

Supplementary Information

Morphology Under Control: Engineering Biodegradable Stomatocytes

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Experimental Materials and Methods

Materials

All chemicals were used as received unless otherwise stated. For the synthesis of poly(ethylene glycol)-*b*-poly(D,L-lactide) (PEG-PDLLA), Poly(ethylene glycol) 1K and 2K were purchased from JenKem technology and freeze-dried before use. D,L-lactide was purchased from Acros and used as supplied. All other chemicals were supplied by Sigma-Aldrich and were used without any purification. Ultra pure MilliQ water obtained from Merck Millipore Q-Pod system (18.2 MΩ) with a 0.22 μm Millipore® Express 40 filter was used for the polymersome self-assembly and their dialysis. Dialysis Membranes MWCO 12-14000 g mol⁻¹ Spectra/Por® were used for dialysis during the stomatocyte formation. Sodium chloride was purchased from Merck.

Methods

Nuclear Magnetic Resonance (NMR): Proton nuclear magnetic resonance measurements were performed on a Bruker 400 Ultrashield™ spectrometer equipped with a Bruker SampleCase autosampler, using CDCl₃ as a solvent and TMS as internal standard.

Gel permeation chromatography (GPC): GPC was conducted using a Shimadzu Prominence-*i* GPC system with a PL gel 5 μm mixed D and mixed C column (Polymer Laboratories) with PS standard and equipped with a Shimadzu RID-20A differential refractive index detector and THF used as an eluent with a flow rate of 1 mL min⁻¹.

Differential scanning calorimetry (DSC): DSC measurements were performed using TA Instruments Multicell DSC.

Dynamic Light Scattering (DLS): DLS measurements were performed on a Malvern instrument Zetasizer (model Nano ZSP) equipped with an autosampler. Zetasizer software was used to process and analyse the data.

Cryogenic transmission electron microscopy (cryo-TEM): Experiments were performed using a FEI Tecnai G2 Sphere (200 kV electron source) equipped with LaB₆ filament utilizing a cryoholder or a FEI Titan (300 kV electron source) equipped with autoloader station. Samples for cryo-TEM were prepared by treating the grids (Lacey carbon coated, R2/2, Cu, 200 mesh, EM sciences) in a Cressington 208 carbon coater for 40 seconds. Then, 3 μl of the polymersome solution was pipetted on the grid and blotted in a Vitrobot MARK III at 100% humidity. The grid was blotted for 3 seconds (offset -3) and directly plunged and frozen in liquid ethane. Processing of TEM images was performed

with ImageJ, a program developed by NIH and available as public domain software at <http://rsbweb.nih.gov/ij/>.

Asymmetric Flow Field-Flow Fractionation and multi-angle light scattering (AF4-MALS): The asymmetric flow field-flow fractionation – UV – Quels (AF4-UV-Quels) experiments were performed on a Wyatt Dualtec AF4 instrument connected to a Shimadzu LC-2030 Prominence-*i* system with Shimadzu LC-2030 autosampler. The AF4 was connected to a Wyatt DAWN HELEOS II light scattering detector (MALS) installed at different angles (12.9 °, 20.6 °, 29.6 °, 37.4 °, 44.8 °, 53.0 °, 61.1 °, 70.1 °, 80.1 °, 90.0 °, 99.9 °, 109.9 °, 120.1 °, 130.5 °, 149.1 °, and 157.8 °) using a laser operating at 664.5 nm and a Wyatt Optilab Rex refractive index detector. Detectors were normalized using Bovine Serum Albumin (BSA). The processing and analysis of the LS data and radius of gyration (R_G) calculations were performed on Astra 7 software (using the Berry model, which is recommended for particles of size > 50 nm). All AF4 fractionations were performed on an AF4 short channel with regenerated cellulose (RC) 10 KDa membrane (Millipore) and spacer of 350 μm .

Experimental Procedures

Ring opening polymerization (ROP) of poly(ethylene)glycol-poly(D,L-lactide) PEG-PDLLA block copolymers: PEG_n-PDLLA_m was synthesized by ring opening polymerization starting from PEG-macro initiators and using DBU as a catalyst as described previously¹ yielding PEG_n-PDLLA_m polymers with 80 % yield. Copolymer composition was calculated by comparing the protons of PEG (3.65-3.7 ppm) and its terminal methyl unit (singlet at 3.37 ppm) to PDLLA CH (multiplet at 1.4-1.8 and 5.11-5.28 ppm). All PDI values were calculated to be less than 1.2 using PS standards for calibration. Differential Scanning Calorimetry measurements were performed to determine glass transition temperatures (T_g) by scanning from -20 °C to 60 °C at a rate of 0.5 °C/min. and 3 heating/cooling steps. The average polymer compositions and values for PDI and T_g are given in Table S1.

