

Supporting Information

Feedback-induced Phase Separation of Hollow Condensates to Create Biomimetic Membraneless Compartments

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Table of Contents

Experimental Procedures	3
Materials and reagents	3
Instrumental and characterization	3
Synthesis of sELP-ON conjugates	3
Preparation of sELP-ON condensates	3
Preparation of the observation chamber	3
Bright-field and confocal microscope imaging.....	3
Fluorescence recovery after photobleaching (FRAP).....	3
Surface-wetting	3
Droplet fusing	4
Dynamic light scattering (DLS) and zeta potential measurements	4
Transition temperature (T_i) measurement	4
Fluorescent labeling of proteins	4
Encapsulation of molecule dyes and biomacromolecules	4
sELP-ON coacervate droplets for GOx-HRP cascade.....	4
Supplementary Figures S1-S14 and Tables S1-S2	5
Supplementary References.....	11

Experimental Procedures

Materials and reagents

Short elastin-like polypeptides were purchased from ChinaPeptides Co., Ltd. Oligonucleotides, acryl/bis 30% solution (29:1), TBE premixed powder were (10×), Tris-HCl solution (2 M, pH=7.8), ethylene diamine tetraacetic acid solution (EDTA, 0.5 M, pH=8.0) and bovine serum albumin (BSA, 98.0%) were purchased from Sangon Biotech (Shanghai) Co., Ltd. Sodium L-ascorbate (99.0%), copper(II) sulfate pentahydrate ($\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$, 99.9%), 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulfonic acid) (ABTS, 99.0%), glucose oxidase (GOx, 95.0%) and horseradish peroxidase (HRP, 95.0%) were purchased from J&K China Chemical Co. Ltd. 1,6-hexanediol (98.0%), urea (99.0%), rhodamine 6G (95.0%), rhodamine B (95.0%), fluorescein (98.0%) and carboxyfluorescein (98.0%) were purchased from Shanghai Titan Scientific Co., Ltd. Bovine hemoglobin (Hb, 99.5%) was purchased from Shanghai yuanye Bio-Technology Co., Ltd. Sodium chloride (NaCl, 99.9%), magnesium chloride hexahydrate ($\text{MgCl}_2 \cdot 6\text{H}_2\text{O}$, 99%) and Nile red (98.0%) were purchased from Aladdin Industrial Corporation. Sulforhodamine B (95.0%), bodipy 493/503 (95.0%) and pyranine (90.0%) were purchased from Shanghai Canspec Scientific Instruments Co., Ltd. Calcein (95.0%), 5-fluorescein isothiocyanate (FITC, 90.0%) and rhodamine B 5-isothiocyanate (RBITC, 95.0%) were purchased from Dibai Chemical Co., Ltd. Lucifer yellow cadaverine (L. Yellow, 95.0%) was purchased from Bidepharm Co., Ltd. Porcine insulin (98.8%) was purchased from Xuzhou Wanbang Jinqiao Pharmaceutical Co., Ltd. (Jiangsu, China). FITC-Dextran (40 kDa, 95.0%) was purchased from JiQi Biotech Co., Ltd. N,N,N',N'-tetramethylethylenediamine (TEMED, 98.0%), ammonium persulfate (APS, 98.0%) and lysozyme (95.0%) were purchased from Shanghai Macklin Biochemical Co., Ltd. Porcine liver esterase (PLE, 15 U/mg) and FITC-IgG were purchased from Thermo Fisher Scientific Inc. Gelred (10000×) were purchased from Beyotime Biotechnology. All aqueous solutions were prepared with ultrapure water (18.25 MΩ·cm) supplied by Milli-Q Advantage A10 (Merck Millipore). All chemicals were used as received without further purification.

Instrumental and characterization

The concentration of oligonucleotides and sELP-ON conjugates were measured using a Thermo Fisher Nanodrop 2000. Denatured polyacrylamide gel electrophoresis (PAGE) was carried out in 1×TBE buffer (89 mM Tris-borate, 2 mM EDTA, pH=8.2-8.4) at 120 V (constant voltage) for 1 h using a Bio-Rad Mini-PROTEAN® Tetra Vertical Electrophoresis Cell and visualized using DLAB GelSMART gel imaging system. UV-vis absorption and absorbance measurements were carried out with a Shimadzu UV-visible 2600 spectrometer. Dynamic light scattering (DLS) and zeta potential measurements were performed using a Malvern Zetasizer Pro instrument using a He-Ne laser (the wavelength $\lambda = 633$ nm, 4 mW) as the light source with scattering angle of 175°. The real-time fluorescence and absorbance of enzymatic reactions were measured in a BioTek Synergy NEO microplate reader. Confocal fluorescence microscope measurements were performed using a Nikon A1R inverted laser-scanning confocal microscope equipped with a 60× water objective. The microscope slide glasses (26 mm × 76 mm) and cover glasses (18 mm × 18 mm × 0.17 mm) were purchased from Beyotime Biotechnology.

