

Membrane-Fusogen Distance is Critical for Efficient Coiled-Coil-Peptide-Mediated Liposome Fusion

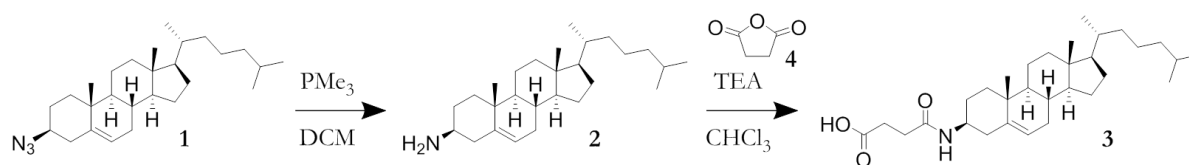
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Supporting Information

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Synthetic procedure for Cholesteryl-amino-hemisuccinate



Scheme S1. Synthetic procedure towards cholesteryl- 4-amino-4- oxobutanoic acid (3). 3-azido-5-cholestene (1) is reduced to 3-amino-5-cholestene (2), which in turn is condensed with succinic anhydride (4) to yield final cholesterol anchor 3.

3-Amino-5-cholestene (2): To a solution of 3-Azido-5-cholestene (1) (15.8 g, 38.4 mmol) in dichloromethane, 4 equivalents of trimethylphosphine (1M solution in toluene) was added and the reaction was stirred overnight. When 3-Azido-5-cholestene was consumed as evidenced by TLC (eluent: hexane), the reaction was concentrated *in vacuo* to yield 2 (14.8 g, 100%) as a pure compound. ¹H NMR (300 MHz, CDCl₃) δ 5.27 (s, 1H), 2.60 (m, 1H), 2.17-0.85 (m, 40H), 0.64 (s, 3H); ¹³C NMR (75 MHz, CDCl₃) δ 141.8, 120.7, 56.8, 56.2, 52.0, 50.3, 43.4, 42.3, 39.8, 39.5, 38.2, 36.6, 36.2, 35.8, 32.7, 31.9 (× 2), 28.3, 28.0, 24.3, 23.9, 22.8, 22.6, 21.0, 19.5, 18.7, 11.9; IR (film) ν 3354, 3260, 3154, 2936, 2896, 2849, 1464, 1437, 1381 cm⁻¹; LRMS (ESI⁺) *m/z* 386.4 ([M+H]⁺, C₂₇H₄₅N calcd. 386.4).

Cholesteryl-4-amino-4-oxobutanoic acid (3): To a solution of 2 (10g, 26 mmol) in chloroform (150mL), 4 (7.8g, 78 mmol) was added. Triethylamine (8mL, 57 mmol) was added drop-wise to the solution and the reaction mixture was stirred overnight under nitrogen atmosphere at room temperature, until the starting material was completely consumed as evidenced by TLC (eluent: hexane: ethylacetate 1:1). The reaction mixture was transferred to a separatory funnel, and the organic layer was washed with water and brine, was dried with MgSO₄ and was concentrated *in vacuo* to afford cholesterol amide acid (10g) as a yellow solid. The crude cholesterol amide acid was purified by recrystallization in Acetonitrile: MeOH (1:1) at 4°C and was vacuum filtered to afford 3 as an off-white solid (8.7g, 76%) ¹H NMR (400MHz, CDCl₃) δ 5.36 (s, 1H), 3.69 (s, 1H), 3.60 (m, 1H) 2.62 (m, 2H), 2.44 (m, 2H), 2.20-1.72 (m, 7H), 1.67-1.03 (m, 20H) 1.01 (s, 3H) 0.92 (d, *J* = 6.5 Hz, 3H), 0.87 (dd, *J* = 6.6, 1.7 Hz, 6H), 0.69 (s, 3H). TLC-MS (ESI⁺) *m/z* 486.4 ([M+H]⁺, C₃₁H₅₁NO₃ calcd. 486.4 g/mol).

