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Confinements of Thermoresponsive Dendronized Polymers to Proteins

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Fig. S1 (A) Molecular structures of OEG-based dendronized polymethacrylate homopolymers **PG1** and dendronized chitosans (**DCS**). (B) Synthetic procedure for the copolymer **PG1S** through free radical copolymerization of **MS=S** with dendritic monomer **M3** in the presence of AIBN at 65 °C for 3 h in DMF.

sample	feed ratio ^a	actual copolymerization ratio ^b	GPC results ^c	
	[M3]/[MS=S]	[M3]/[MS=S]	<i>M</i> _n ×10 ⁻⁵	Ð
PG1	١	\	2.3	1.5
PG1S	20:1	21.5	2.2	2.3

Table S1 Conditions for and results from the polymerizations

^a Polymerizations were performed with AIBN as the initiator at 65 °C. ^b Copolymer compositions were determined through calculation according to proton integrations from ¹H NMR spectra in Fig. S2. ^c Determined by GPC with DMF as the eluent containing 0.1 wt% LiBr, M_n and \tilde{D} represent number-average molecular weight and polydispersity, respectively.



Fig. S2 ¹H NMR spectra of **PG1S** in DMSO- d_6 at 60 °C, C = 10 mg·mL⁻¹.



Fig. S3 (A) Absorbance of solutions for **PG1S** with Mb at 420 nm over reaction time. Heat effects by ITC in the titration of (B) **PG1** and (C) **DCS** to Mb in traditional mode at 25 °C.



Fig. S4 (A) Cloud point (T_{cp} s) for **PG1/proteins** mixture with different concentrations of proteins. [**PG1**] = 2.5 mg·mL⁻¹. (B) Plots of transmittance *vs* temperature for **DCS**, C = 2.5 mg·mL⁻¹. (C) Hydrodynamic radii (R_h , intensity radii) of **PG1/BSA**, **PG1/LYZ**, and **DCS** between 25 °C and 45 °C, respectively. C = 0.05 mg·mL⁻¹.



Fig. S5 R_h of (A) proteins and **PG1/proteins**, and (B) **PG1S-Mb**, **DCS/Mb**, and **DCS** at different temperatures. (C) Fluorescence spectrum of ANS (420–650 nm) in samples at 25 °C and 40 °C, respectively. PDI in Figure A and B represents the polydispersities of the aggregates.



Fig. S6 FT-IR spectra of (A) PG1S, PG1S-Mb, PG1, and PG1/Mb, at 25 °C, and (B) PG1S at different temperatures. All samples were prepared in D_2O .



Fig. S7 Changes of frequency Δf (red curves) and dissipation ΔD (blue curves) in QCM cell over time, after the injection of (A) Mb and PBS solution at 25 °C (below: schematic representation for immobilization process of proteins on QCM cells), and (B) **DCS** at 25 °C and 60 °C, respectively.



Fig. S8 Far-UV CD spectra of (A, C) BSA and (B, D) LYZ in presence of **PG1** with different concentrations and at different temperatures (polymer/protein = 40, w/w), respectively.



Fig. S9 Absorbance spectra of Mb from **PG1/Mb** (A) with different concentrations of polymer (polymer/protein, from 0 to 50, w/w) and (B) at different temperatures (polymer/protein = 40, w/w). (C) Maximum absorbance (A_{max}) at 410 nm of Mb from **PG1/Mb** solution after heating and cooling between 25 °C and 37 °C for 8 cycles. Inset in Figure B: photographs of the solutions below and above the T_{cp} .