Preparation of polymersomes: In a 15 mL vial, PEG₂₂-PDLLA₉₅ and PEG₄₄-PDLLA₉₅ block copolymers (1:1 w/w, 20 mg) were dissolved in 2 mL of THF and dioxane (1:4 v/v) and the vial was sealed with a rubber septum. The solution was stirred at 700 rpm for a minimum of 30 minutes prior to the addition of MilliQ (2 mL, 1 mL h⁻¹) *via* a syringe pump. A needle was inserted into the septum to release pressure. The resulted cloudy suspension was transferred into a prehydrated dialysis bag (SpectraPor, MWCO: 12-14 kDa, 2 mL/cm). Dialysis was performed against MilliQ at room temperature for max. 24 hours with a water change after 1 hour.

Preparation of discs: Polymersomes were prepared according to the abovementioned procedure. Afterwards, dialysis was performed against MilliQ at 5 °C for max. 24 hours with a water change after 1 hour.

General method for stomatocytes preparation: First, a batch of polymersomes was prepared following the previously mentioned procedure. Polymersomes were dialyzed against pre-cooled salt solution (50 mM NaCl, unless stated otherwise) at 5 °C for max. 24 hours.

Reverse engineering of stomatocytes into polymersomes: Stomatocytes were prepared according to above described procedure. Stomatocytes were transferred into a prehydrated dialysis bag. Stomatocytes were dialyzed against 50 vol % organic solvent (THF/dioxane (1:4)). Aliquots of 250 μL were withdrawn every hour and quenched with water to a total volume of 2 mL. Samples were directly used for cryo-TEM imaging and DLS.

Supplementary Figures and Tables

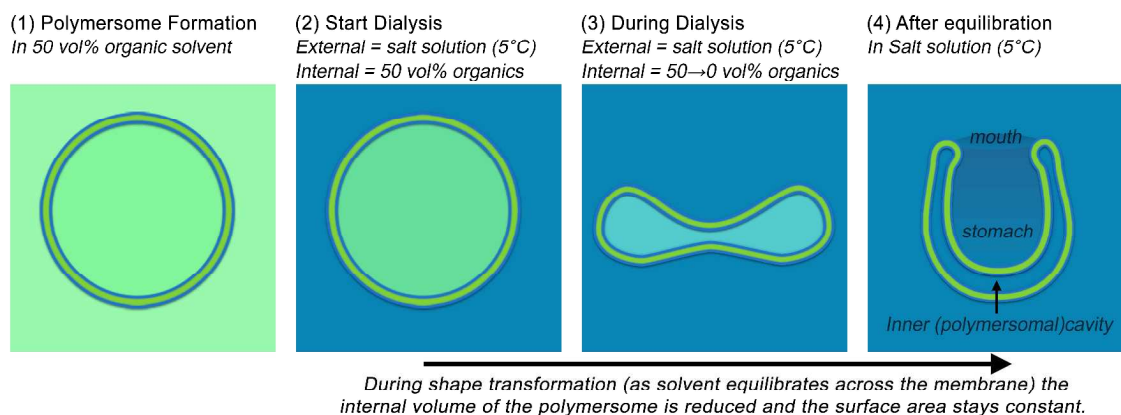


Figure S1: Description of the shape transformation process utilized in this work, whereby polymersomes (having been fabricated in 50 vol% of organic solvent) are dialyzed against salt solution at 5 °C. During the early stages of dialysis, as organic solvent re-equilibrates across the polymersomal membrane, shape transformation occurs as a result of osmotically-induced volume reduction and consequent folding of the membrane. The color scheme here is intended to highlight the solvent composition inside and outside of the polymersome *inner* cavity during this process. Shape transformation of spherical into bowl-shaped (stomatocytes) polymersomes proceeds *via* an oblate pathway as a result of induced *negative* C_0 .²

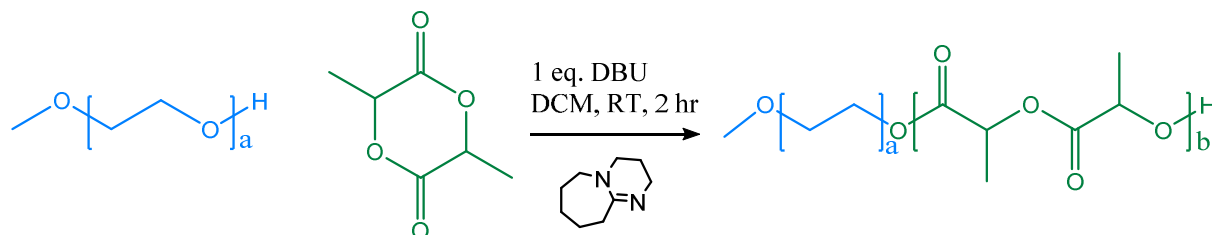


Figure S2: Organocatalyzed synthesis of block copolymer PEG_a-PDLLA_b with a=22/44 and b=45/70/95/120.