Synthesis of sELP-ON conjugates

sELP-ON conjugates were synthesized as previously described.^{1,2} CuSO_4 solution (1 mM, 5 μL) and sodium L-ascorbate solution (2 mM, 5 μL) were added into a 1.5 mL Eppendorf tube and incubated for 3 min. And then, 50% (v/v) DMSO aqueous solution (200 μL), alkynyl-sELP solution (10 mg/mL, 50 μL , dissolved in DMSO), and azido-oligonucleotides solution (100 μM , 50 μL) were added into the tube. The mixed solution was stirred at 37 °C for 6 hours under nitrogen. After reaction, sELP-ON conjugates were purified using 10% denaturing polyacrylamide gel electrophoresis (PAGE) and concentrated using Amicon filtering tubes 3 kDa (Amicon bioseparations-Millipore U.S.A.). sELP-ON conjugates were stored in Tris-HCl solution (pH=7.8, 20 mM) at -40 °C.

Preparation of sELP-ON condensates

The sELP-ON conjugate was concentrated to a final concentration of 30 μM using ultrafiltration, which underwent liquid-liquid phase separation (LLPS) to the formation of sELP-ON microdroplets. To facilitate the subsequent hollow condensate formation, a gradual addition of NaOH solution (0.1 M) was employed to elevate the pH value of the solution and induce the phase transition from microdroplets to hollow condensates.

Preparation of the observation chamber

The observation chambers were prepared as previously described.³ The microscope glasses were soaked in 5% (w/v) bovine serum albumin (BSA) solution for over 4 hours to prevent surface-wetting of the sELP-ON coacervates on the glass surface before being used. After the BSA coating, the glasses were washed with distilled water and dried under an airflow. The BSA-coated slide glasses and cover slips were assembled using double-sided tape. The sample solutions were immersed into the slit between the slide glass and the cover slips, and the edges were sealed with Devcon 5 Minute Epoxy to avoid evaporation of the sample solution during the observation period.

Bright-field and confocal microscope imaging

Samples in the observation chamber were visualized using a confocal laser scanning microscope equipped with a 60× water objective and a stage heater at 37 °C. Blue channel, excitation: 405 nm, collection: 425-475 nm; green channel, excitation: 488 nm, collection: 500-550 nm; red channel, excitation: 561 nm, collection: 570-620 nm. Three-dimensional reconstructions from Z scans were performed using NIS-Viewer software.

Fluorescence recovery after photobleaching (FRAP)

For FRAP measurements, the sELP-ON coacervate droplets were stained by Gelred probe. Images at attenuated laser intensity (10% intensity) were taken before photobleaching. The photobleached area was chosen in the center of a single droplet of at least 4 μm in diameter. A 2-frame pre-bleach sequence was used, followed by a 2-frame bleach at 100% 561 nm. After the area's fluorescence emission intensity was significantly decreased, the laser was then switched back to the attenuated intensity and took a frame every 10 s for at least 180 s.

Surface-wetting

For Surface-wetting measurements, the microscope glasses were not treated with BSA. Surface-wetting imaging was performed immediately after adding the sample solution into the observation chamber. The suspended droplets were selected for surface-wetting imaging and took a frame every 10 s for 300 s.

Droplet fusing

For droplet fusing measurements, the touching sELP-ON coacervate droplets were chosen for fusing imaging and took a frame every 10 s for at least 300 s.

Dynamic light scattering (DLS) and zeta potential measurements

sELP-ON coacervate samples (20 μ M) were injected into a disposable zeta cuvette to measure the hydration dynamic diameters and zeta potentials at 37 °C. Samples were pre-incubated at 37 °C for at least 120 s before measurements and the reported data represent an average of at least three measurements.

Transition temperature (T_c) measurement

The sELP-ON conjugate samples were heated until phase separation before measurement. Immediately, the absorbance at 350 nm of sELP-ON conjugate samples was monitored using a UV-visible spectrometer equipped with a temperature sensor during cooling. The absorbance at each temperature was recorded. Origin software was used to analyze data and draw graphs. For NaCl concentration and pH effect test, sELP-ON conjugates were dissolved in buffer solution (20 mM Tris-HCl, pH=7.8), and the NaCl stock solution (5 M) was used to adjust NaCl concentration, HCl (1 M) and NaOH (1 M) were used to adjust pH.

Fluorescent labeling of proteins

Proteins were labeled as previously described.⁴ FITC or RBITC (10 mM, 20 μ L, dissolved in DMSO) was added into proteins solution (30 μ M, 1000 μ L, dissolved in 0.1 M NaHCO₃-Na₂CO₃, pH=9.5), the mixed solution was stirred at 37 °C for 4 hours and incubated at 4 °C for 8 hours under dark. And then, Amicon filtering tubes were used to remove excess FITC or RBITC.

