## Fabrication of Artifical Toroid Nanostructures by Modified

# β-sheet Peptide

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### 1. Peptide synthesis:



Fmoc-tyrosine derivative with tri(ethylene glycol) monomethyl ether and Fmoc-L-Phenylalanine-4'-azobenzene (Fmoc-Phe-Azo) were synthesized according to previous method. [1]



All the peptides were synthesized with a microwave-equipped liberty peptide synthesizer (CEM). Rink amide resin was utilized and loaded using standard methods HBTU activation. Peptide synthesis was carried out using standard Fmoc methods.

### 2. Measurements:

**Circular Dichroism Spectroscopy**. Peptide solutions were analyzed by CD spectroscopy to characterize secondary structure. Spectra were recorded on a JASCO model J-810 spectropolarimeter. CD spectra were obtained from 260-190 nm with a 1.0 nm step, 1.0 nm bandwidth, and a 3 s collection time per step at 25  $^{\circ}$ C in a 0.1 cm path length quartz cuvette (Hellma). Scans were repeated five times and averaged. The JASCO software was used for background subtraction, conversion to molar ellipticity per amino acid residue.

**Fourier Transform Infrared Spectroscopy**. FT-IR spectra were obtained using a Bruker Equinox 55 FT-IR spectrophotometer. The aqueous solution of the peptide samples was cast onto gold plate. Absorbance spectra were obtained from 1750 to 1550 cm<sup>-1</sup> with a 4 cm<sup>-1</sup> resolution. 502 scans were acquired and background spectra were also collected and subtracted.

**Dynamic Light Scattering (DLS)**. DLS experiment was performed at room-temperature with LV/CGS-3 Compact Goniometer System equipped with He-Ne laser operating at 632.8 nm. The scattering angle was  $90^{\circ}$ . Bbefore measurement, the sample was centrifuged at  $16,110 \times g$  for 20 min to sediment and dust particles. The size distribution was determined by using a constrained regularization method.

**Transmission Electron Microscopy.** Samples (2  $\mu$ L) were spotted directly onto a carbon-coated copper grid and dried completely. Then 2 % (w/v) uranyl acetate solution (2  $\mu$ L) was added and allowed to stand for 1 min. Then, the excess amount of solution was carefully removed by capillary action (filter paper). The cryogenic transmission electron microscopy experiments (cryo-TEM) were performed with a thin film of solution of the peptide (5 $\mu$ L) transferred to a lacey-supported grid. The thin aqueous films were prepared under controlled temperature and humidity conditions (97-99 %) within a custom-built environment chamber to prevent evaporation of water from sample solution. The excess amount of liquid was blotted with filter paper for 2-3 s, and the thin aqueous films were rapidly vitrified by plunging them into liquid ethane (cooled by liquid nitrogen) at its freezing point. The grid was transferred onto a Gatan 626 cryoholder using a cryotransfer device. After that they were transferred to a JEOL-JEM 2100 TEM instrument. Direct imaging was carried out at a temperature of approximately -175 °C and with a 120 kV accelerating voltage, while acquiring the images with a SC 1000 CCD camera (Gatan, Inc., USA). The data were analyzed with Digital Micrograph software.

**Peptide self-assembly.** Peptides were dissolved in unbuffered filtered water (Millipore 0.22  $\mu$ m filter, 18  $\Omega$ ) by the addition of water to the lyophilized powder followed by sonication (3 min) to obtain homogeneous and optically transparent solutions.

#### 3. Peptide Characterization.

The molecular weights of the peptides were confirmed by MALDI-TOF mass spectrometry.

Peptide **P1** (MALDI-TOF-MS) m/z 2070.08 (2070.43 calcd for  $[M+H^+]$ ), 2092.17 (2092.42 calcd for  $[M+Na^+]$ ), 2108.21 (2108.53 calcd for  $[M+K^+]$ ).

Peptide **P2** (MALDI-TOF-MS) m/z 2071.24 (2071.38 calcd for  $[M+H^+]$ ), 2093.01 (2093.37 calcd for  $[M+Na^+]$ ), 2109.18 (2109.48 calcd for  $[M+K^+]$ ).

Peptide **P3** (MALDI-TOF-MS) m/z 2135.82 (2135.51 calcd for  $[M+H^+]$ ), 2157.79 (2157.5 calcd for  $[M+Na^+]$ ), 2173.85 (2173.61 calcd for  $[M+K^+]$ ).

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Fig. S1 MS spectra of peptides P1 (top), P2 (middle) and P3 (bottom).



Fig. S2 HPLC spectra of peptides P1 (top), P2 (middle) and P3 (bottom).

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Fig. S3 DLS spectra of peptide P1 obtained from 40  $\mu$ M aqueous solution.



Fig. S4 TEM image of P1 obtained from 40  $\mu$ M aqueous solution with NaCl concentration of 200 mM.



Fig. S5 TEM image of P2 obtained from 40 µM aqueous solution.



Fig. S6 cryo-TEM image of peptide P2 obtained from 40  $\mu M$  aqueous solution.



Fig. S7 TEM image of peptide P2 obtained from 40  $\mu M$  aqueous solution.