Lipopeptide analysis

Purity was confirmed using LC-MS analysis equipped with Gemini 3 μ C18 column coupled with Finnigan LCQ advantage max (Thermo) ESI-MS analyzer and the results are summarized in Table S1. LCMS eluents were A) H₂O + 0.1% v/v TFA, B) MeCN + 0.1 v/v TFA, with a flow rate of 1 mL min⁻¹. Gradient was applied between 2 and 12 min, from 10% B in A to 90% B in A.

Table S1. Formulae and masses of synthesized lipopeptides.

peptide	formula	Calc. mass	Found mass
CP ₀ E ₃	C ₁₄₄ H ₂₃₆ N ₂₆ O ₄₀	2971.57	2972.14, [M+H] ⁺
CP ₀ K ₃	C ₁₄₉ H ₂₅₂ N ₃₀ O ₃₃	2991.78	2992.60, [M+H] ⁺
CP ₄ E ₃	C ₁₅₄ H ₂₅₅ N ₂₇ O ₄₅	3204.83	3206.63, [M+H] ⁺
CP ₄ K ₃	C ₁₅₉ H ₂₇₁ N ₃₁ O ₃₈	3225.04	1611.48, [M+2H] ⁺²
CP ₈ E ₃	C ₁₆₃ H ₂₇₃ N ₂₇ O ₄₉	3395.06	1696.40, [M+2H] ⁺²
CP ₈ K ₃	C ₁₆₈ H ₂₈₉ N ₃₁ O ₄₂	3415.28	1706.40, [M+2H] ⁺²
CP ₁₂ E ₃	C ₁₇₁ H ₂₈₉ N ₂₇ O ₅₃	3571.27	1190.53, [M+3H] ⁺³
CP ₁₂ K ₃	C ₁₇₆ H ₃₀₅ N ₃₁ O ₄₆	3591.49	1197.07, [M+3H] ⁺³
CP ₁₆ E ₃	C ₁₇₉ H ₃₀₅ N ₂₇ O ₅₇	3747.48	1249.27, [M+3H] ⁺³
CP ₁₆ K ₃	C ₁₈₄ H ₃₂₁ N ₃₁ O ₅₀	3767.70	1255.93, [M+3H] ⁺³
LP ₀ E ₃	C ₁₄₇ H ₂₅₆ N ₂₆ O ₄₃ P ⁻	3106.78	1563.44, [M+H+NH ₄] ⁺²
LP ₀ K ₃	C ₁₅₀ H ₂₇₁ N ₂₉ O ₃₇ P ⁻	3103.96	1553.01, [M+2H] ⁺²
LP ₂ E ₃	C ₁₅₃ H ₂₆₇ N ₂₇ O ₄₆ P ⁻	3251.94	1084.98, [M+3H] ⁺³
LP ₂ K ₃	C ₁₅₆ H ₂₈₂ N ₃₀ O ₄₀ P ⁻	3249.12	1084.03, [M+3H] ⁺³
LP ₄ E ₃	C ₁₅₇ H ₂₇₅ N ₂₇ O ₄₈ P ⁻	3340.04	1114.33, [M+3H] ⁺³
LP ₄ K ₃	C ₁₆₀ H ₂₉₀ N ₃₀ O ₄₂ P ⁻	3337.22	1113.72, [M+3H] ⁺³
LP ₈ E ₃	C ₁₆₆ H ₂₉₃ N ₂₇ O ₅₂ P ⁻	3530.28	1178.04, [M+3H] ⁺³
LP ₈ K ₃	C ₁₆₉ H ₃₀₈ N ₃₀ O ₄₆ P ⁻	3527.46	1177.09, [M+3H] ⁺³
LP ₁₂ E ₃	C ₁₇₄ H ₃₀₉ N ₂₇ O ₅₆ P ⁻	3706.49	1236.74, [M+3H] ⁺³
LP ₁₂ K ₃	C ₁₇₇ H ₃₂₄ N ₃₀ O ₅₀ P ⁻	3703.67	1235.79, [M+3H] ⁺³

Full data of 30min content mixing assay

Content mixing experiments were followed for 30 minutes in a continuous fashion and showed characteristic 1st order kinetics in the initial minutes (*Figure S1, and insets*). Therefore, fluorescence increase after 4 minutes is used to estimate the relationship of L_{PEG} and the fusion efficiency. Error bars and absolute values after 4 and 30 minutes were obtained by at least 3 individual experiments and are shown in Figure S, as a function of m , n . In representative negative control experiments, peptide-containing liposomes (m , $n = 4$ or 8) were mixed with plain liposomes and the fluorescence increase was recorded for 30 minutes.