Polymer	Average polymer composition (¹ H-NMR)	PDI (GPC)	T _g (DSC)
PEG ₂₂ -PDLLA ₉₅	PEG ₂₂ -PDLLA ₉₈	1.17	29 °C
PEG ₄₄ -PDLLA ₉₅	PEG ₄₄ -PDLLA ₉₅	1.11	26 °C
PEG ₂₂ -PDLLA ₇₀	PEG ₂₂ -PDLLA ₇₂	1.09	32 °C
PEG ₄₄ -PDLLA ₇₀	PEG ₄₄ -PDLLA ₆₈	1.15	36 °C
PEG ₂₂ -PDLLA ₁₂₀	PEG ₂₂ -PDLLA ₁₂₅	1.06	35 °C
PEG ₄₄ -PDLLA ₁₂₀	PEG ₄₄ -PDLLA ₁₂₁	1.07	21 °C

Table S1: Characterization of PEG_a-PDLLA_b block copolymers.

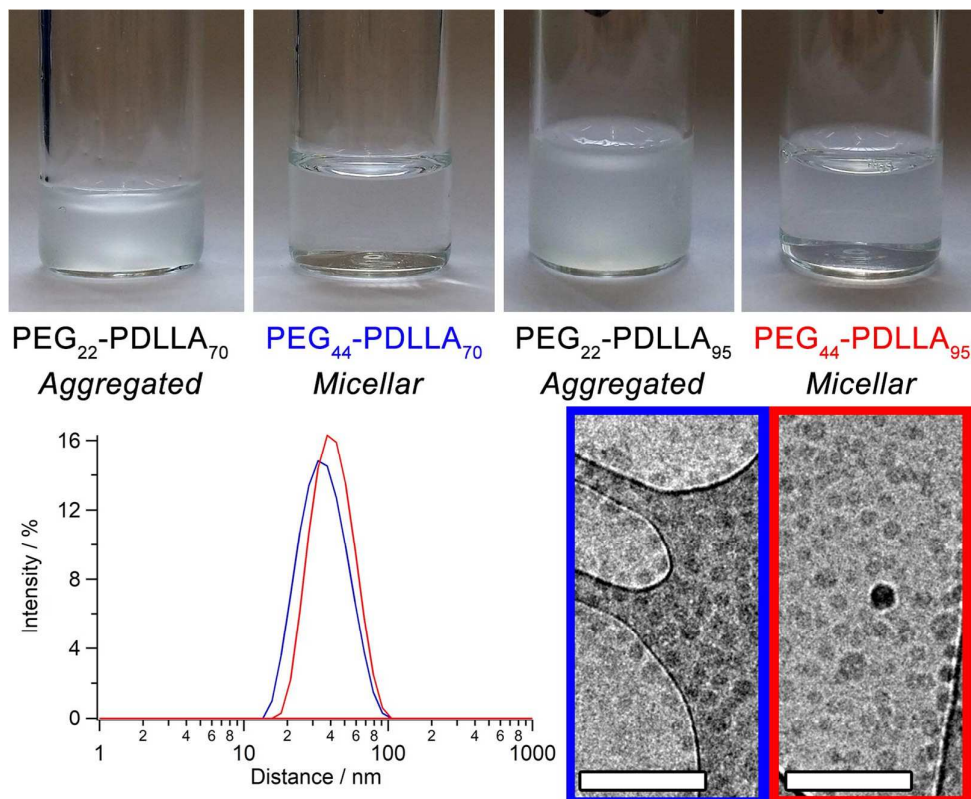


Figure S3: (Top) Photographs showing the outcome of the self-assembly of PDLLA₇₀ and PDLLA₉₅ based copolymers. (Bottom) DLS intensity profile and cryo-TEM images confirming micelles formation from PEG₄₄-PDLLA₇₀ (blue) and PEG₄₄-PDLLA₉₅ (red) with average hydrodynamic diameters (D_h) of 39 and 47 nm, respectively. Scale bars correspond to 200 nm.

