

SUPPLEMENTAL MATERIAL

Structure and dynamics of parallel β -sheets, hydrophobic core, and loops in Alzheimer's A β fibrils

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Fig. S1. The conformations of the infinite fibril segments for the S1, S2 and S3 systems after 10 ns of MD at 298 K. Top views of systems (A) S1- C_{2z} , (B) S1- C_{2x} , (C) S2- C_{2z} , and (D) S3- C_{2z} . Lateral view of systems (E) S1- C_{2z} , (F) S1- C_{2x} , (G) S2- C_{2z} , and (H) S3- C_{2z} .

Fig. S2. Secondary structure of the S2- C_{2z} system along MD trajectories at (A) 298 K and (B) 398 K. Boxed regions indicate formation of helical structure (black).

Fig. S3. Secondary structure of the S3- C_{2z} system along MD trajectories at (A) 298 K and (B) 398 K.

Fig. S4. Structures of A β_{1-40} dodecameric fibril segments during the NPT MD simulations at 298, 348 and 398 K. System S1- C_{2z} with -0.5 initial staggering.

Fig. S5. Structures of A β_{1-40} dodecameric fibril segments during the NPT MD simulations at 298, 348 and 398 K. System S1- C_{2x} with +0.5 initial staggering.

Fig. S6. Structures of A β_{1-40} dodecameric fibril segments during the NPT MD simulations at 298, 348 and 398 K. System S2- C_{2z} with -1.5 initial staggering.

Fig. S7. Structures of A β_{1-40} dodecameric fibril segments during the NPT MD simulations at 298, 348 and 398 K. System S2- C_{2z} -AGAA with -1.5 initial staggering and mutations V24A, S26A and N27A.

Fig. S8. Structures of A β_{1-40} dodecameric fibril segments during the NPT MD simulations at 298, 348 and 398 K. System S3- C_{2z} with -1.5 initial staggering.

Figures

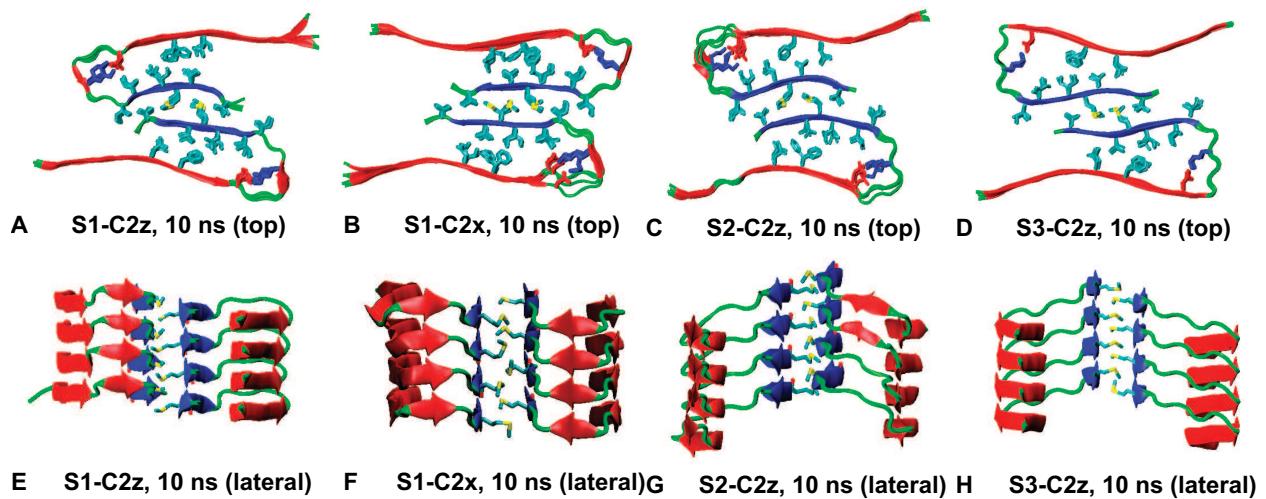


Fig. S 1:

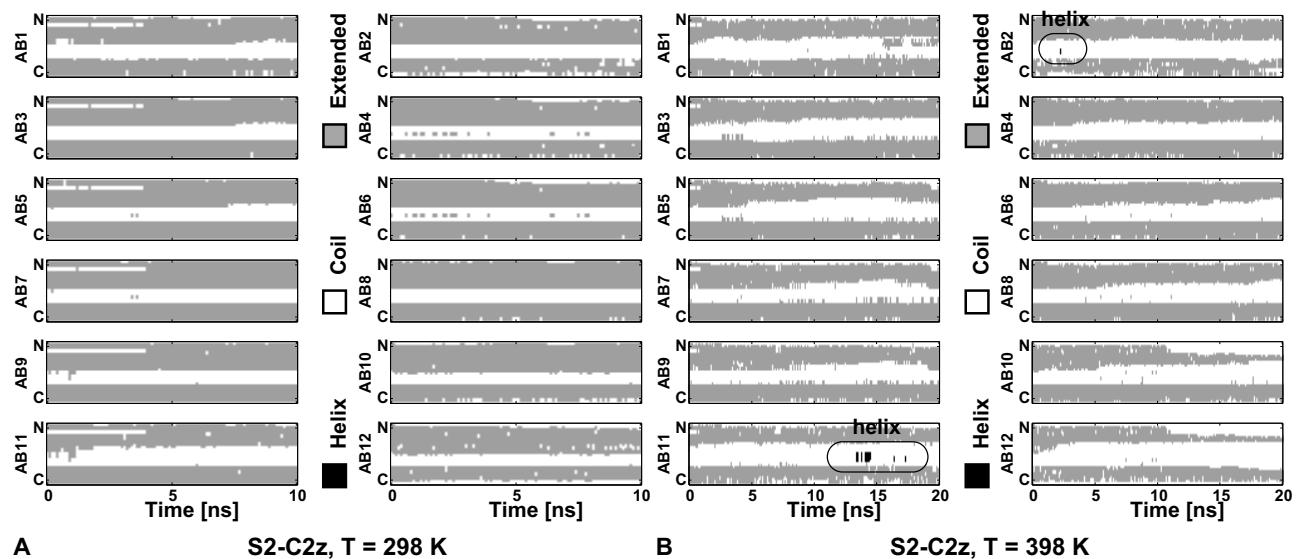


Fig. S 2:

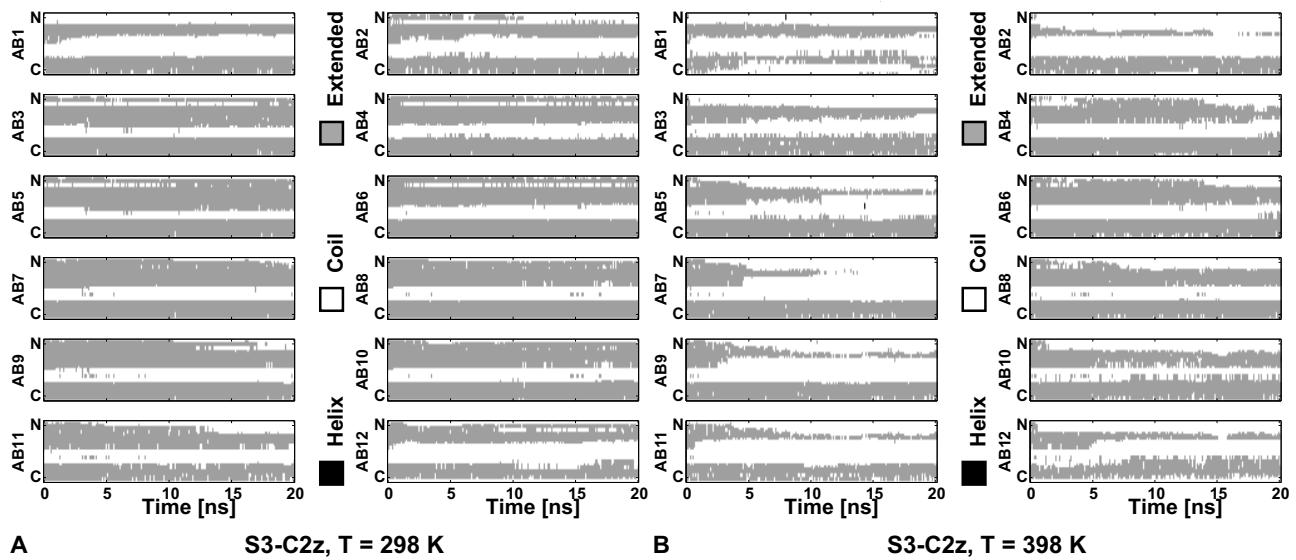


Fig. S 3:

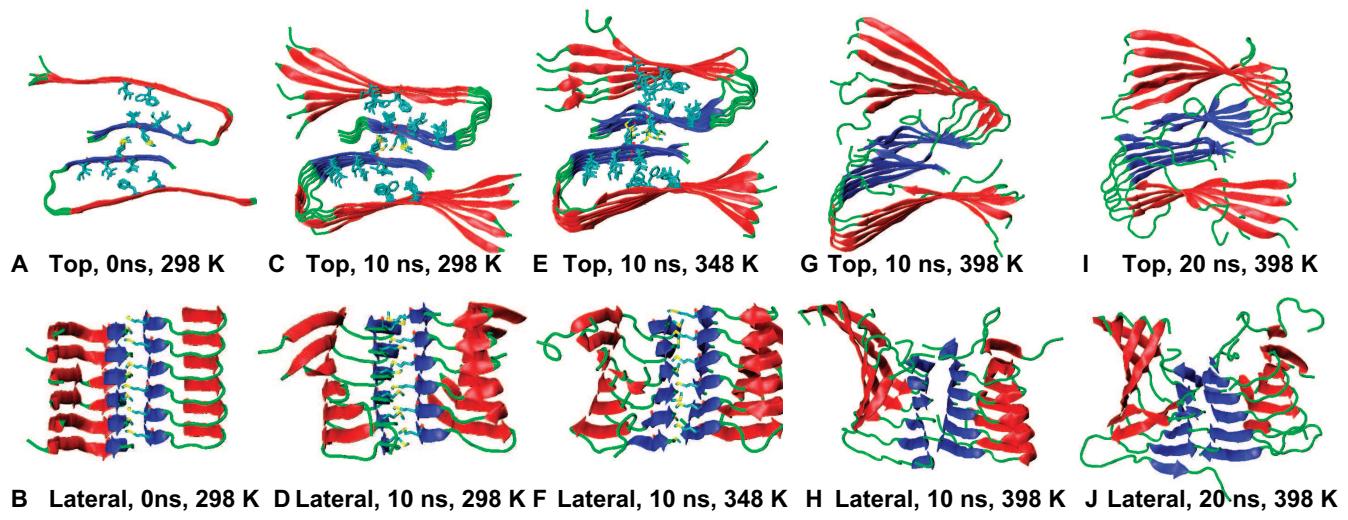


Fig. S 4:

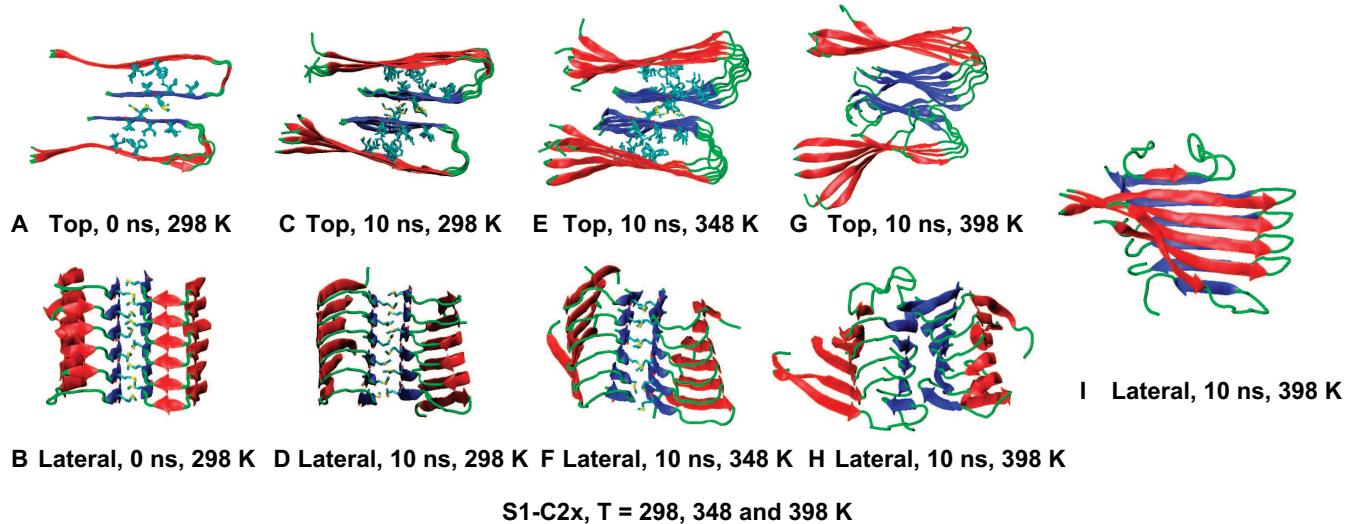


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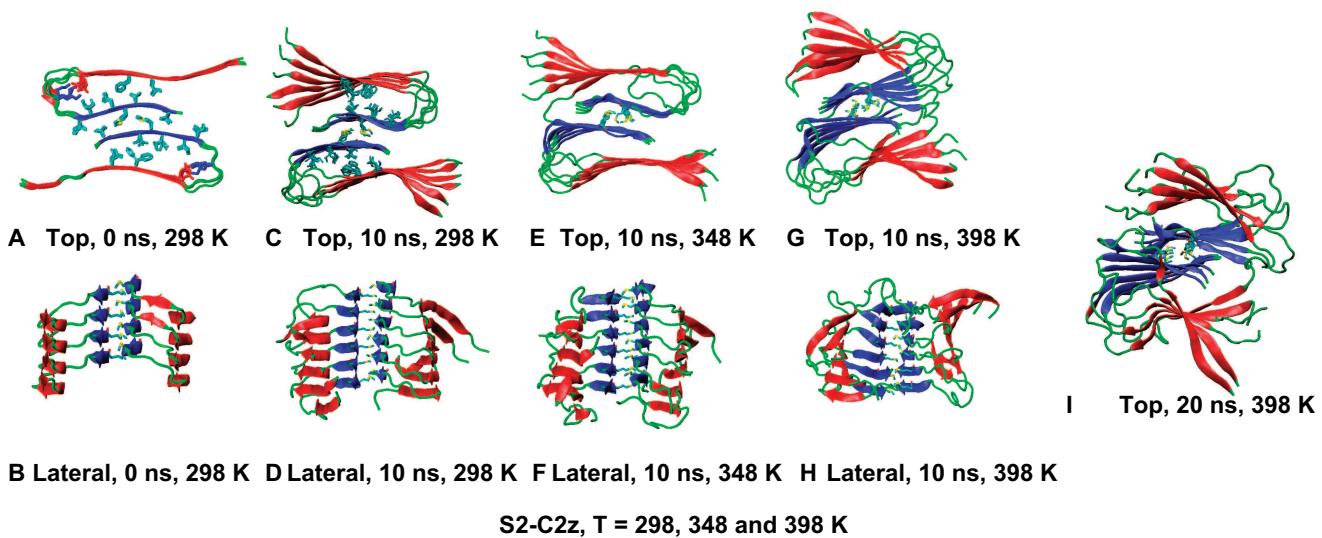


