Supporting information

Macroscale Assembly of Peptide Nanotubes

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(1) X-ray Powder Diffraction



Fig. S1. Comparison of X-ray diffraction patterns of nanotubes before (a) and after (b) sulfate bundling. The typical cross- β conformation with ~ 5Å and ~ 10 Å reflections are conserved. The presence of 9.4 Å reflection suggests the antiparallel β -sheet structure.

(2) Comparison of derived structure factors for nanotubes in salt solutions using form factors for a core-shell cylinder and a rectangular cross-section cylinder

The diameter of the nanotubes prior to bundling, 52 nm, and the repeat distance of 45.6 nm based on the first order peak at Q ~ 0.0138 Å⁻¹ were used to determine the cross-section of the nanotubes. The tubes inside the bundles do not have a perfect square cross-section. Rather it fits better with a structure that is between a circle and a square, having square sides of length (l) and round edges of radius (r):

$$2r + l = 45.6 (1)$$

 $2r + 4l = 52 \text{ nm}$

Solving these simultaneous equations gives r = 11.09 nm and l=23.42 nm

We compared the extracted structure factors using the form factors for core-shell cylinders with a circular (r=26 nm) and a square cross-section with a = 41 nm (Fig. 2S). In both cases, the peak positions are similar and consistent with a lamellar order.

Core-shell circular cylinder form factor:



Generalized core-shell square cylinder form factor:



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Fig. S2. Structure factor evaluations for different sulfate titrations using the form factors for core-shell cylinder with a circular cross-section (r=26 nm) and a square cross-section with a 41 nm width for the 9 mM sulfate titration. The peak intensity (red curve with markers for the 9 mM sulfate) is more pronounced, but the peak positions still correspond to the lamellar