

Supplementary Information

Accelerated Amyloid Beta Pathogenesis by Bacterial Amyloid FapC

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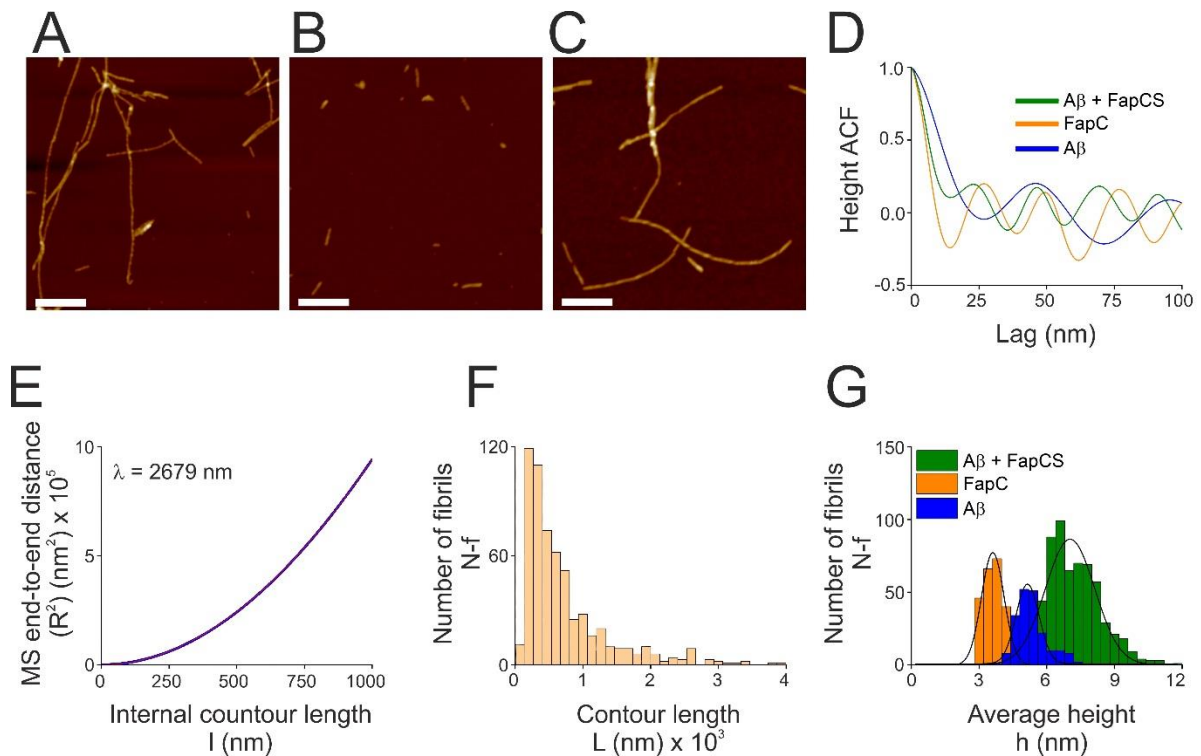
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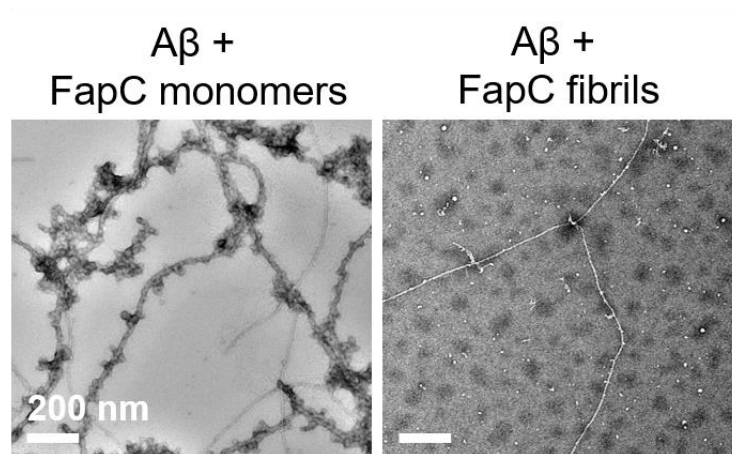
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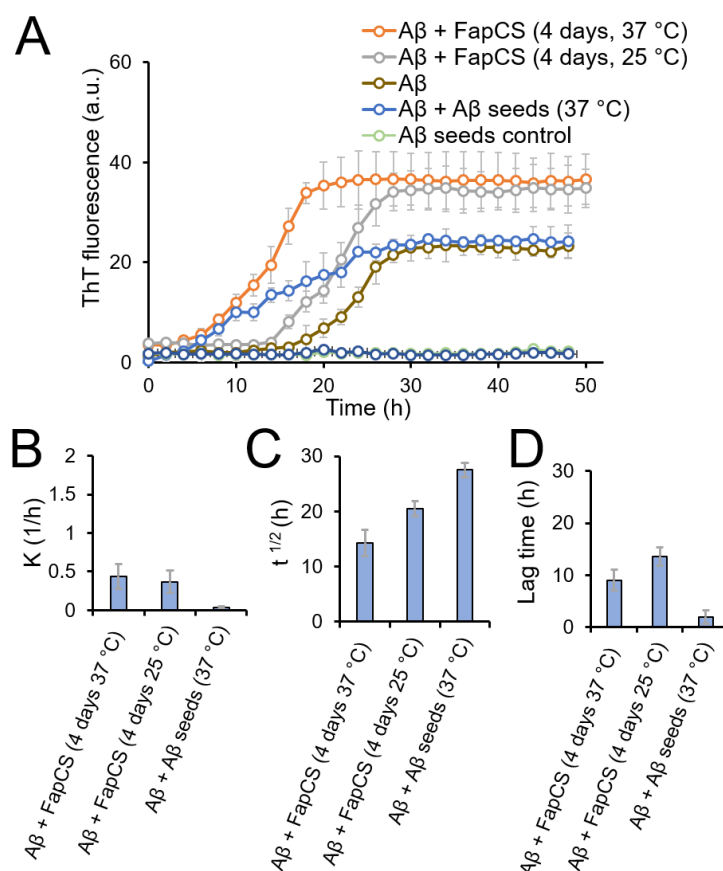