Encapsulation of molecule dyes and biomacromolecules

Before imaging, sELP-ON coacervate droplets were incubated with molecule dyes or biomacromolecules for 2 hours at 37 °C. The concentrations of all molecule dyes were 2 μ M. The concentrations of all proteins were 0.2 μ M. The concentration of FITC-Dextran was 33 mg/L. The concentrations of nucleic acids were 0.5 μ M. For two-component uptake experiments, the concentrations of molecule dyes were 1 μ M and the concentrations of proteins were 0.1 μ M.

sELP-ON coacervate droplets for GOx-HRP cascade

sELP-ON coacervate droplets were pre-incubated with 2.5 μ M GOx/HRP for 2 hours at 37 °C. And then, 20 mM L-Arg and 40 mM glucose were added to coacervate samples. For confocal imaging, the fluorescence images were taken every 1 minute for a total of 30 minutes.

Supplementary Figures S1-S14 and Tables S1-S2

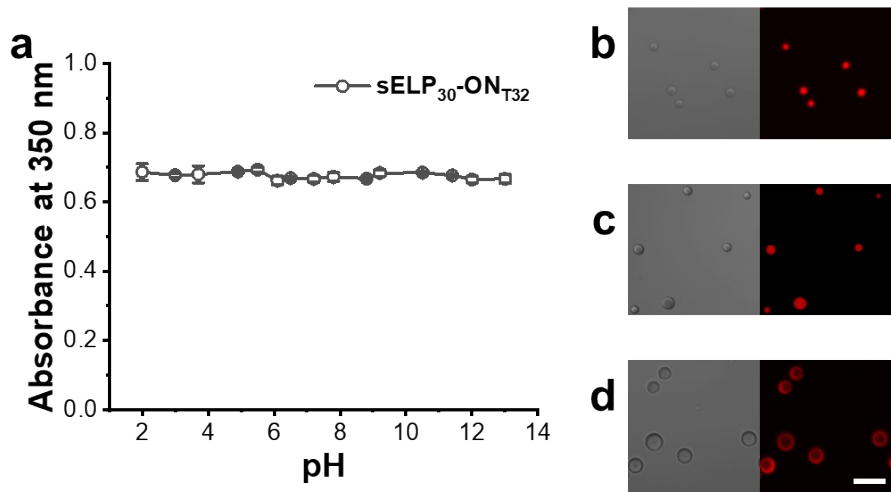


Figure S1. (a) Absorbance profile at 350 nm of the sELP₃₀-ON_{T32} by varying pH from 2 to 13 under 25 °C. DIC and fluorescence micrographs of Gelred-labeled sELP₃₀-ON_{T32} condensate structures under 25 °C at (b) pH 2, (c) pH 7.8, and (d) pH 13. Scale bar: 10 μm.

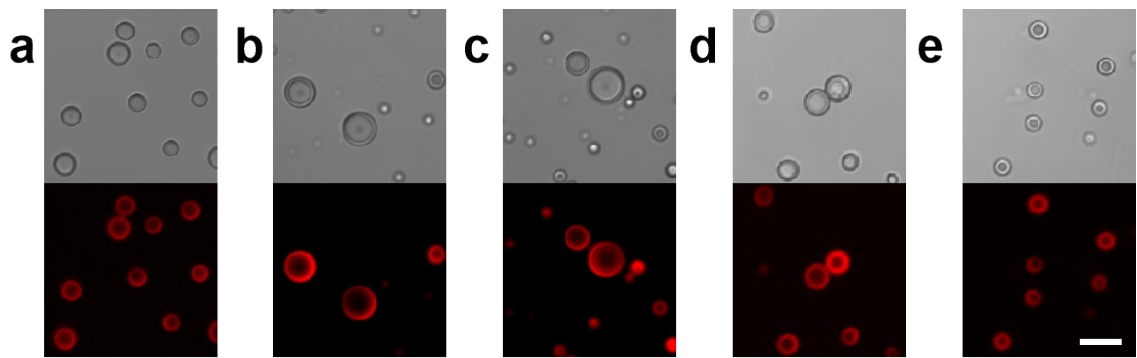


Figure S2. DIC and fluorescence images of Gelred-labelled sELP-ON hollow condensate under 25 °C at pH 13. (a) sELP₃₀-ON_{A32}, (b) sELP₃₀-ON_{C32}, (c) sELP₃₀-ON₃₂, (d) sELP₃₀-ON_{T22} and (e) sELP₂₅-ON_{T22}. Scale bar: 10 μm.

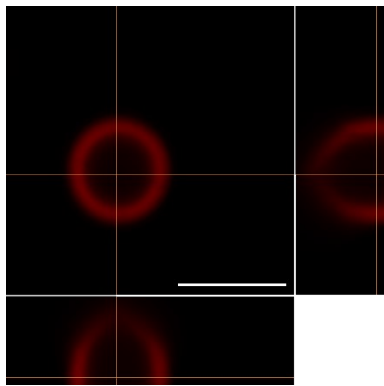


Figure S3. Reconstructed 3D fluorescence images of a Gelred-labelled sELP₃₀-ON_{T22} hollow condensate. Scale bar: 10 μm.

