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Supporting Information for the Manuscript

Interactions of aromatic residues in amyloids: A Survey of PDB Crystallographic Data

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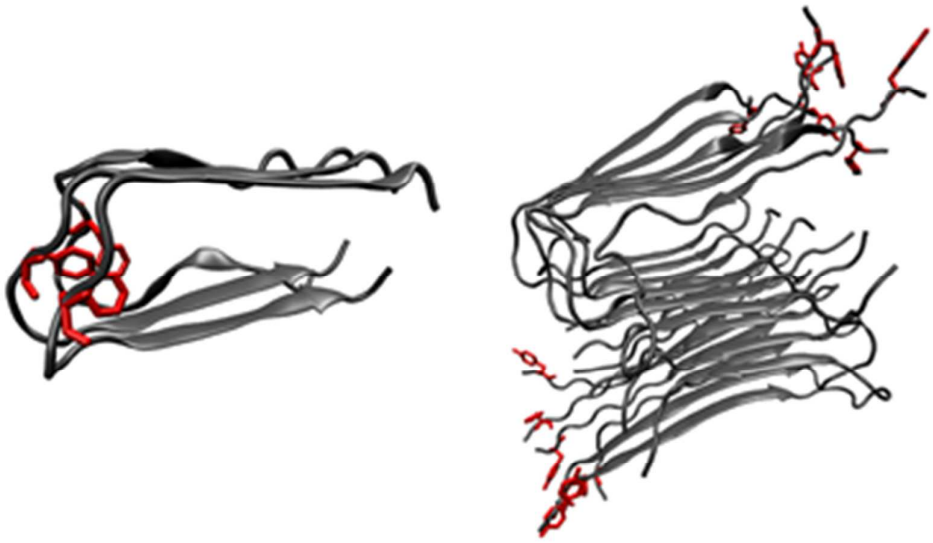

Fig. S1. Turn and coil aromatic amino acids (in red). PDBids: 2e8d, 2lmo.

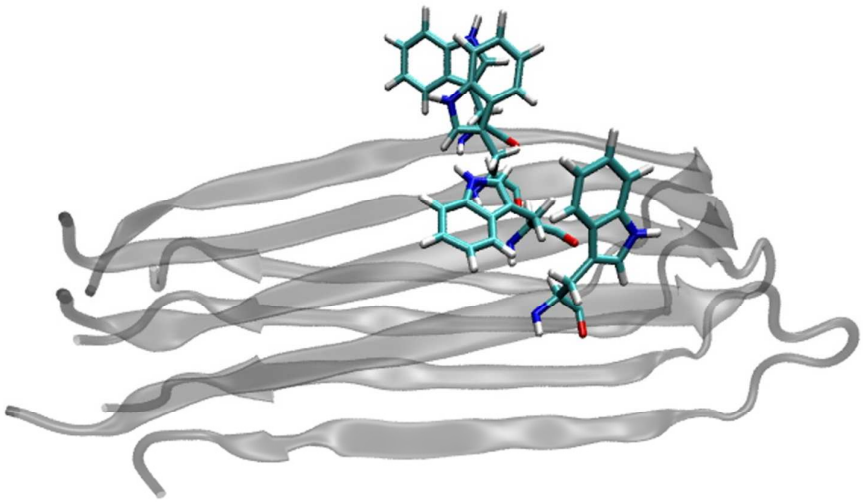

Fig. S2. The structure with Trp-Trp intrasheet interations, PDBid 2nnt.

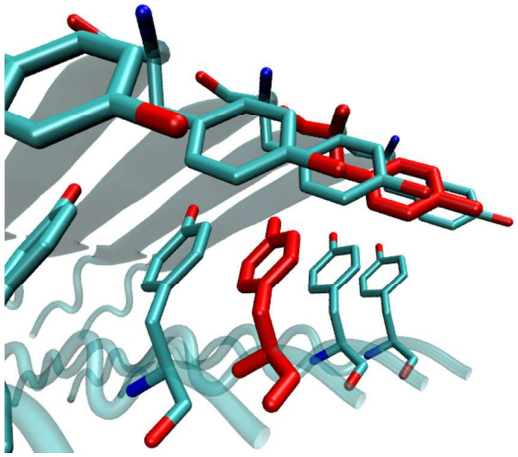

Fig. S3. The 20-framed NMR structure, PDBid 2m5n. An example of the high-offset interactions as a consequence of geometry, rings in red. P_1/P_2 = 36.09 °, R = 1.78 Å, r = 5.91 Å, d = 6.17 Å. Surrounding interactions are intrasheet aromatic-aromatic interactions.