Supplementary Figure 1. AFM and morphological statistics for FapCS-seeded A β , A β and FapC fibrils. AFM images of (A) mature FapC fibrils (37 °C, 1 week), (B) FapCS and (C) A β fibrils. Scale bar for all AFM images is 200 nm. (D) Estimation of periodicity of FapCS-seeded A β , A β and FapC fibrils calculated from the autocorrelation function (ACF) of AFM images. (E) Mean squared end-to-end distance versus internal contour length providing the persistence length of FapCS-seeded A β fibrils. (F) Contour length distribution of FapCS-seeded A β fibrils. (G) Average height distributions of FapCS-seeded A β , FapC and A β fibrils.

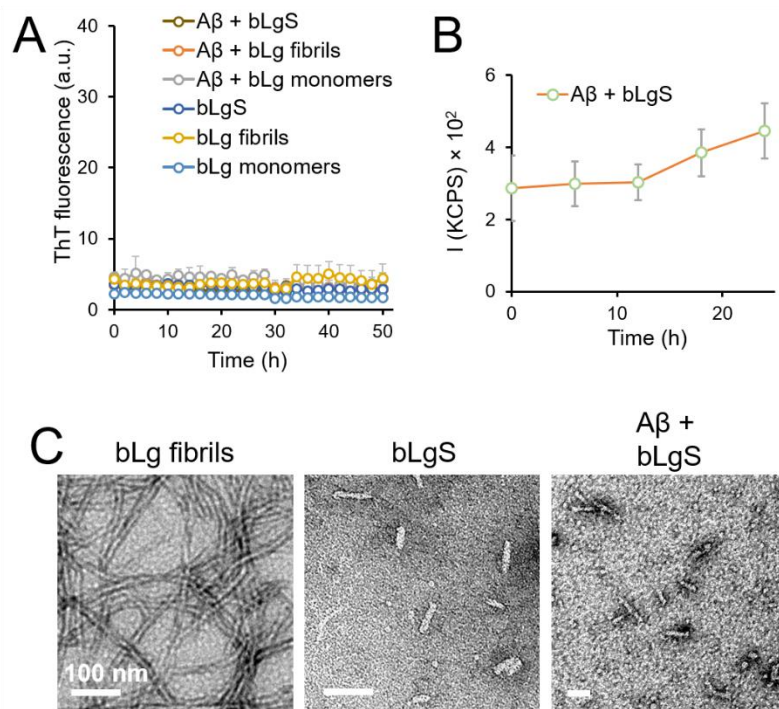


Supplementary Figure 2. TEM images for A β fibrillized in the presence of FapC monomers and preformed FapC fibrils. Thick A β fibrils coated with FapC monomers were observed under TEM. Incubation time was 50 h at 37 °C. A β , when incubated with