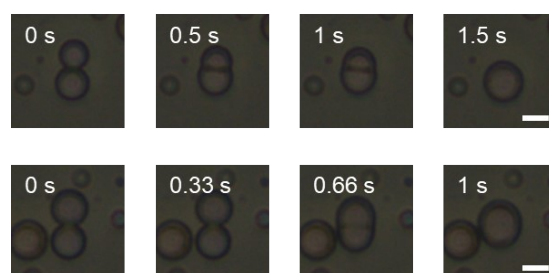


Figure S4. Time-lapsed bright-field images of fusion measurement of two adhering sELP₃₀-ON_{T22} vesicle-like hollow condensates. Scale bar: 5 μ m.

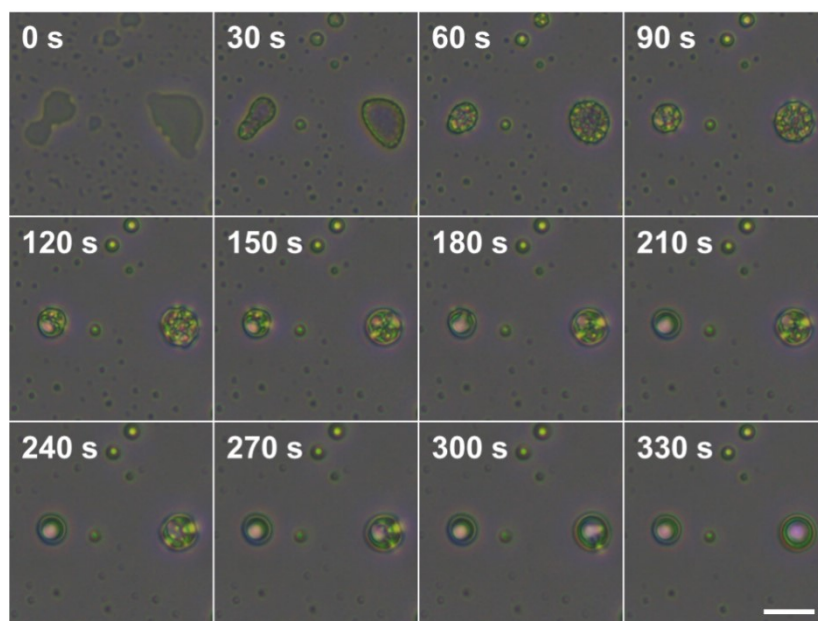


Figure S5. Time-lapsed bright-field images of wetted sELP₃₀-ON_{T22} microdroplets transforming into unwetted hollow condensates after adjusting the pH to 13 using NaOH. Scale bar: 10 μ m.

