

Supplementary Information

Alzheimer's Protective Cross-interaction Between Wild-type and A2T Variants Alters A β 42 Dimer Structure

PD^{1*}, AC¹, GB²

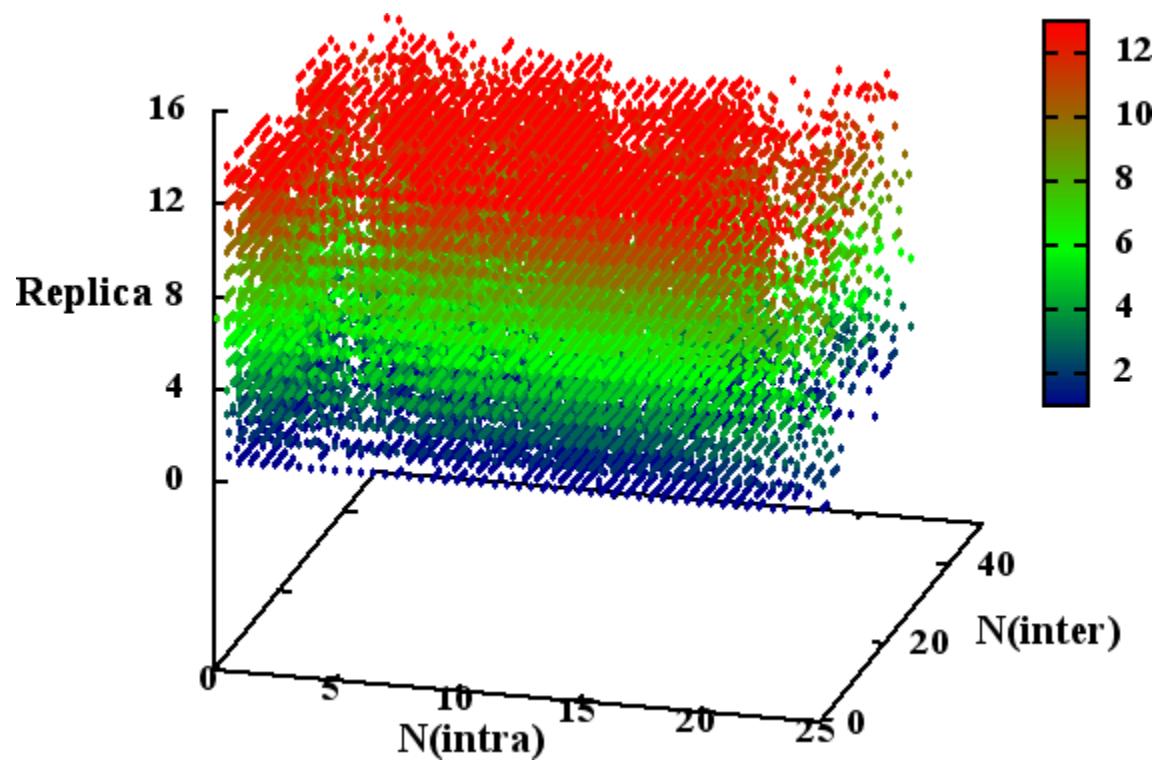


Figure S1. Conformational sampling as a function of intra- and inter-chain contacts, as obtained from the first 13 replicas for the heterodimer system.