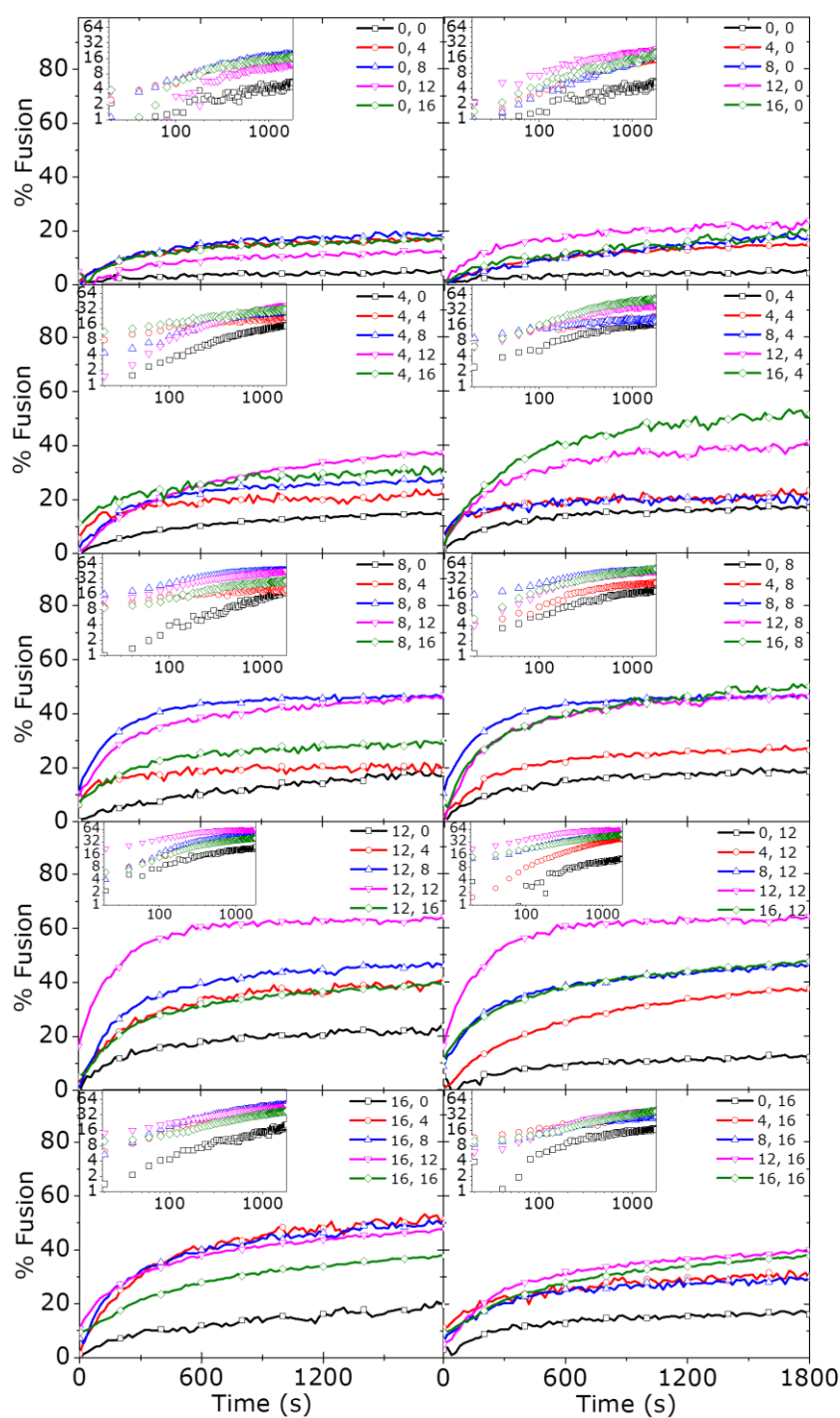


Figure S1. Content mixing of CP_mE_3 and CP_nK_3 decorated liposomes as a function of m , n . Insets with log scale illustrate 1st order kinetics until at least the 4th minute. [Total lipid] = 0.1 mM with 1 mol% lipopeptide, PBS pH 7.4.

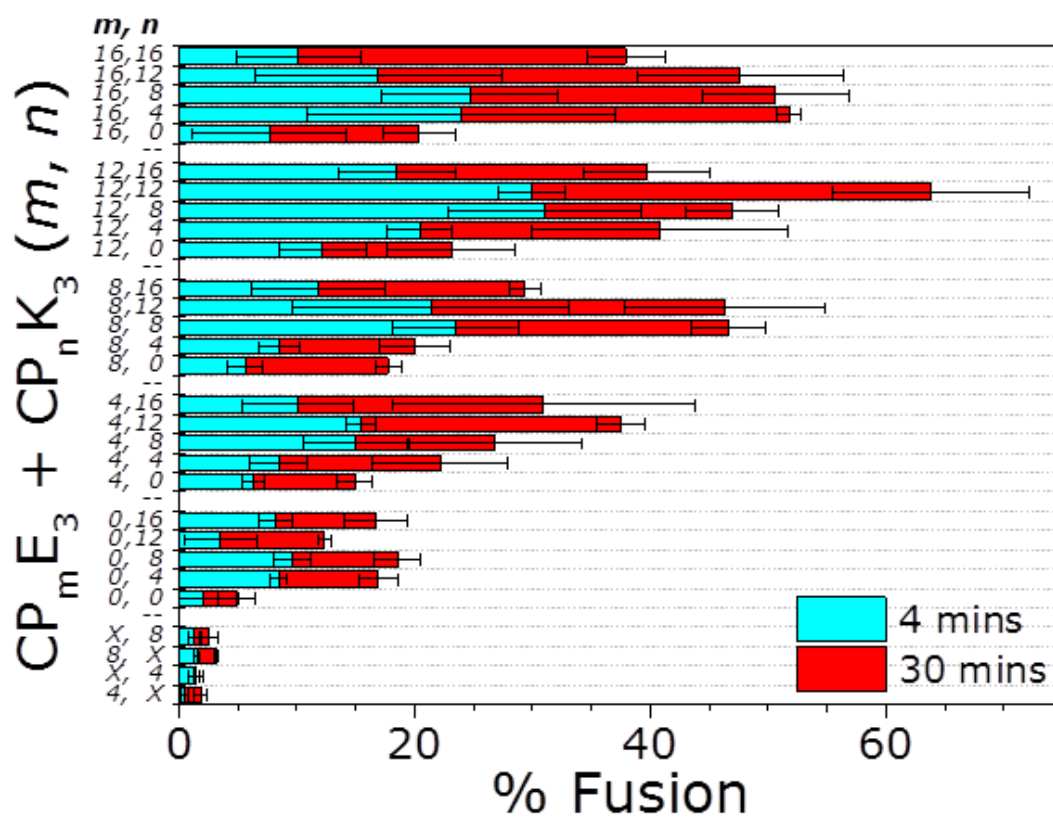


Figure S2. Time resolved rate of fusion (content mixing) with lipopeptide CP_mE_3 or CP_nK_3 decorated liposomes as a function of m, n , indicated by an increase in sulforhodamine B emission after 4 and 30 minutes. For control experiments the omitted peptide-functionalized liposomes are represented by X . [Total lipid] = 0.1 mM with 1 mol% lipopeptide, in PBS pH 7.4.