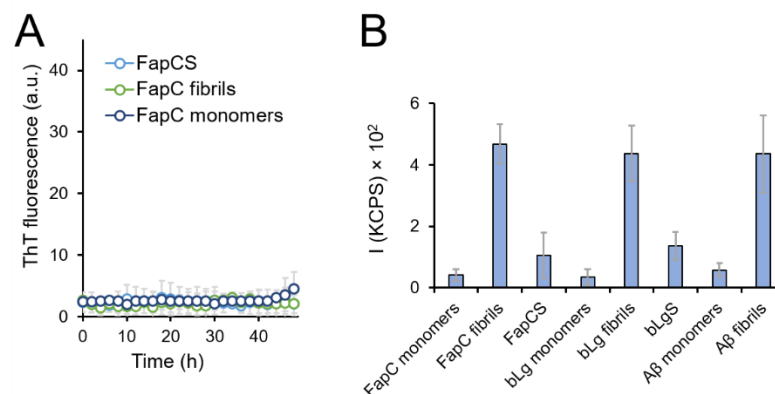
preformed FapC fibrils, did not coat FapC fibrils. However, the fibrillization of A β was significantly suppressed.



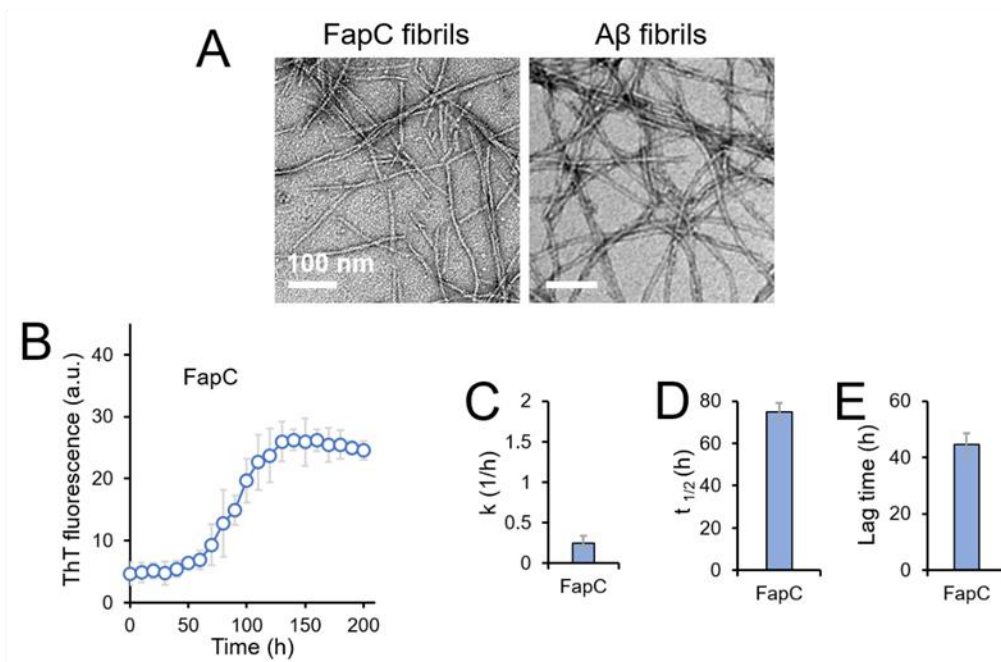
Supplementary Figure 3. ThT kinetic assay and parameters of A β with FapCS from younger FapC fibrils, with A β seeds. FapCS (5 μ M) from younger FapC fibrils (fibrillized at 37 and 25 °C for 4 days) also accelerated A β (50 μ M) fibrillization, however, not to an extent observed for FapCS from mature FapC fibrils (fibrillized at 37 °C for 1 week) (n=3). Self-seeded A β (A β + A β seeds, 10:1 molar ratio) shortened the lag phase to 1.9 h from 23 h for A β alone while the fibrillization rate constant k is decreased to 0.03 h⁻¹.