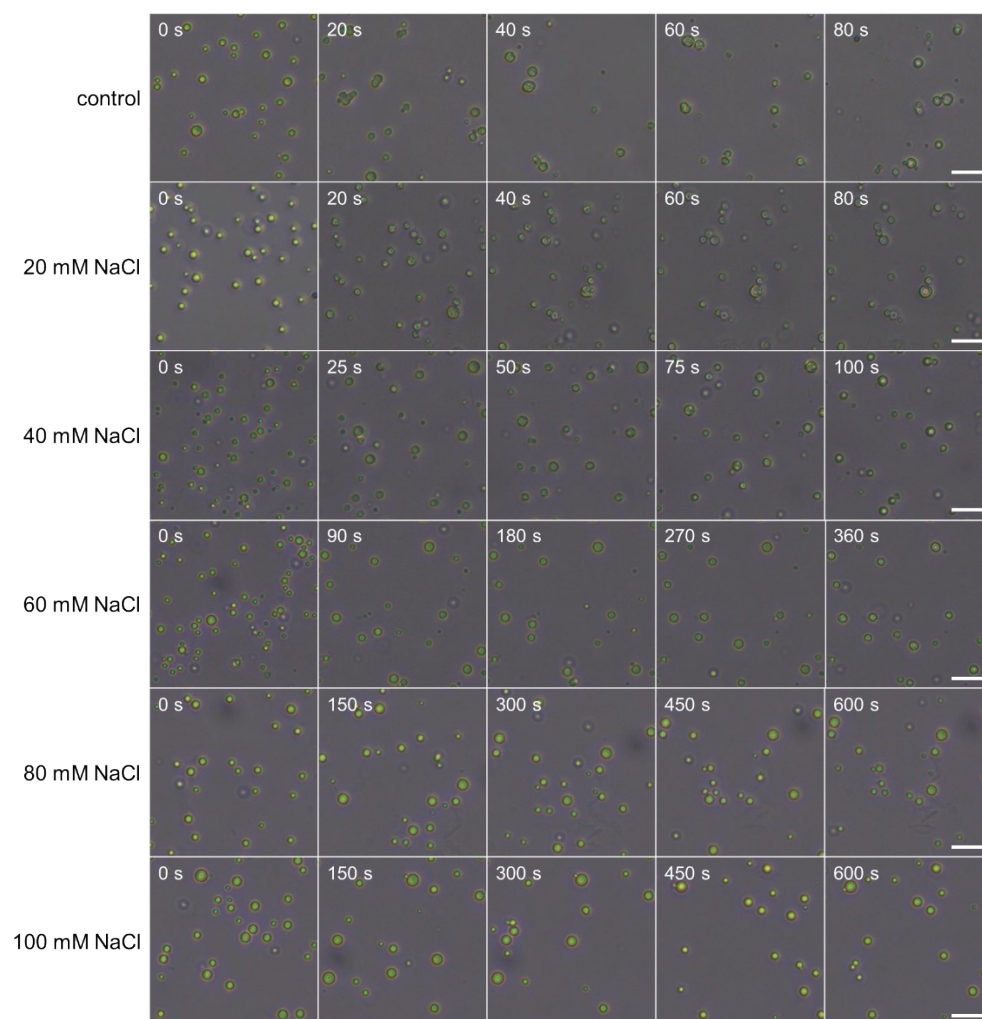


Figure S6. Time-lapsed bright-field images of sELP₃₀-ON_{T22} condensates after the addition of NaOH in the presence of 0-100 mM NaCl. Scale bar: 10 μ m.

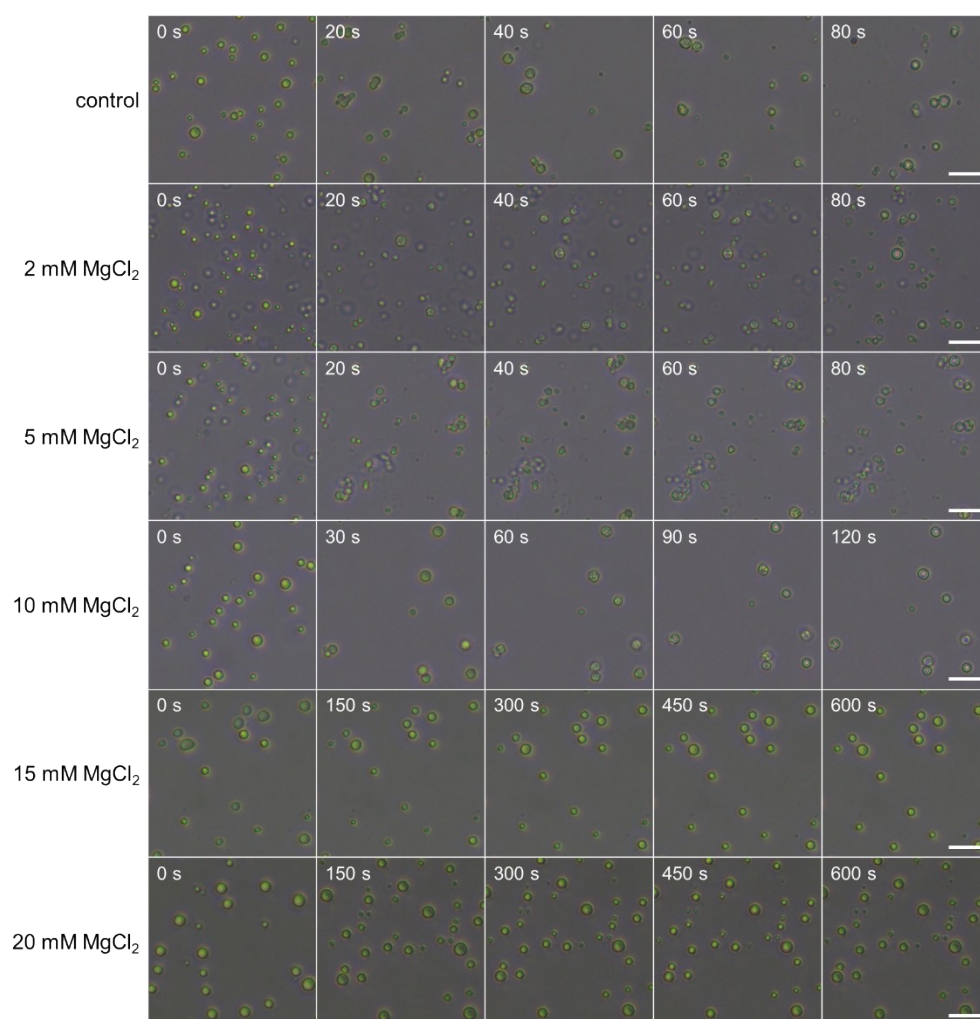


Figure S7. Time-lapsed bright-field images of sELP₃₀-ON₇₂₂ condensates after the addition of NaOH in the presence of 0-20 mM MgCl₂. Scale bar: 10 μ m.

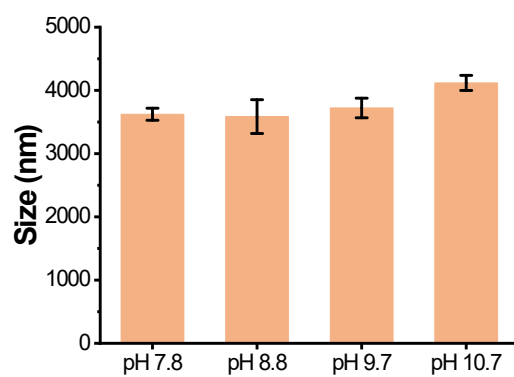


Figure S8. Hydrodynamic diameters of sELP₃₀-ON₇₂₂ condensates in the presence of 20 mM NaCl under 37 °C at pH 7.8, 8.8, 9.7, and 10.7 respectively.