Influence of CPE / CPK ratio on fusion efficiency

A change in peptide functionalization ratio reveals the influence of peptide equilibria on the fusion process. Again, results are shown as a function of m, n . (Figure S2)

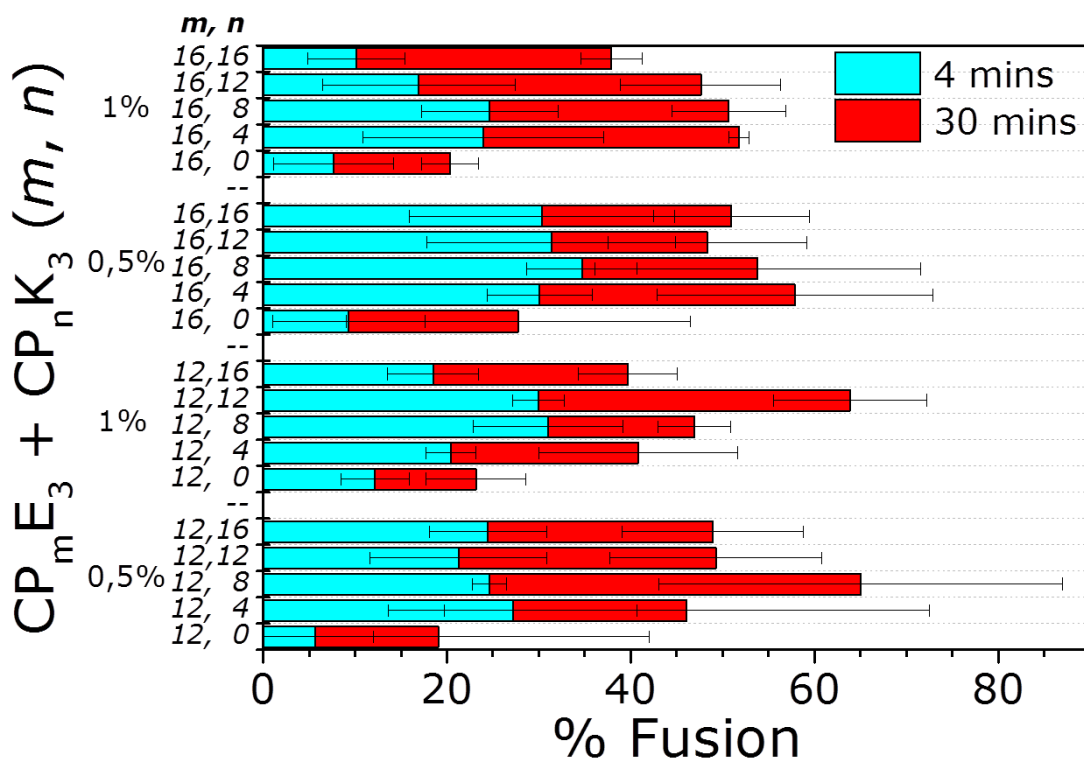


Figure S2. Time resolved rate of fusion (content mixing) with peptide CP_mE_3 and CP_nK_3 decorated liposomes as a function of m, n , and $[CP_mE_3]$, as indicated by an increase in sulforhodamine B emission after 4 and 30 minutes. $[Total\ lipid] = 0.1\ mM$ with $1\ mol\% CP_nK_3$ and $1\ mol\%$ or $0.5\ mol\% CP_mE_3$, PBS pH 7.4.

CD spectroscopy

Characteristics ($\theta_{222}/\theta_{208}$ and percentage helicity at θ_{222}) of the spectra as summarized in Table 2, are supported with the recorded spectra between 207nm and 250 nm (Figure S3). Below 207 nm, spectra were too noisy due to scattering of the signal by liposomes.