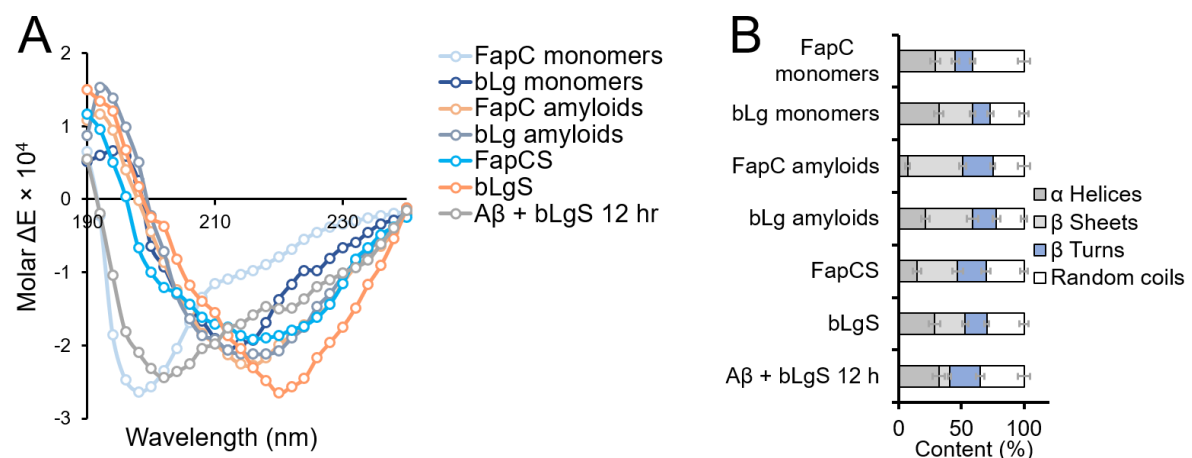
Supplementary Figure 4. Seeds from functional amyloids of β -lactoglobulin (bLgS) and their impact on A β fibrillization. bLgS were produced via sonication and were cross-seeded with A β . (A) ThT assay indicated that bLgS, bLg monomers and preformed amyloids (5 μ M) inhibited A β (50 μ M) fibrillization (n=3). (B) SLS of A β fibrillized in the presence of bLgS (n=3). (C) TEM micrographs of bLg fibrils, bLgS and A β + bLgS.



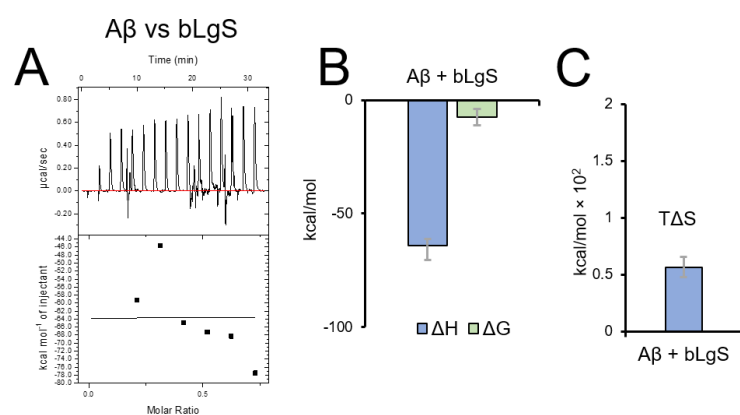
Supplementary Figure 5. (A) ThT assay of FapCS, FapC fibrils and FapC monomers (5 μ M), as controls, for 50 h incubation at 37 $^{\circ}$ C (n=3). (B) SLS of controls of FapC monomers and fibrils, FapCS, bLgS, bLg monomers and fibrils, and A β monomers and fibrils (n=3).