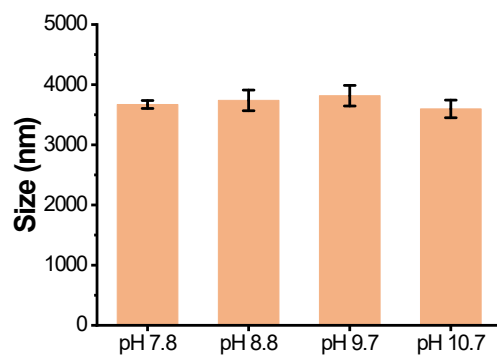


Figure S9. Hydrodynamic diameters of sELP₃₀-ONT₂₂ condensates in the presence of 20 mM MgCl₂ under 37 °C at pH 7.8, 8.8, 9.7, and 10.7 respectively.

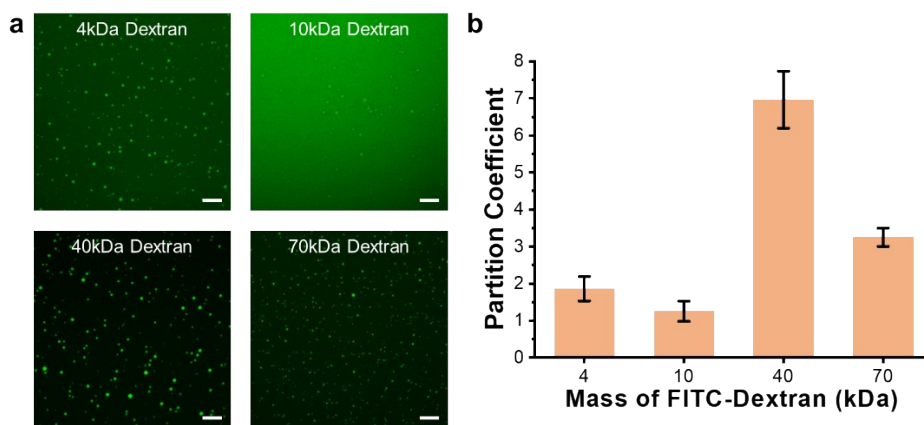


Figure S10. (a) CLSM images of sELP₃₀-ONT₂₂ microdroplets incubated with 4 kDa, 10 kDa, 40 kDa, and 70 kDa FITC-dextran. Scale bar: 20 μ m. (b) Partition coefficient of 4 kDa, 10 kDa, 40 kDa, and 70 kDa FITC-dextran in sELP₃₀-ONT₂₂ microdroplets. At least 300 microdroplets were calculated by imageJ software.

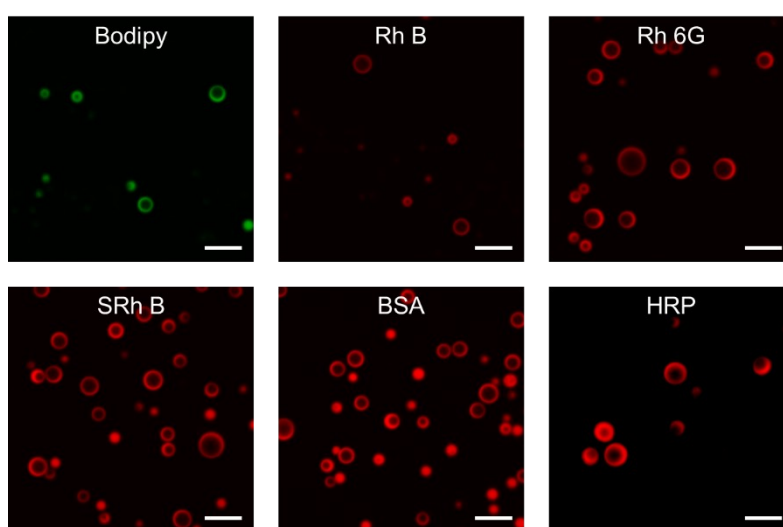


Figure S11. CLSM images of sELP₃₀-ONT₂₂ hollow condensates transformed by sELP₃₀-ONT₂₂ microdroplets preloaded with Bodipy, RhB, Rh 6G, SRh B, BSA, and HRP after the addition of NaOH. Scale bar: 10 μ m.