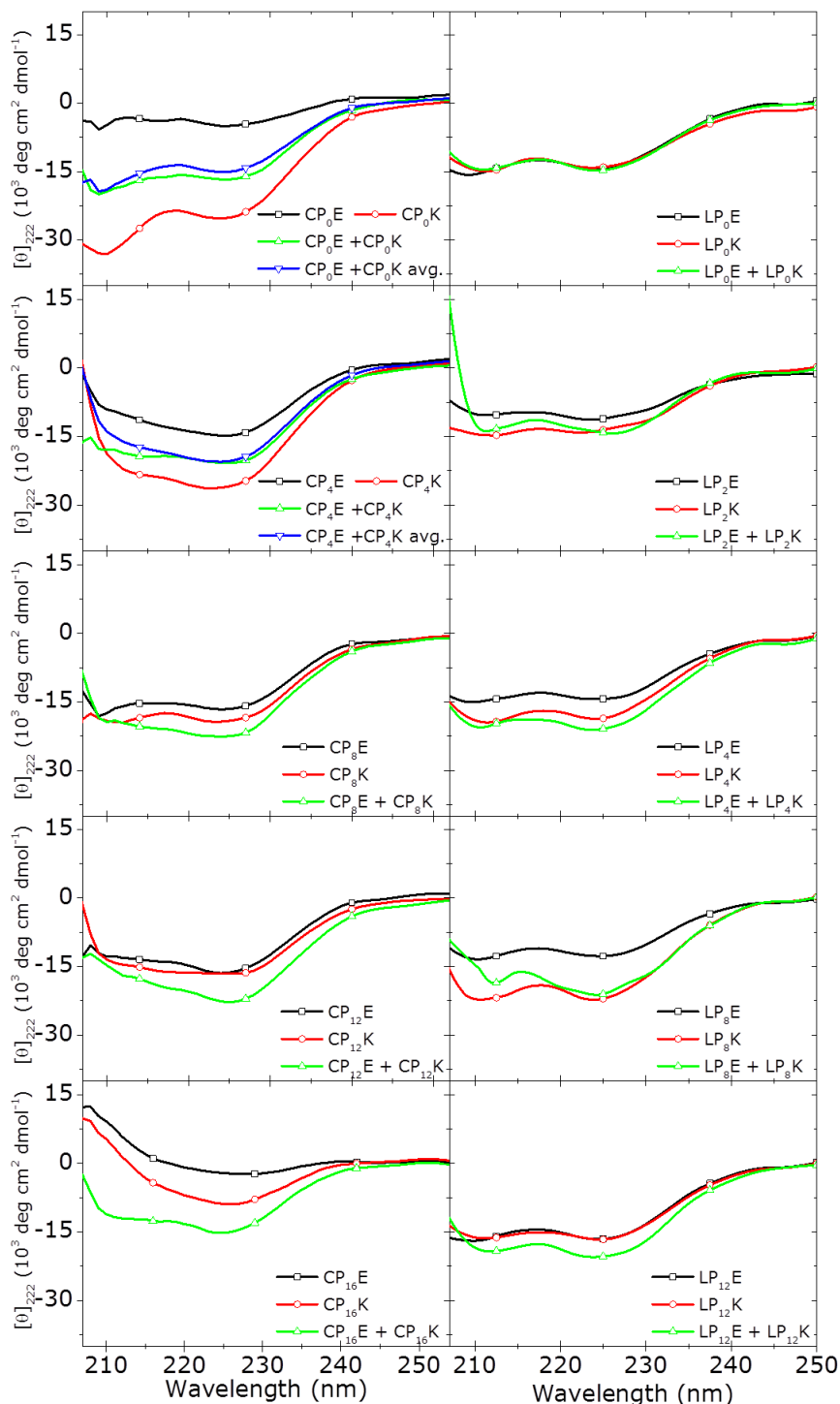


Figure S3. Peptide secondary structure of E- (black) and K- (red) decorated liposomes before and after mixing (green), measured by circular dichroism. Left series is with cholesterol anchor, right series contain a DOPE anchor. Only symmetric pairs ($m = n$) are used in this study. With CP_0 and CP_4 anchor coiled coil formation is not evident, so the calculated average of the individual E_3 and K_3 spectra is added (blue). [Total lipid] = 0.5 mM with 1mol% lipopeptide, PBS pH 7.4, and cuvette path length of 5mm.