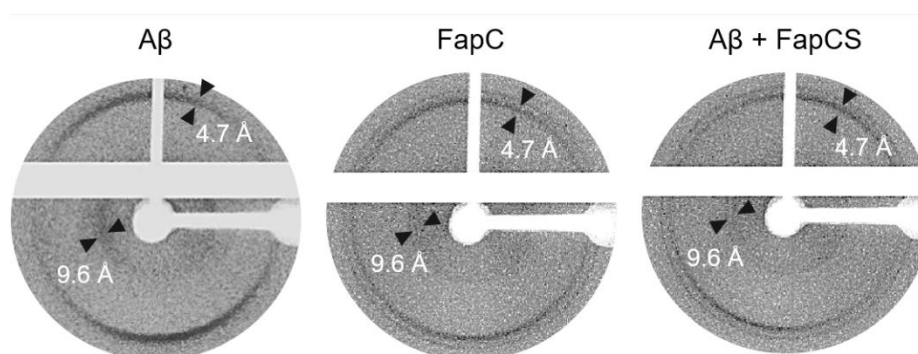
Supplementary Figure 6. (A) TEM of FapC and A β fibrils. (B) ThT assay of FapC (50 μ M) at 37 $^{\circ}$ C (n=3). It took 150 h for FapC to reach the saturation phase. (C) Fibrillization rate constant k , (D) half-life $t_{1/2}$ and (E) lag time of FapC fibrillization (n=3).



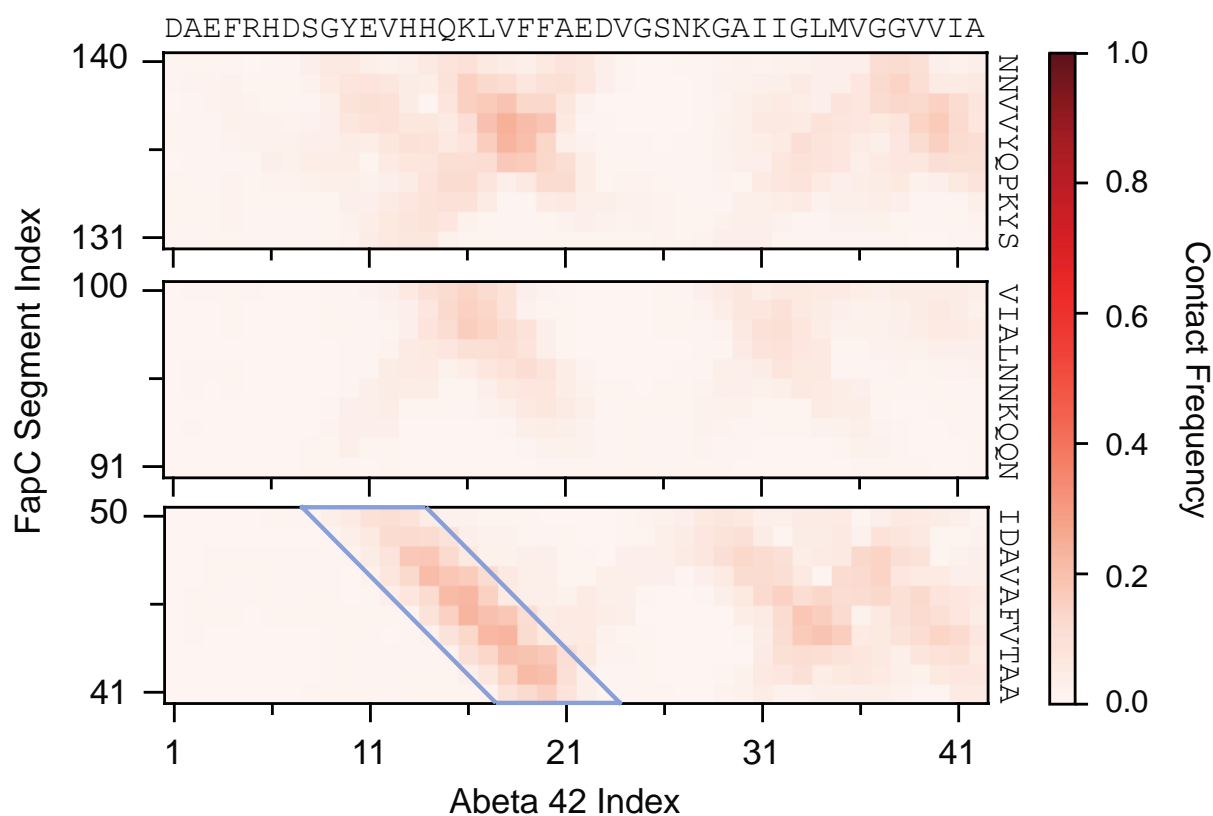
Supplementary Figure 7. Secondary structure of FapC and bLg controls and A β in the presence of bLgS. (A) CD spectra and (B) secondary structure (%) of monomers and fibrils of bLg and FapC, bLgS, FapCS and A β in the presence of bLgS at 12 h incubation at 37 $^{\circ}$ C (n=3). After incubation with A β , the α -helical content of bLgS was slightly increased.