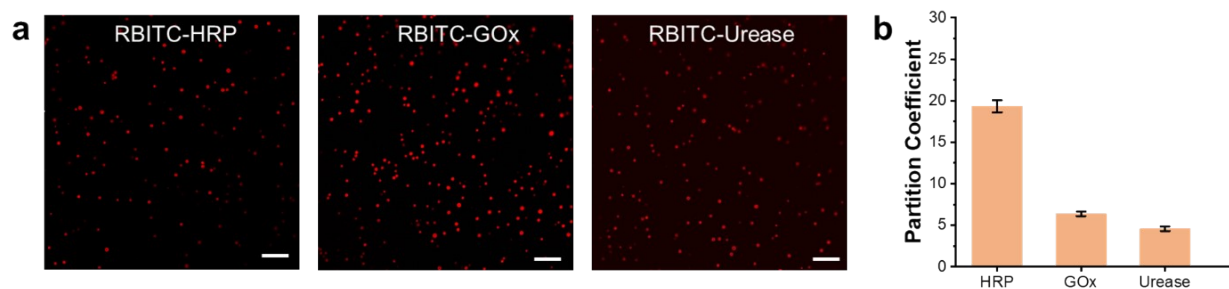


Figure S12. (a) CLSM images of sELP₃₀-ONT₂₂ microdroplets incubated with RBITC-HRP, RBITC-GOx, and RBITC-Urease, respectively. Scale bar: 20 μm. (b) Partition coefficient of RBITC-HRP, RBITC-GOx, and RBITC-Urease in sELP₃₀-ONT₂₂ microdroplets. At least 300 microdroplets were calculated by imageJ software.

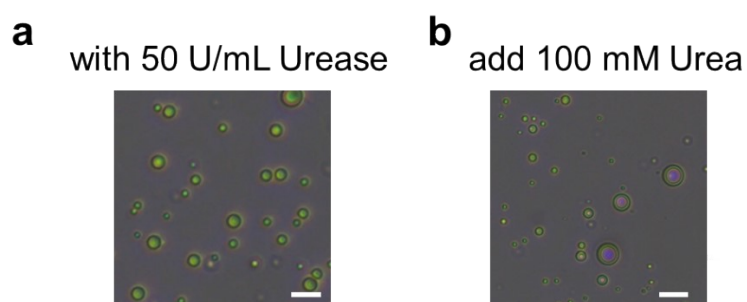


Figure S13. Bright-field images of sELP₃₀-ONT₂₂ condensates after (a) the addition of 50 U/mL urease and (b) the following addition of 100 mM urea. Scale bar: 10 μm.

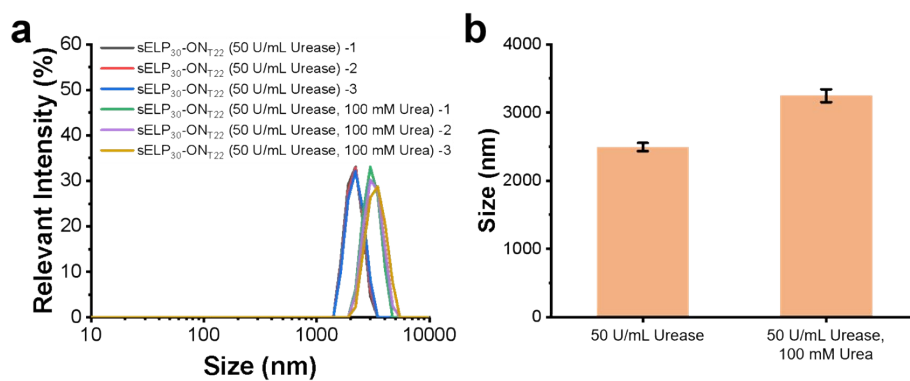


Figure S14. (a) Three repeated results of dynamic laser scattering analysis of sELP₃₀-ONT₂₂ microdroplets containing 100 U/mL urease before and after the addition of 100 mM urea. (b) The corresponding hydrodynamic diameters of sELP₃₀-ONT₂₂ microdroplets containing 50 U/mL urease before and after addition of 100 mM urea. The error bars indicated three repeated experiments.

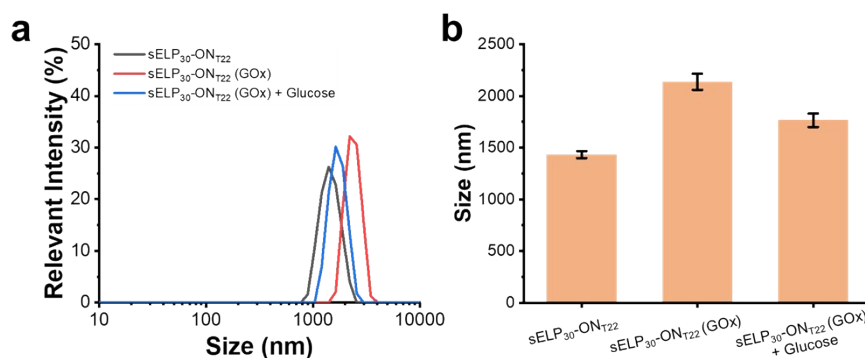


Figure S15. (a) Dynamic laser scattering analysis of sELP₃₀-ONT₂₂ microdroplets containing 2.5 μ M GOx before and after the addition of 40 mM glucose. (b) The corresponding hydrodynamic diameters of sELP₃₀-ONT₂₂ microdroplets containing 2.5 μ M GOx before and after addition of 40 mM glucose. The error bars indicated three repeated experiments.

