Supporting Information

Synthesis and self-assembly of hyperbranched multiarm copolymer peptide conjugates based on light-induced metal-free ATRP

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Characterization: ¹H NMR measurements were carried out on a 400 MHz Varian UNITY-plus NMR spectrometer. The M_n and the dispersity were measured by a Hitachi gel permeation chromatography (GPC) with DMF as the mobile phase and PMMA as the standards. The GPC was equipped with a Hitachi L-2490 refractive index detector and a Viscotek 270 dual detector. Scanning electron microscopy (SEM) was measured with an FEI Apreo S LoVac electron microscope. Transmission electron microscopy (TEM) was measured with a Tecnai G2 F20 electron microscope under 200 kV operating voltage. The hydrodynamic diameters (D_h), the polydispersities (PDI) and the Zeta potentials of the assemblies were measured by a Malvern Zetasizer Nano-ZS. The UV-vis absorbance spectra were collected on a Shimadzu UV-2450 spectrophotometer. Confocal images of the aggregates were obtained with a ZEISS LSM800 CLSM.

Table S1.1 orymetrization recipe of the hyperoranened porymers							
	DEGMA (mL)	BMA (g)	PTH (mg)	DMF (mL)	Time (h)		
h ₁ PDEGMA	2.70	0.4	4.4	3.1	11		
h ₂ PDEGMA	2.15	0.4	3.6	2.5	13		
h ₃ PDEGMA	1.35	0.4	2.4	1.7	12		

Table S1. Polymerization recipe of the hyperbranched polymers

Table S2. Po	lymerization	recipe o	of the hy	vperbranched	multiarm co	opolymers
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	hPDEGMA (g)	DSMA (g)	PTH (mg)	DMF (mL)
h1PDEGMA-star-PDSMA	0.15	0.233	1.3	1.5
h ₂ PDEGMA-star-PDSMA	0.15	0.328	1.7	1.5
h ₃ PDEGMA-star-PDSMA	0.15	0.179	1.0	1.5



Figure S1. ¹H NMR spectrum of PTH in CDCl₃.



Figure S2. ¹H NMR spectrum of DSMA in CDCl₃.



Figure S4. ¹H NMR spectrum of PDEGMA-*b*-PDSMA in CDCl₃.



Figure S5. ¹H NMR spectrum of BMA in CDCl₃.



Figure S6. ¹H NMR spectrum of h₁PDEGMA in CDCl₃.



Figure S7. ¹H NMR spectrum of h₂PDEGMA in CDCl₃.



Figure S8. ¹H NMR spectrum of h₃PDEGMA in CDCl₃.



Figure S9. GPC curves of h2PDEGMA (black line) and h2PDEGMA-star-PDSMA63 (red line).



Figure S10. GPC curves of h₃PDEGMA (black line) and h₃PDEGMA-star-PDSMA₅₆ (red line).



Figure S11. Fitting result of the GPC curve of h_1 PDEGMA. The M_n s of the two peaks are 12.8 and 23.7 kDa.



Figure S12. Fitting result of the GPC curve of h_2 PDEGMA. The M_n s of the two peaks are 12.6 and 28.8 kDa.



Figure S13. Fitting result of the GPC curve of h_3 PDEGMA. The M_n s of the two peaks are 12.0 and 22.6 kDa.



Figure S14. ¹H NMR spectrum of h₁PDEGMA-star-PDSMA₅₆ in CDCl₃.



Figure S15. ¹H NMR spectrum of h₂PDEGMA-star-PDSMA₆₃ in CDCl₃.



Figure S16. ¹H NMR spectrum of h₃PDEGMA-star-PDSMA₅₆ in CDCl₃.



Figure S17. Fitting result of the GPC curve of h_1 PDEGMA-*star*-PDSMA₅₆. The M_n s of the two peaks are 7.5 and 34.5 kDa.



Figure S18. Fitting result of the GPC curve of h_2 PDEGMA-*star*-PDSMA₆₃. The M_n s of the two peaks are 7.4 and 34.2 kDa.



Figure S19. Fitting result of the GPC curve of h_3 PDEGMA-*star*-PDSMA₅₆. The M_n s of the two peaks are 6.5 and 30.4 kDa.



Figure S20. UV-vis absorbance spectra of 2-mercaptopyridine in water as a byproduct of $h_1PDEGMA$ -*star*-PGS₅₆ (green line), $h_2PDEGMA$ -*star*-PGS₆₃ (red line), and $h_3PDEGMA$ -*star*-PGS₅₆ (black line).



Figure S21. UV-vis absorbance spectrum of 2-mercaptopyridine in water as a byproduct of h_1 PDEGMA-*star*-PT7₅₆.