₀ (HUVEC)	
	µg/ml	µg/ml	µg/ml	
Free DOX	0.072	0.06	0.075	
DMO-1@DOX	1.61	0.77	>10	
DMO-2@DOX	0.65	1.75	>10	
DMO-3@DOX	0.83	1.91	>10	

Table S2.IC₅₀ value of free DOX and DMO@DOX prodrug.^a

a: IC₅₀ value of free DOX, DMO-1@DOX, DMO-2@DOX and DMO-3@DOX determined in HeLa cells, MCF-7 cells and HUVEC cells by PrestoBlue assay, and cells were incubated with samples for 72 h.