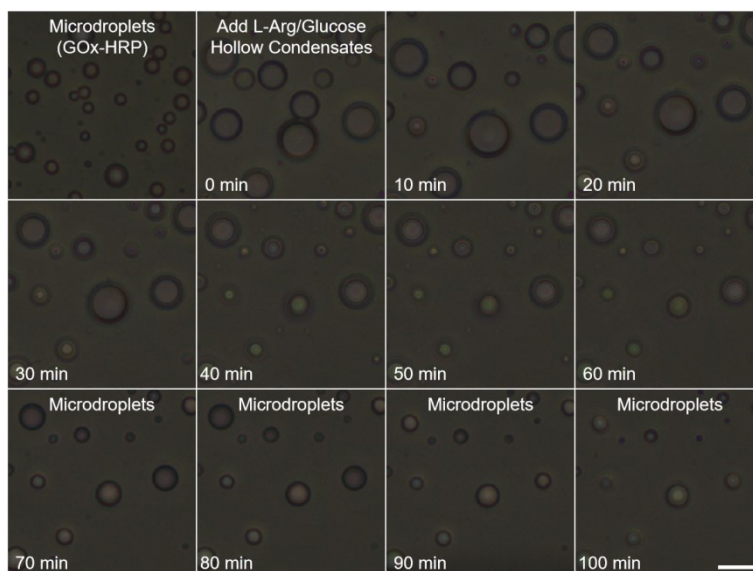


Figure S16. Time-lapsed bright-field images of sELP₃₀-ONT₂₂ microdroplets preloaded with 2.5 μ M GOx-HRP rapidly transforming into hollow condensates and gradually returning to microdroplets after the addition of 20 mM L-Arg and 40 mM glucose. Scale bar: 10 μ m.

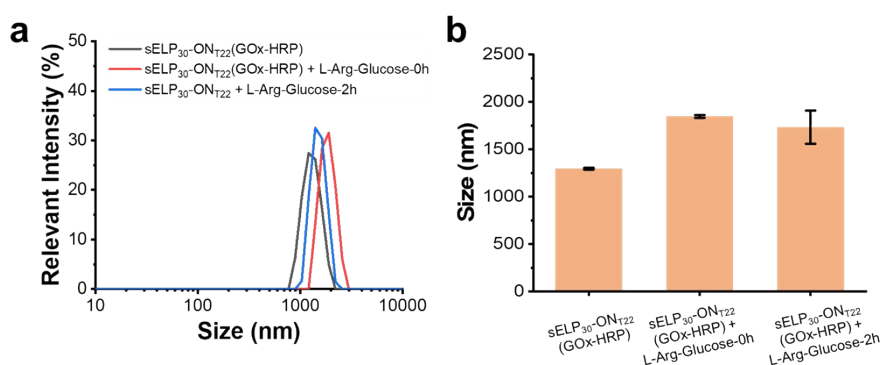


Figure S17. (a) Dynamic laser scattering analysis of sELP₃₀-ONT₂₂ microdroplets containing 2.5 μ M GOx/HRP before and after the addition of 20 mM L-Arg and 40 mM glucose. (b) The corresponding hydrodynamic diameters of sELP₃₀-ONT₂₂ microdroplets containing 2.5 μ M GOx/HRP before and after addition of 20 mM L-Arg and 40 mM glucose. The error bars indicated three repeated experiments.

Table S1. Elastin-like polypeptides sequences used in this study.

sELP	Sequence (N'-C')
sELP ₃₀	VPFGFVPFGFVPFGFVPFGFVPFGFVPFGF-(L-Pra ^[a])-NH ₂
sELP ₂₅	VPFGFVPFGFVPFGFVPFGFVPFGFVPFGF-(L-Pra)-NH ₂

[a] L-Propargylglycine.

Table S2. Oligonucleotides polypeptides sequences used in this study.

Oligonucleotide	Sequence (5'-3')	Base number
Azido-ON _{T32}	Azido-TTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	32
Azido-ON _{A32}	Azido-AAAAAAAAAAAAAAAAAAAAAAAAAAAA	32
Azido-ON _{C32}	Azido-CCCCCCCCCCCCCCCCCCCCCCCCCCCC	32
Azido-ON ₃₂	Azido-TCCCCTATAGTGAGTCGTATTAATTTTCGCGGG	32
Azido-ON _{T22}	Azido-TTTTTTTTTTTTTTTTTTTTTT	22

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- 2 S. Yao, Y. Liao, R. Pan, W. Zhu, Y. Xu, Y. Yang and X. Qian, *Chinese Chem. Lett.* 2021. DOI: 10.1016/j.cclet.2021.08.116.
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- 4 W. Mu, Z. Ji, M. Zhou, J. Wu, Y. Lin and Y. Qiao, *Sci. Adv.* 2021, **7**, eabf9000.