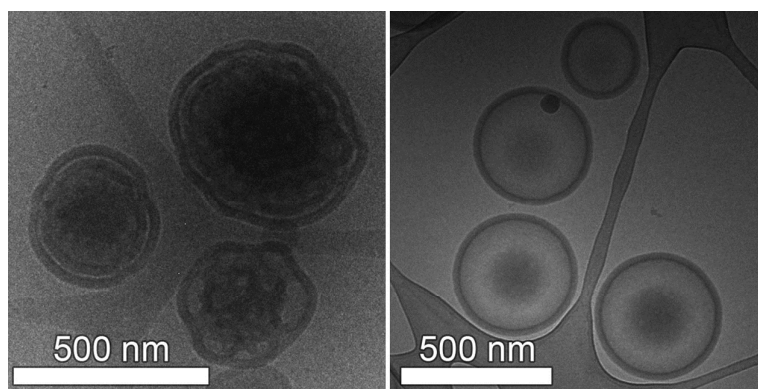


Figure S4: Cryo-TEM images of blended PEG_{22/44}-PDLLA₁₂₀ and non-blended PEG₄₄-PDLLA₁₂₀ prepared via dialysis against 50 mM NaCl at 5 °C, displaying sponge-phase particles (left) and polymersomes (right).

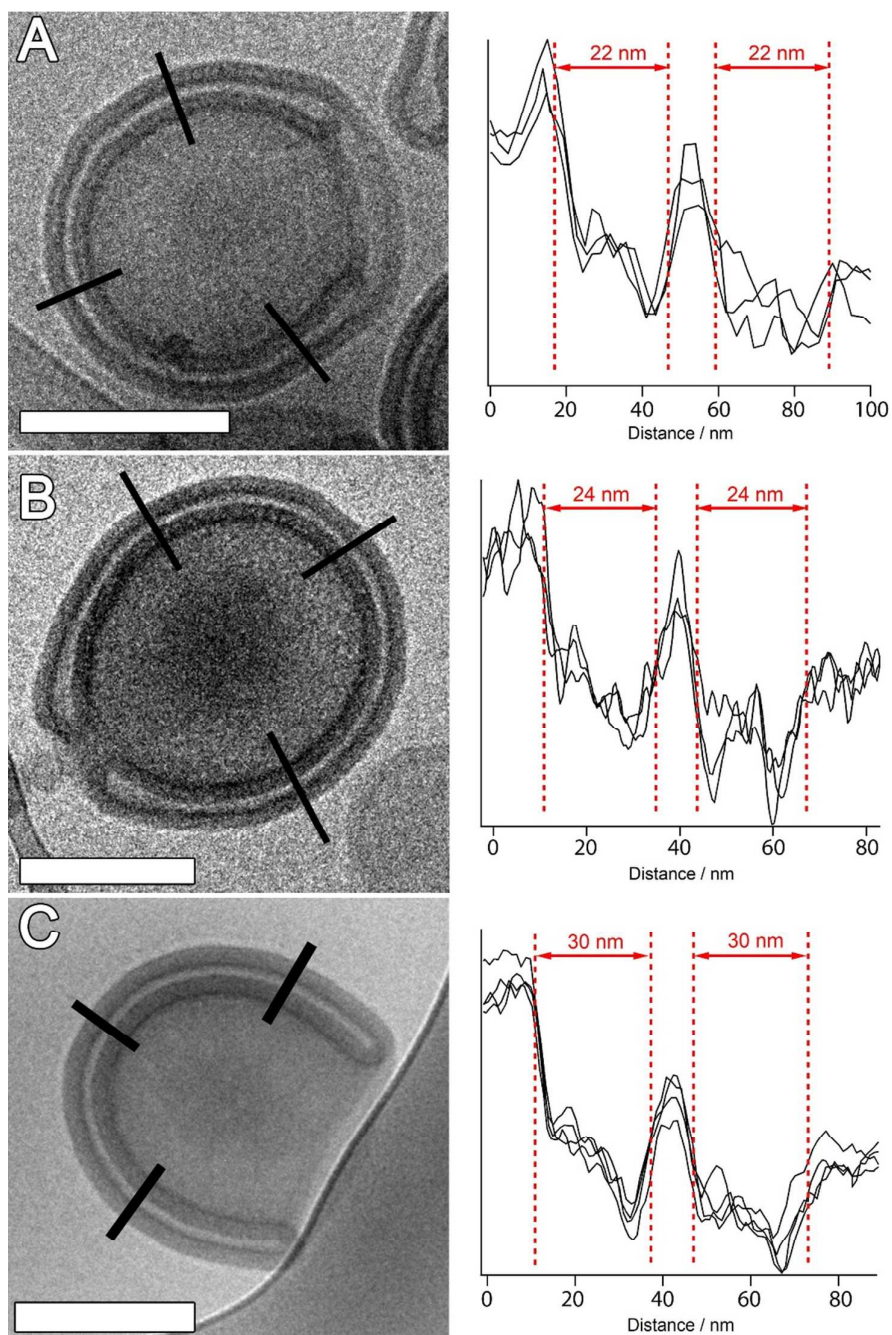
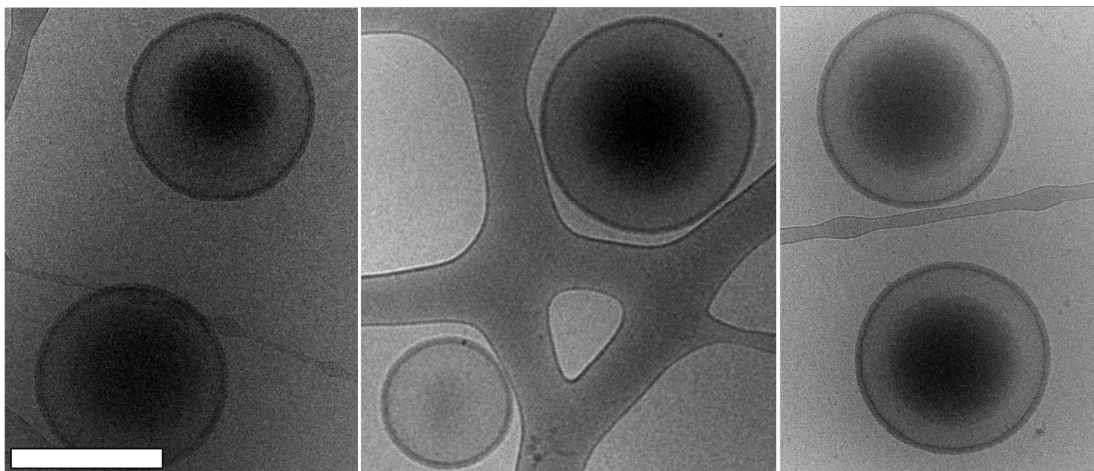
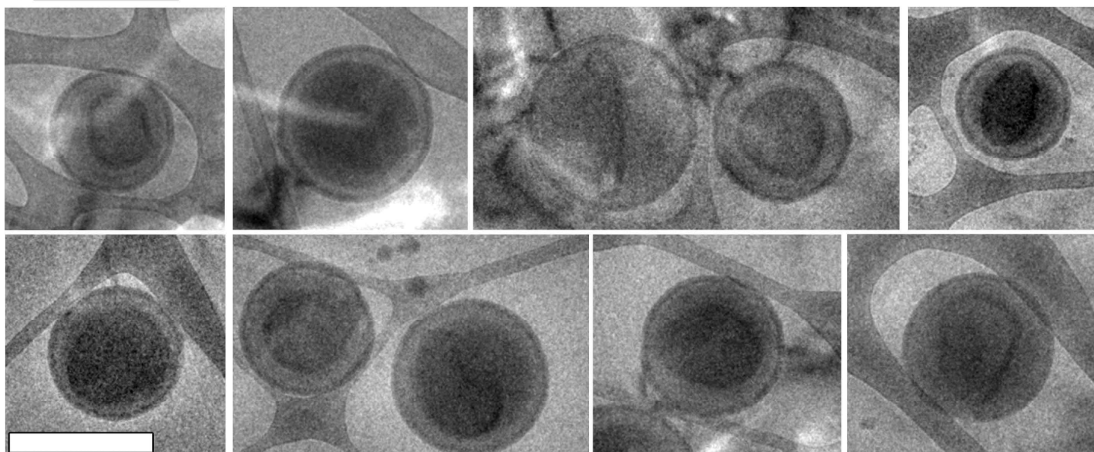


