Electronic Supplementary Information

Cholesterol-sensing role of phenylalanine in the interaction of human islet

amyloid polypeptide with lipid bilayers

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Fig. S1. Comparison between the kinetics of fibrillar assembly of hIAPP and hIAPP_{F15L} in 10 mM Tris-HCl buffer with 100 mM NaCl at pH 7.4 (A) and in 25 mM phosphate buffer with 50 mM NaCl at pH 7.4 (B) in the presence of DOPC and DOPC/20%Chol LUVs monitored by ThT fluorescence.



Fig. S2. CD spectra of hIAPP and hIAPP_{F15L} (15 μ M) at various incubation time recorded in phosphate buffer at pH 7.4, room temperature.

Peptide	Time	Secondary structure (%)				
	(min)	Helix	Strand	Turn	Unordere	
hIAPP	0	16.7	26.9	25.6	30.8	
	60	20.3	25.2	24.7	29.7	
	150	5.7	31.7	26.4	36.2	
hIAPP _{F15L}	0	10.1	32.7	23.2	34.1	
	60	14.3	31.4	21.0	33.0	
	210	7.4	38.3	21.0	33.2	

Table S1 Secondary structure data of hIAPP and hIAPP_{\rm F15L} in phosphate buffer at various time.



Fig. S3. CD spectra of hIAPP and hIAPP_{F15L} in phosphate buffer in the presence of DOPC, DOPC/20%Chol, DOPC/DPPC 1:2, DOPC/DPPC/Chol 1:2:1, DPPC and DPPC/20%Chol LUVs at various incubation time.

Table S2 Secondary structure data of hIAPP and $hIAPP_{F15L}$ in phosphate buffer in the presence of DOPC, DOPC/20%Chol, DOPC/DPPC 1:2, DOPC/DPPC/Chol 1:2:1, DPPC and DPPC/20%Chol LUVs at various incubation time.

Peptide	Lipid	Time (min)	Secondary structure (%)			
			Helix	Strand	Turn	Unordered
		0	16.1	31.7	21.8	30.4
	DOPC	15	7.8	33.8	26.5	31.9
hIAPP		30	6.8	37.5	34.2	31.6
		0	21.2	19.8	26.1	32.9
	DOPC/20%Chol	60	23.2	22.9	21.8	32.2
		120	17.4	29.9	21.0	31.7
hIAPP _{F15L}		0	19.2	26.4	21.8	32.6
	DOPC	15	14.5	32.6	21.1	31.8
		45	8.9	35.7	21.9	33.5
		0	18.9	26.6	22.4	32.1
	DOPC/20%Chol	90	15.3	24.5	24.3	35.8
		135	7.8	33.2	29.2	29.9
hIAPP		0	17.2	30.3	17.5	34.9
	DOPC/DPPC 1:2	15	17.5	26.5	19.4	36.7
		30	2.9	44.9	21.0	31.1
		0	37.7	11.1	22.1	29.1
	DOPC/DPPC/Chol 1:2:1	15	20.0	23.9	22.2	33.9
		30	7.1	38.4	23.9	30.8
hIAPP _{F15L}	DOPC/DPPC 1:2	0	7.3	36.7	22.0	34.2
		15	4.8	41.4	21.9	31.9
		45	4.5	40.2	21.4	33.9
		0	7.6	33.3	24.9	34.3
	DOPC/DPPC/Chol 1:2:1	45	7.1	30.3	24.9	37.6
		90	4.1	41.4	21.5	33.0
hIAPP		0	46.5	9.8	22.0	21.5
	DPPC	45	29.5	21.2	22.2	27.2
		60	14.2	34.7	22.3	28.9
		0	28.5	20.3	21.2	30.0
	DPPC/20%Chol	30	17.2	31.1	21.0	30.6
		60	14.2	34.7	22.3	28.9
hiapp _{fisl}		0	29.5	23.5	21.1	25.9
	DPPC	90	24.9	31.5	18.6	25.1
		120	5.3	42.5	22.4	29.9
		0	23.8	17.3	25.8	33.1
	DPPC/20%Chol	90	11.3	28.8	23.4	36.5
		105	8.6	36.9	25.5	29.1



Fig. S4. TEM images of hIAPP and $hIAPP_{1-19/F15L}$ measured after incubation in phosphate buffer for 30 min (A and B, respectively) and 5 h (C and D, respectively).



Fig. S5. ¹H-NMR signals in the regions of 6.95-7.40 ppm (aromatic protons of F15 and H18-H_{δ}) and 3.96-4.03 ppm (V17-H_{α}) used for calculation of dissociation constants of hIAPP₁₋₁₉ binding with DOPC (A) and DOPC/20%Chol (B) LUVs. The ¹H-NMR spectra were recorded in phosphate buffer at 25°C at various lipid-to-peptide ratios (L/P).



Fig. S6. ¹H-NMR signals in the regions of 6.95-7.03 ppm (H18-H_{δ}), 3.96-4.03 ppm (V17-H_{α}) and 2.97-3.02 ppm (K1-H_{ϵ}) used for calculation of dissociation constants of hIAPP_{1-19/F15L} binding with DOPC (A) and DOPC/20%Chol (B) LUVs. The ¹H-NMR spectra were recorded in phosphate buffer at 25°C at various lipid-to-peptide ratios (L/P).



Fig. S7. ¹H-NMR signals in the regions of 6.95-7.40 ppm (aromatic protons of F15 and H18-H_{δ}) and 3.96-4.03 ppm (V17-H_{α}) used for calculation of dissociation constants of hIAPP₁₋₁₉ binding with DPPC (A) and DPPC/20%Chol (B) LUVs. The ¹H-NMR spectra were recorded in phosphate buffer at 25°C at various lipid-to-peptide ratios (L/P).



Fig. S8. ¹H-NMR signals in the regions of 6.95-7.03 ppm (H18-H_{δ}), 3.96-4.03 ppm (V17-H_{α}) and 2.97-3.02 ppm (K1-H_{ϵ}) used for calculation of dissociation constants of hIAPP_{1-19/F15L} binding with DPPC (A) and DPPC/20%Chol (B) LUVs. The ¹H-NMR spectra were recorded in phosphate buffer at 25°C at various lipid-to-peptide ratios (L/P).