Electronic Supplementary Information for:

## Tubulation onto peptide vesicle by phase-separation of a binary mixture of amphiphilic right-handed and left-handed helical peptides

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## **Materials and Methods**

**Preparation of Molecular Assemblies.** Polypeptide (12 mg) was dissolved in ethanol (120  $\mu$ L). Then an aliquot (30  $\mu$ L) of the peptide solution was injected into a buffer (1 mL, 10 mM Tris-HCl, pH 7.4) with stirring at 0 °C. After 30 min, the dispersion was purified by Sephacryl S-100 column (1.5 × 30 cm, GE healthcare Bio-Sciences) using 10 mM Tris-HCl buffer (pH 7.4) as an eluent to remove ethanol. Molecular assemblies of different compositions were prepared similarly.

**Circular Dichroism (CD).** CD measurements were carried out on a JASCO J600 spectropolarimeter with an optical cell of 0.1 cm optical path length at room temperature. The sample concentration in 10 mM Tris-HCl buffer (pH 7.4) was 0.375 mM (per amino acid residue).

**Transmission Electron Microscopy (TEM).** TEM images were taken using a JEOL JEM-2000EXII at an accelerating voltage of 100 kV. For the observation, a drop of dispersion was mounted on a carbon-coated Cu grid and stained negatively with 2% uranyl acetate, followed by suction of the excess fluid with a filter paper.

**Synthesis of sSLL.** sSLL was prepared by the same method<sup>[1]</sup> as SLL but the capping reagent was lipoic acid instead of glycolic acid. After complete consumption of Sar NCA was confirmed, lipoic acid (50 mg, 5 eq.), HATU (90 mg, 5 eq.) and triethylamine (64  $\mu$ L, 7.5 eq.) were added to the solution. After stirring for 12 h, another lipoic acid (25 mg, 2.5 eq.), HATU (45 mg, 2.5 eq.) and triethylamine (32  $\mu$ L, 3.8 eq.) were added to the solution. After stirring for 12 h, the solution was condensed, and the residue was purified by a Sephadex LH20 column with methanol as an eluent to afford sSLL.

<sup>1</sup>H NMR(400 MHz, MeOH-d) δ (ppm) 8.2–7.7 (m, 11H, amide), 7.4–7.3 (br, 1H, amide), 4.6–3.8 (br, 56H, LeuC<sup>α</sup>*H*, SarC*H*<sub>2</sub>), 3.66 (s, 3H, OC*H*<sub>3</sub>), 3.55 (quin, 1H, SSC*H*), 3.3–2.8 (m, 79H, Sar N-C*H*<sub>3</sub>, SSC*H*<sub>2</sub>, CH<sub>2</sub>C*H*<sub>2</sub>CO), 2.5–2.4 (br, 2H, SSCH<sub>2</sub>C*H*<sub>2</sub>C*H*<sub>2</sub>), 2.0–1.7 (br, 8H, C*H*<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CO, CH<sub>2</sub>C*H*<sub>2</sub>CH<sub>2</sub>CCO, CH<sub>2</sub>C*H*<sub>2</sub>CH<sub>2</sub>CCO, CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CCO, CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CCO, CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CCO, CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CCO, CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CO, CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CO), 1.7–1.3 (m, 36H, LeuC*H*<sub>2</sub>, LeuC<sup>γ</sup>*H*, AibC*H*<sub>3</sub>), 1.1–0.8 (m, 36H, Leu(C*H*<sub>3</sub>)<sub>2</sub>).

Mixtures of **SLL** and **SDL** in the molecular assemblies were checked by CD whether the helix contents were affected by mixing in the assemblies.



**Fig. S1.** CD spectra with changing the mixing ratio of **SLL** and **SDL** in the molecular assemblies. The molecular assemblies were formed from varying mole ratios of amphiphilic polypeptides, **SLL** and **SDL**, in 10 mM Tris-HCl buffer (0.18 mg / 1 mL).

We can improve the purity of the nano round-bottom flask assembly by modifying the preparation process. At first, two types of molecular assemblies, the nanotube of **SLL** (50 wt%) and the nano planar sheet composed of an equimolar mixture of **SLL** and **SDL** (25 and 25 wt%), were prepared separately. Then, both molecular assemblies were mixed together and heated at 90 °C for 1 h to obtain the round-bottom flask assembly with a high yield of 38 %. The dispersion was purified by a size exclusion chromatography of Sephacryl S-100 to obtain nearly pure round-bottom flask assembly. In this case, the planar sheet has a chance to fuse with the nanotube to yield the nano round-bottom flask or is just transformed by itself to vesicle. However, the yield of the round-bottom flask was found to be 38 % (the number of the round-bottom flask/(the number of the round-bottom flask + the number of the nanotube) = 168/443 in six TEM images), which coincides with the calculated value for quantitative production without the vesicle transformation (Fig. S2). Priority of the round-bottom flask assembly over the vesicle under the present preparation conditions remains to be solved.



**Fig. S2.** TEM images (negative staining with uranyl acetate) of molecular assemblies prepared from a mixture of the nanotube prepared from **SLL** and the planar sheet prepared from an equimolar mixture of **SLL** and **SDL** after heat treatment (90 °C, 1 h). The nanotube and the planar sheet assemblies were

prepared in 10 mM Tris-HCl buffer (pH 7.4) (3 mg/1 mL) by the ethanol injection method with and without heat treatment, respectively.

The calculation method is as follows:

The occupied surface area of a molecule, **SLL**, is same to that of **SDL** in assembly. So, the percentage of constituent molecules, **SLL** or **SDL**, of the morphology can be calculated by the surface area. Now, **SLL/SDL** = 75/25 in this system, because the nanotube of **SLL** (50 wt%) and the planar sheet composed of an equimolar mixture of **SLL** and **SDL** (25 and 25 wt%), were prepared, and then, both molecular assemblies were mixed. The surface area of the spherical part (diameter = 180 nm) of round-bottom flask type is 101736 nm<sup>2</sup>. This part is composed of a mixture of **SLL** (25 wt%) and **SDL** (25 wt%). On the other hand, the surface area of the neck part (diameter 60 nm and length 200 nm) is 37680 nm<sup>2</sup>, suggesting that this part is composed of **SLL** (19 wt%), by (25 + 25)\*(37680/101736). Therefore, It is considered that the rest **SLL** (31 wt%) formed the free nanotube in this system, indicating that the ratio of the round-bottom flask (the round-bottom flask/(the round-bottom flask + the free nanotube)) is 19/(19 + 31).