Figure S5: Cryo-TEM images of stomatocytes with varying membrane thickness and corresponding line profile: (A) PEG_{22/44}-PDLLA₇₀ stomatocytes, (B) PEG_{22/44}-PDLLA₉₅ stomatocytes and (C) PEG_{22/44}-PDLLA₁₂₀ sponge-phase particles. All scale bars correspond to 200 nm.

A) Pre-Dialysis



B) 0 mM NaCl



C) 10 mM NaCl

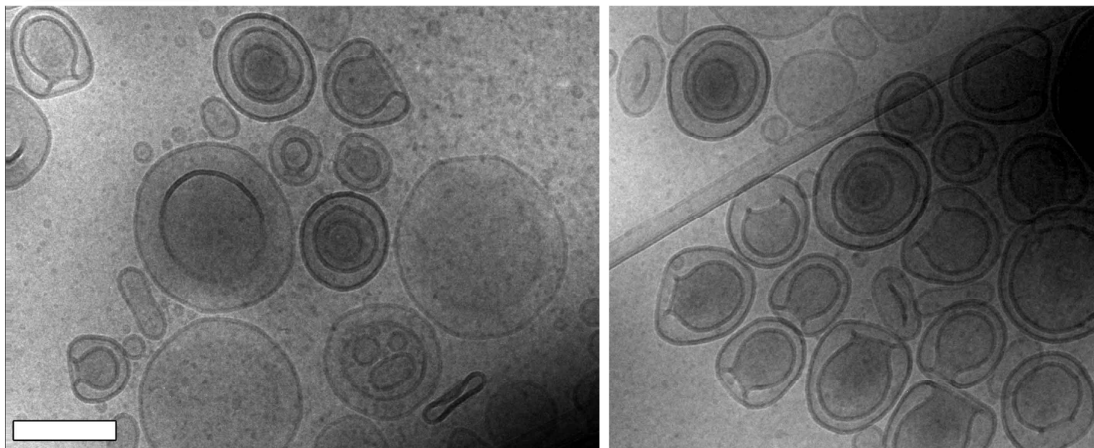
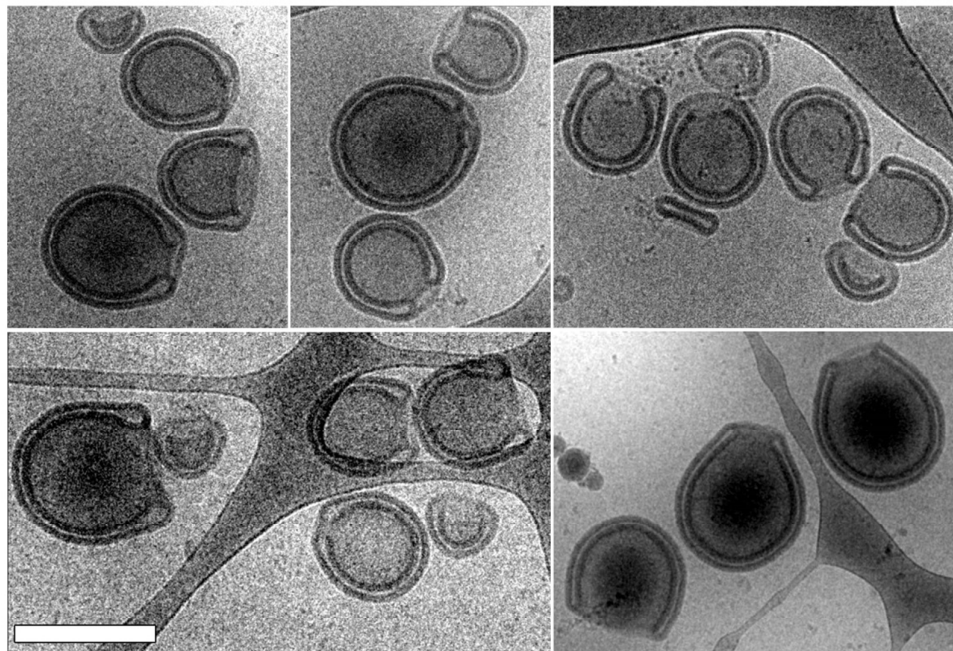


Figure S6: Cryo-TEM images of a) PEG_{22/44}-PDLLA₉₅ polymersomes before dialysis. Shape transformation of polymersomes (A) after dialysis against MilliQ at 5 °C (B) and against 10 mM NaCl (C) at 5 °C. All scale bars correspond to 500 nm.

A) 50 mM NaCl



B) 100 mM NaCl

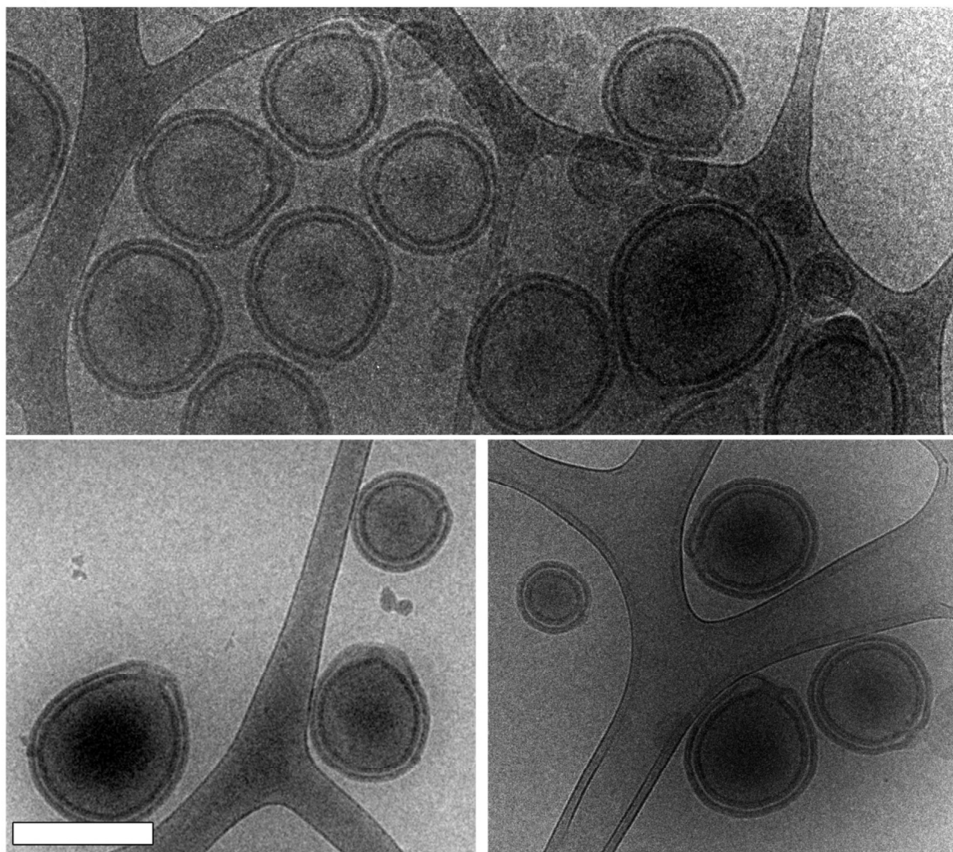
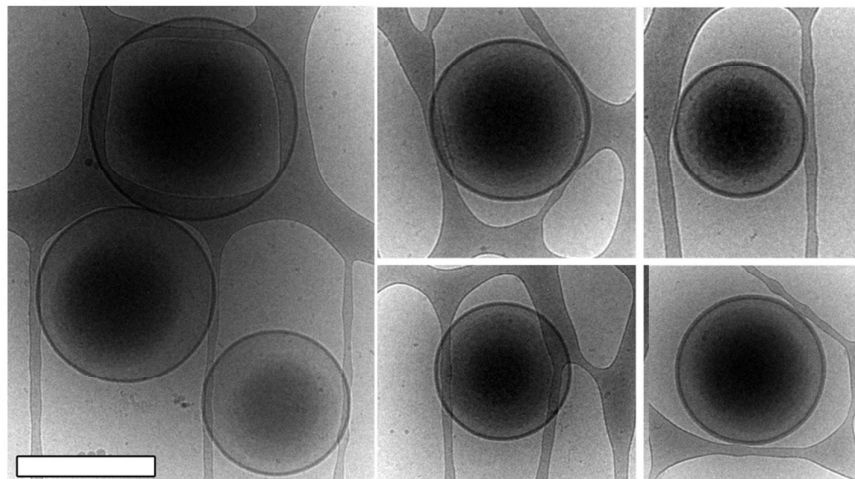
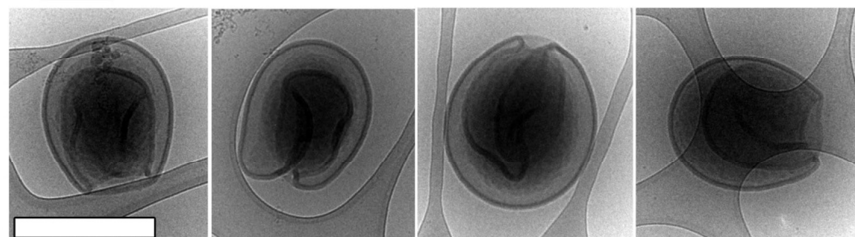


Figure S7: Cryo-TEM images of stomatocytes prepared from PEG_{22/44}-PDLLA₉₅ obtained *via* dialysis against 50 mM NaCl (A) and 100 mM NaCl (B) at 5 °C. Please note decreasing neck size at higher salt concentrations. All scale bars correspond to 500 nm.

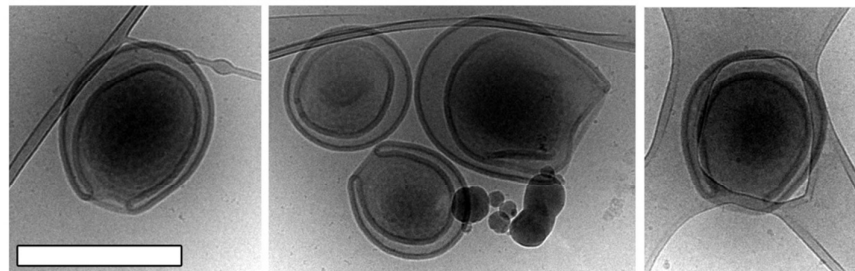
A) 0 min



B) 5 min



C) 15 min



D) 30 min

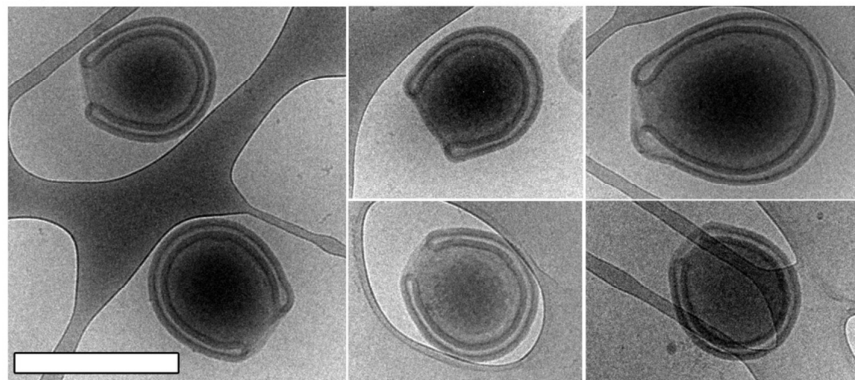
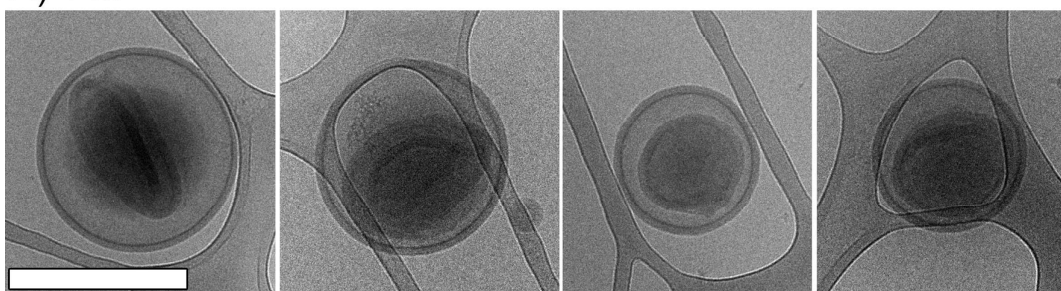


Figure S8: Cryo-TEM images of the shape transformation of polymersomes towards stomatocytes at different time intervals. Stomatocytes were obtained via dialysis of polymersomes of PEG_{22/44}-PDLLA₉₅ against 50 mM NaCl at 5 °C. All scale bars correspond to 500 nm.

Start (min)	End (min)	Mode	Cross flow start (mL min ⁻¹)	Cross flow end (mL min ⁻¹)
0	2	Elution	0.20	0.20
2	3	Focus	-	-
3	5	Focus + inject	-	-
5	6	Focus	-	-
6	36	Elution	0.20	0.10
36	43	Elution	0.10	0.00
43	44	Elution	0.00	0.00
44	45	Elution + inject	0.00	0.00
45	50	Elution	0.00	0.00

Table S2: General method for AF4 fractionation. 20 μL Aliquots of polymersome samples were injected into the AF4 short channel and the conditions were programmed as follows: detector flow (1.5 mL min⁻¹), focus flow (0.70 mL min⁻¹) and injection flow (0.2 mL min⁻¹)

A) 1 hr



B) 2.5 hr

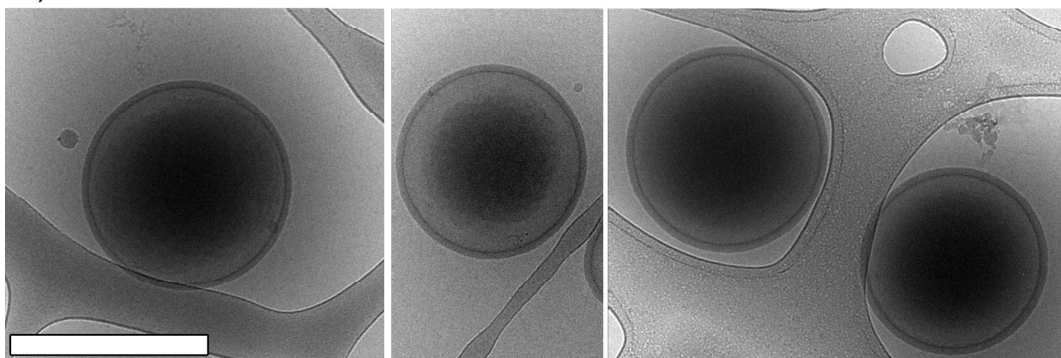


Figure S9: Cryo-TEM images tracking the reverse dialysis process as stomatocytes are reverse engineered to spherical polymersomes after dialysis against 50 vol% organic solvent. All scale bars correspond to 500 nm.

References

1. Abdelmohsen, L. K. E. A. *et al.* Formation of Well-Defined, Functional Nanotubes via Osmotically Induced Shape Transformation of Biodegradable Polymersomes. *J. Am. Chem. Soc.* **138**, 9353–9356 (2016).
2. Rikken, R. S. M. *et al.* Shaping Polymersomes into Predictable Morphologies via Out-of-Equilibrium Self-assembly. *Nat. Commun.* **7**, (2016)