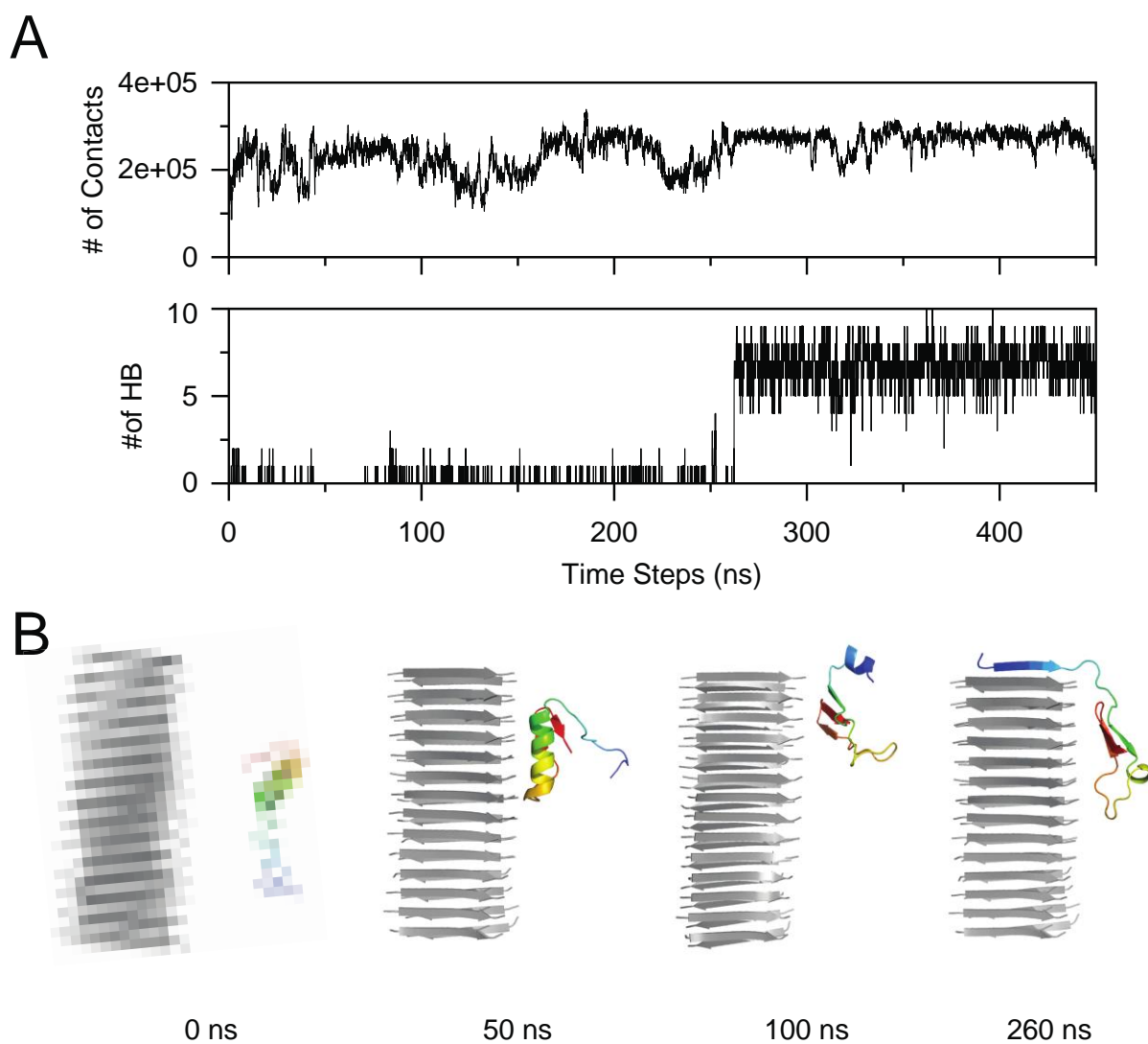
Supplementary Figure 8. (A) Isothermal titration calorimetry (ITC) of A β with bLgS. (B) Binding enthalpy (ΔH) and free energy (ΔG) and (C) entropic factor ($T\Delta S$) of A β with bLgS ($n=3$).



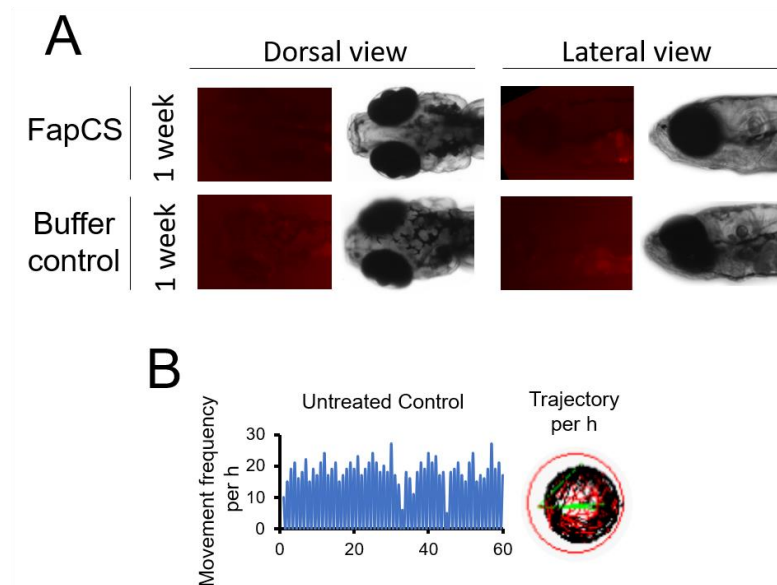
Supplementary Figure 9. X-ray fiber diffraction of A β , FapC and A β + FapCS. All three samples showed similar inter-sheet and inter-strand distance of 9.6 and 4.7 Å, respectively.



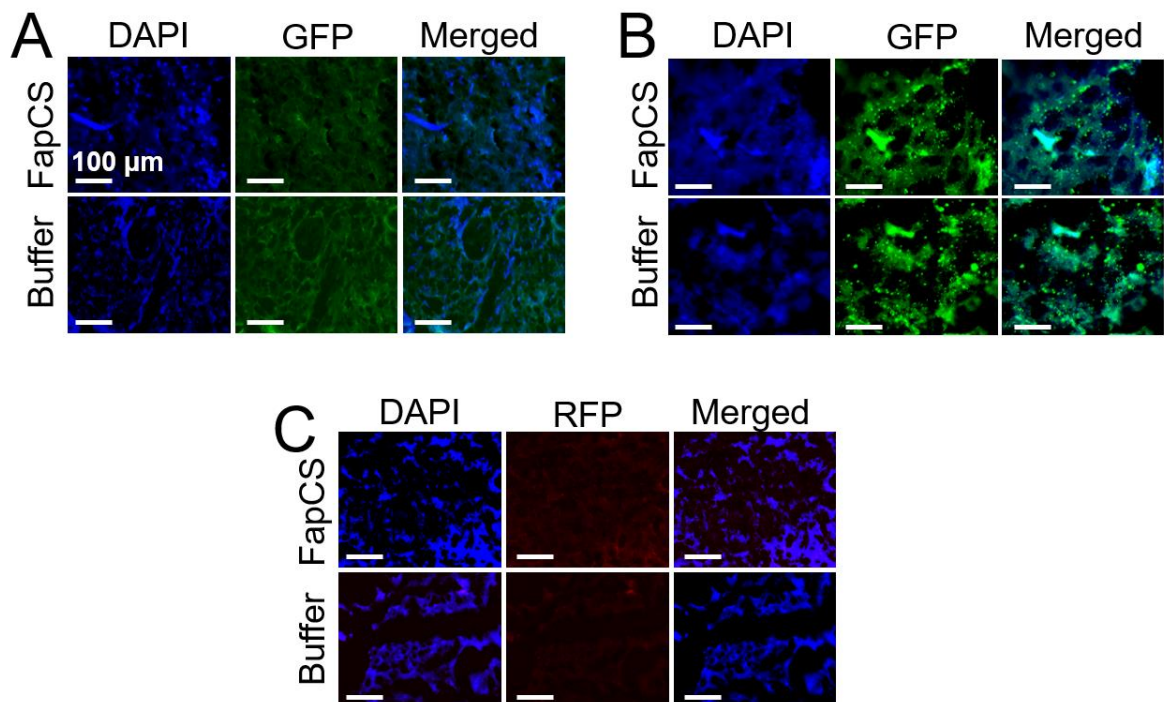
Supplementary Figure 10. The inter-molecular contact frequency maps for the binding of three hotspot fragments (FapC41-50, FapC91-100, FapC131-140) with an A β 42 monomer. Anti-parallel β -sheet formed between FapC41-50 and the amyloidogenic region of A β 16-22 is highlighted with a blue box.



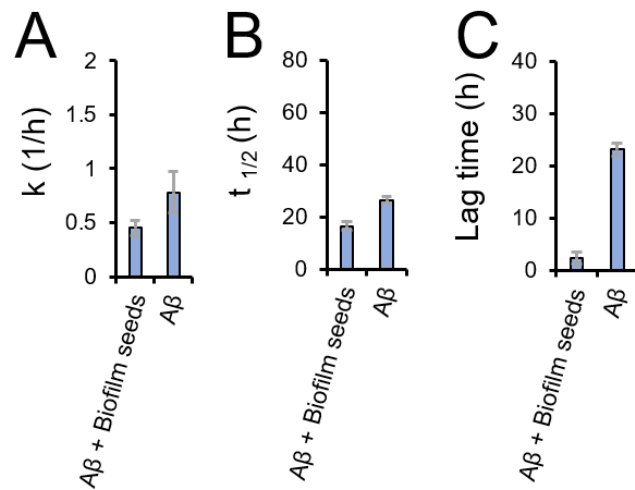
Supplementary Figure 11. (A) Time evolution of the number of atomic contacts between an A β monomer and a FapC nanofibril (top) and the inter-molecular H-bond backbone of an A β fibril (bottom) from a representative trajectory. (B) Snapshots along the simulation trajectory are shown at different times. The FapC nanofibril is shown in grey, while A β 42 in rainbow from its N- (blue) to C-terminus (red).



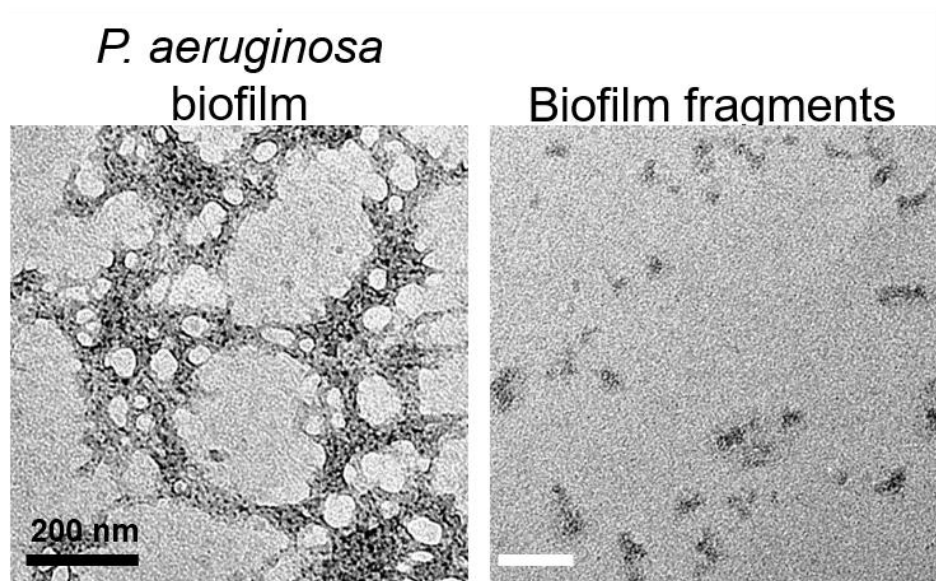
Supplementary Figure 12. (A) Whole-mount fluorescence imaging of FapCS and buffer control at the one-week time point. (B) Movement frequency and trajectory of untreated zebrafish larvae over the period of 1 h observation (n=10 per group and 3 groups per sample).



Supplementary Figure 13. Immunohistochemistry (IHC) of adult zebrafish brain slices for (A) $A\beta$ deposition, (B) depletion of synaptophysin positive cells and (C) TUNEL assay for neuronal cell death for FapCS and buffer control.



Supplementary Figure 14. ThT kinetic parameters of (A) fibrillization rate constant (k), (B) half-life $t_{1/2}$ and (C) lag time for Aβ in the presence and absence of *P. aeruginosa* biofilm fragments (n=3).



Supplementary Figure 15. TEM images for *P. aeruginosa* biofilm and sonicated biofilm fragments/seeds.