## Destructive Mechanism of $A\beta_{1-42}$ Protofibril by Norepinephrine revealed via Molecular Dynamics Simulations

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This material contains four figures: the correlations between  $A\beta_{1-42}$  protofibril experimental NMR chemical shift ( $C_{\alpha}/C_{\beta}$  atoms) and simulated chemical shift data (Figure S1), the J-coupling ( ${}^{3}J(H_{N},H_{\alpha})$ ) constants values of  $A\beta_{42}$  residues calculated from 5OQV and those from simulationgenerated conformational ensembles of  $A\beta_{1-42}$  protofibril system (Figure S2), the secondary structure probabilities of  $A\beta_{1-42}$  protofibril in the  $A\beta$  and  $A\beta+NE^{0}$  systems (Figure S3), the average number of  $\pi$ - $\pi$  stacking between the aromatic residues of  $A\beta_{1-42}$  protofibril and NE<sup>+</sup>/NE<sup>0</sup> (Figure S4), and the interchain contact number of  $A\beta_{1-42}$  protofibril in the absence and presence of NE (in protonated and deprotonated states) (Figure S5).



**Figure S1.** Conformational comparison between experimental data and simulated data. (a–b) Correlations between experimental NMR chemical shift data for  $C_{\alpha}$  and  $C_{\beta}$  atoms and simulated chemical shift data for the A $\beta_{1-42}$  protofibril system.



**Figure S2.** The J-coupling  $({}^{3}J(H_{N},H_{\alpha}))$  constants values of A $\beta_{42}$  residues calculated from 5OQV and those from simulation-generated conformational ensembles of A $\beta_{1-42}$  protofibril system.



Figure S3. Secondary structure probabilities of  $A\beta_{1-42}$  protofibril in the  $A\beta$  and  $A\beta$ +NE<sup>0</sup> systems.



(a)

**Figure S4.** The LS-shaped configuration of  $A\beta_{1-42}$  protofibril shown in (a). The ABCD is used to indicate four chains of  $A\beta_{1-42}$  protofibril. (b) The interchain contact number of  $A\beta_{1-42}$  protofibril in the absence and presence of NE (in protonated and deprotonated states). The interchain contact were averaged over the three pairs of neighboring chains (i.e. chain A-chain B, chain B-chain C and chain C-chain D).



**Figure S5.** Average number of  $\pi$ - $\pi$  stacking between the aromatic residues of A $\beta_{1-42}$  protofibril and NE<sup>+</sup>/NE<sup>0</sup>.