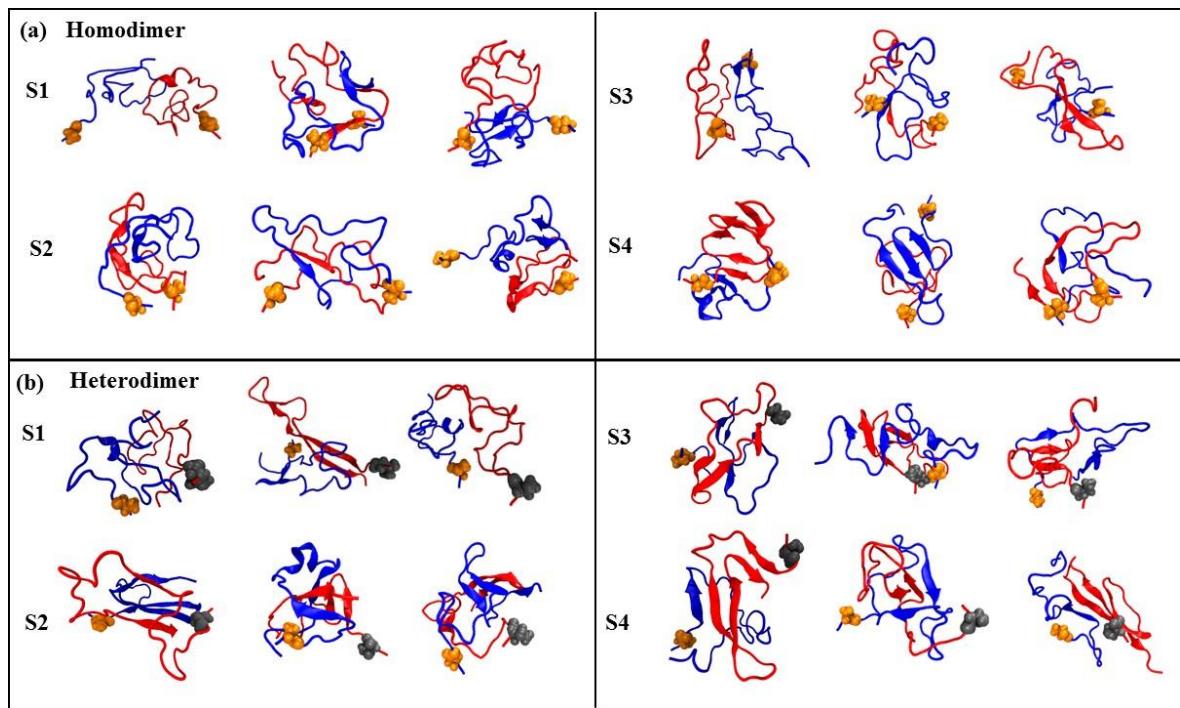


Figure S2: Representative structures of S1-S4 states (top panel: homodimer and bottom panel: heterodimer).

Table S1: Probability (in %) of inter-chain salt-bridges. Those with probability <5% are marked as “-“.

Homodimer		Heterodimer
D7-K28	6.3	-
E11-K28	8.3	-
D23-K28	-	7.8
K16-D23	-	6.0
D7-K16	-	6.8
R5-E11	-	5.7
D3-K28	-	5.2

Table S2: Average $\Delta G^{\text{Binding}_{\text{vdW}}}$ and $\Delta G^{\text{Binding}_{\text{elec}}}$ (in kcal/mol) of S1-S4 dimer structures.

		S1	S2	S3	S4
WT homodimer	$\Delta G^{\text{Binding}_{\text{vdW}}}$	-111.25	-138.32	-116.31	-95.90
	$\Delta G^{\text{Binding}_{\text{elec}}}$	-21.50	-29.51	-6.01	-19.01
A2T heterodimer	$\Delta G^{\text{Binding}_{\text{vdW}}}$	-102.47	-137.46	-122.55	-122.41
	$\Delta G^{\text{Binding}_{\text{elec}}}$	-25.43	-47.68	-46.70	-44.40

Table S3: Summary of cluster analysis: total % of population, number of clusters and size distribution of top 5 clusters for S1-S4 regions of homo- and heterodimer.

Homodimer	S1	S2	S3	S4
% of total population	9.40	22.75	19.75	22.37
# clusters	38	28	60	38
Size of top five clusters (%)	1. 21.64 2. 14.27 3. 11.33 4. 8.22 5. 8.00	1. 20.51 2. 15.17 3. 13.11 4. 12.70 5. 12.05	1. 9.97 2. 9.23 3. 9.14 4. 7.79 5. 7.14	1. 19.09 2. 12.14 3. 10.71 4. 9.31 5. 8.94
Heterodimer	S1	S2	S3	S4
% of total population	22.75	15.82	19.17	13.16
# clusters	42	54	55	31
Size of top five clusters (%)	1. 12.80 2. 10.60 3. 10.53 4. 8.93 5. 8.00	1. 28.41 2. 13.51 3. 9.56 4. 6.59 5. 5.39	1. 17.14 2. 12.47 3. 6.40 4. 6.05 5. 6.05	1. 22.20 2. 13.14 3. 12.99 4. 9.99 5. 9.41