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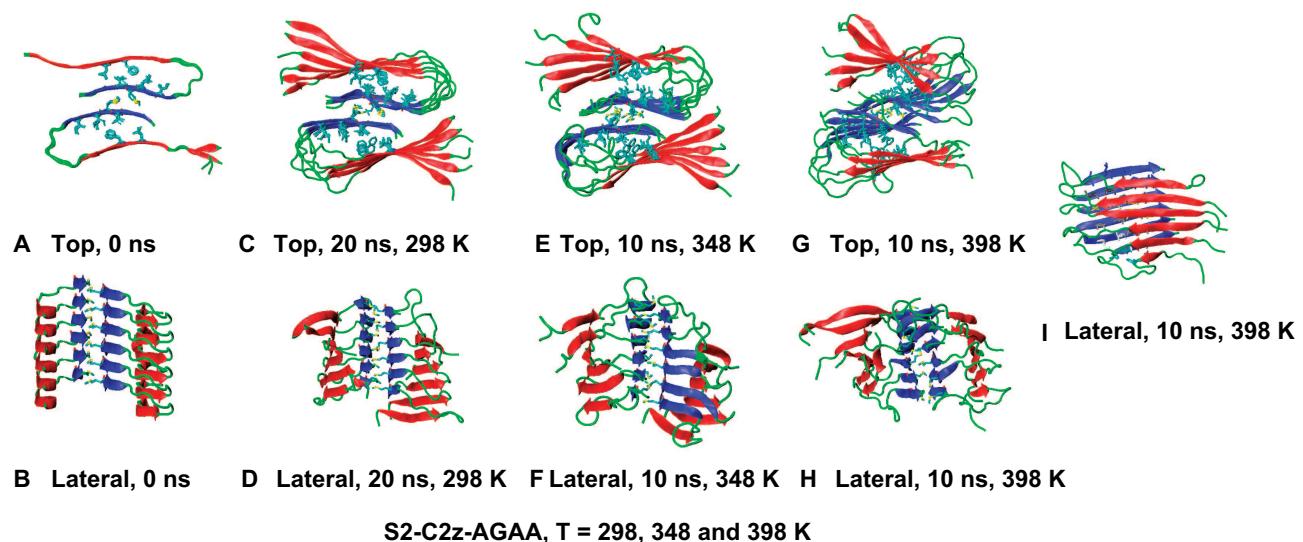


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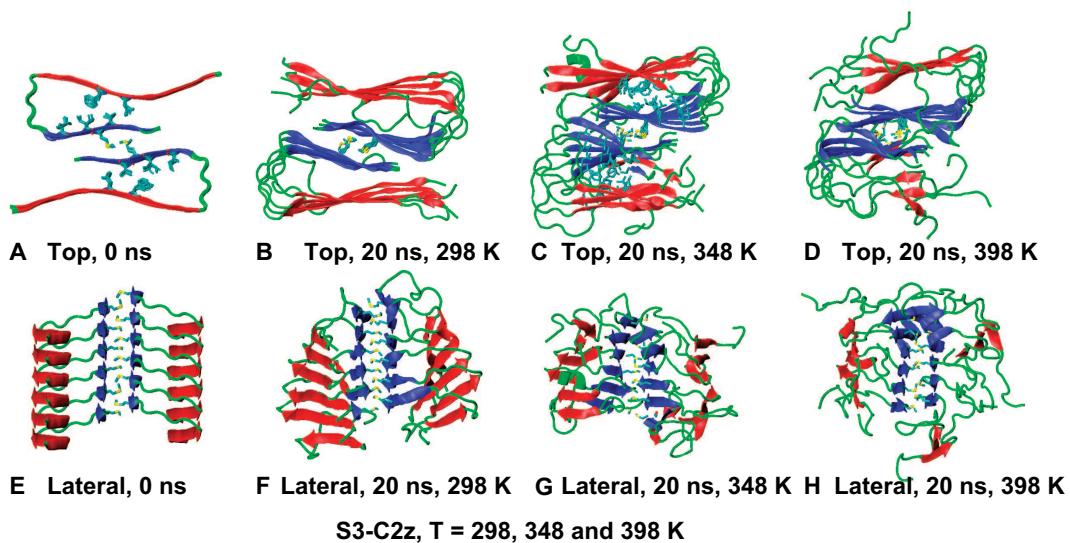


Fig. S 8: