

## **An Orthogonal Array Optimization of Lipid-like Nanoparticles for mRNA Delivery in Vivo**

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### General synthetic procedures and spectral data:

*Synthesis of compound 3.* To a solution of diamine (75 mmol) in  $\text{CHCl}_3$  (70 mL) was added a solution of  $\text{Boc}_2\text{O}$  (14.4 mmol) in  $\text{CHCl}_3$  (30 mL) via an additional funnel over 2.5 h. The resulting suspension was stirred and 100 mL of  $\text{NaHCO}_3$  (1N) was slowly added to form a bi-layer solution. The organic layer was washed with 100 mL of 1N  $\text{NaHCO}_3$  and 20 mL of brine, and then dried over solid  $\text{MgSO}_4$  for 2 h. The solution was then filtered, evaporated and dried under high vacuum in order to afford compound **3**. Yield: n=2, 91%; n=3, commercially available reagent; n=4, 63%; n=5, 95%; n=6, commercially available reagent; n=7, 95%; n=8, 86%.

*Synthesis of compound 4.*<sup>1</sup> To a solution of compound **2** (1.88 mmol) in  $\text{CH}_2\text{Cl}_2$  (30 mL) was added 10 mL of pyridine and cooled in an ice bath. A solution of compound **3** (7.52 mmol) in  $\text{CH}_2\text{Cl}_2$  (30 mL) was added dropwise with stirring. The reaction mixture was then allowed to warm to RT, diluted with 100 mL of  $\text{CH}_2\text{Cl}_2$  and washed twice with 50 mL of water, 50 mL of saturated  $\text{NaHCO}_3$ , and 50 mL of brine. The solution was dried over solid  $\text{MgSO}_4$  for 2 h and concentrated. The residue was purified by column chromatography using a CombiFlash Rf system with a RediSep Gold Resolution silica column (Teledyne Isco) with gradient elution from 100%  $\text{CH}_2\text{Cl}_2$  to  $\text{CH}_2\text{Cl}_2/\text{MeOH}/\text{NH}_4\text{OH}$  (75/22/3 by volume) to give compound **4**. Yield: n=2, 74%; n=3, 68%; n=4, 65%; n=5, 69%; n=6, 81%; n=7, 37%; n=8, 61%.

*Synthesis of compound 1.* To a suspension of compound **4** (1.41 mmol) in  $\text{CH}_2\text{Cl}_2$  (20 mL) was added trifluoroacetic acid (TFA, 1.41 mmol). The mixture was stirred at RT for 1 h and monitored with thin layer chromatography (TLC). Upon completion of the reaction, the solvent was evaporated and the residue was dissolved in MeOH and concentrated. After solidification in EtOAc, compound **1** was dried under an oil pump and afforded in quantitative yield.

<sup>1</sup>H NMR spectra were recorded at 300 or 400 MHz on the Bruker instrument. <sup>1</sup>H NMR chemical shifts were reported as  $\delta$  values in ppm relative to TMS. Mass spectra were obtained on a Micromass Q-TOF micro Mass Spectrometer.

*N*<sup>1</sup>,*N*<sup>3</sup>,*N*<sup>5</sup>-tris(2-(didodecylamino)ethyl)benzene-1,3,5-tricarboxamide (**TT2**): yield (82%). <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>,  $\delta$ ) 8.88 (3H, br), 8.56 (3H, s), 3.81 (6H, m), 3.32 (6H, m), 3.04-3.08 (12H, m), 1.69 (12H, s), 1.20-1.30 (108H, m), 0.88 (18H, tri, *J* = 6.9 Hz). MS (*m/z*): [M + H]<sup>+</sup> calcd. for C<sub>87</sub>H<sub>169</sub>N<sub>6</sub>O<sub>3</sub>, 1346; found, 1346.

*N*<sup>1</sup>,*N*<sup>3</sup>,*N*<sup>5</sup>-tris(3-(didodecylamino)propyl)benzene-1,3,5-tricarboxamide (**TT3**): yield (50%). <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>,  $\delta$ ) 8.53 (3H, br), 8.40 (3H, s), 3.57 (6H, m), 2.67-3.60 (6H, tri, *J* = 5.4 Hz), 2.58-2.48 (12H, tri, *J* = 7.5 Hz), 1.90-1.70 (12H, m), 1.57-1.38 (12H, m), 1.35-1.17 (96H, m), 0.89 (18H, tri, *J* = 6.9 Hz). MS (*m/z*): [M + H]<sup>+</sup> calcd. for C<sub>90</sub>H<sub>175</sub>N<sub>6</sub>O<sub>3</sub>, 1388; found, 1388.

*N*<sup>1</sup>,*N*<sup>3</sup>,*N*<sup>5</sup>-tris(4-(didodecylamino)butyl)benzene-1,3,5-tricarboxamide (**TT4**): yield (31%). <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>,  $\delta$ ) 8.33 (3H, s), 7.39 (3H, br), 3.5-3.42 (6H, m), 2.65-2.46 (18H, m), 1.75-1.55 (12H, m), 1.52-1.37 (12H, m), 1.35-1.17 (102H, m), 0.90 (18H, tri, *J* = 6.9 Hz). MS (*m/z*): [M + H]<sup>+</sup> calcd. for C<sub>93</sub>H<sub>181</sub>N<sub>6</sub>O<sub>3</sub>, 1430; found, 1430.

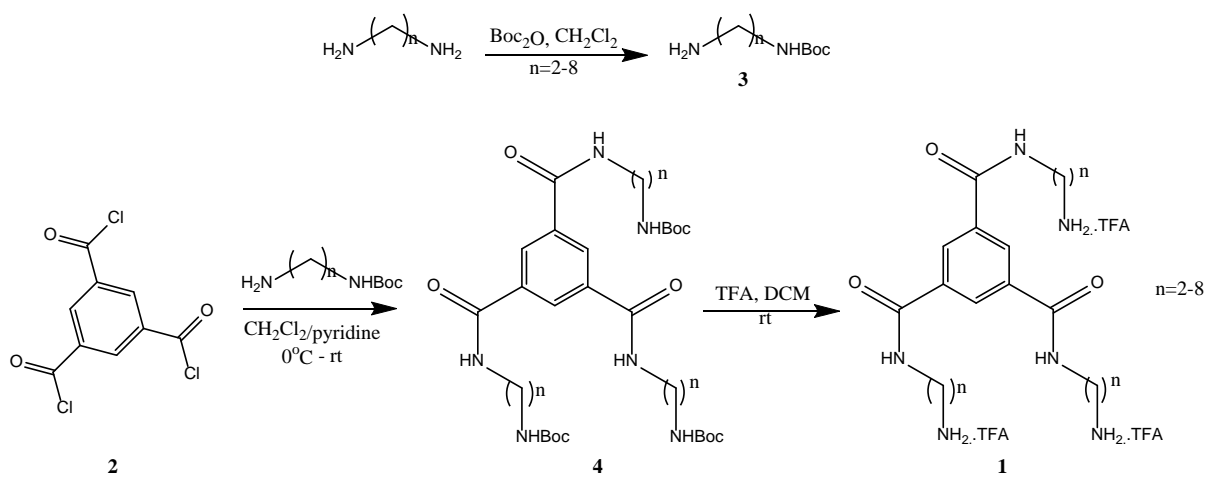
*N*<sup>1</sup>,*N*<sup>3</sup>,*N*<sup>5</sup>-tris(5-(didodecylamino)pentyl)benzene-1,3,5-tricarboxamide (**TT5**): yield (45%). <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>,  $\delta$ ) 8.40 (3H, s), 6.62 (3H, s), 3.56-3.45 (6H, m), 2.47-2.39 (18H, m), 1.76-1.65 (18H, m), 1.52-1.37 (24H, m), 1.35-1.20 (90H, m), 0.89 (18H, tri, *J* = 6.9 Hz). MS (*m/z*): [M + H]<sup>+</sup> calcd. for C<sub>96</sub>H<sub>187</sub>N<sub>6</sub>O<sub>3</sub>, 1472; found, 1472.

*N*<sup>1</sup>,*N*<sup>3</sup>,*N*<sup>5</sup>-tris(6-(didodecylamino)hexyl)benzene-1,3,5-tricarboxamide (**TT6**): yield (53%). <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>,  $\delta$ ) 8.38 (3H, s), 6.52 (3H, br), 3.52-3.45 (6H, m), 2.47-2.39 (18H, m), 1.72-1.63 (18H, m), 1.54-1.36 (30H, m), 1.32-1.21 (90H, m), 0.90 (18H, tri, *J* = 6.9 Hz). MS (*m/z*): [M

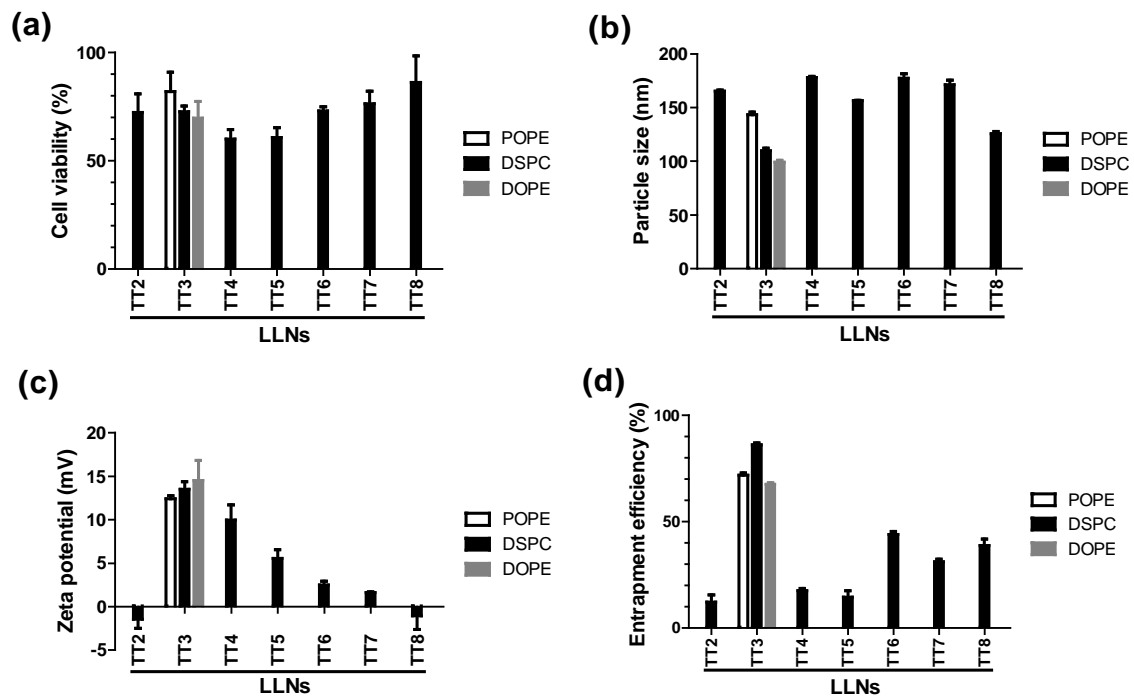
+ H]<sup>+</sup> calcd. for C<sub>99</sub>H<sub>193</sub>N<sub>6</sub>O<sub>3</sub>, 1515; found, 1515.

*N*<sup>1</sup>,*N*<sup>3</sup>,*N*<sup>5</sup>-tris(7-(didodecylamino)heptyl)benzene-1,3,5-tricarboxamide (**TT7**): yield (75%). <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>, δ) 8.40 (3H, s), 6.72 (3H, s), 3.48 (6H, m), 2.50 (18H, m), 1.70-1.56 (6H, m), 1.55-1.34 (30H, m), 1.37-1.27 (108H, m), 0.89 (18H, tri, *J* = 6.9 Hz). MS (*m/z*): [M + H]<sup>+</sup> calcd. for C<sub>102</sub>H<sub>199</sub>N<sub>6</sub>O<sub>3</sub>, 1557; found, 1557.

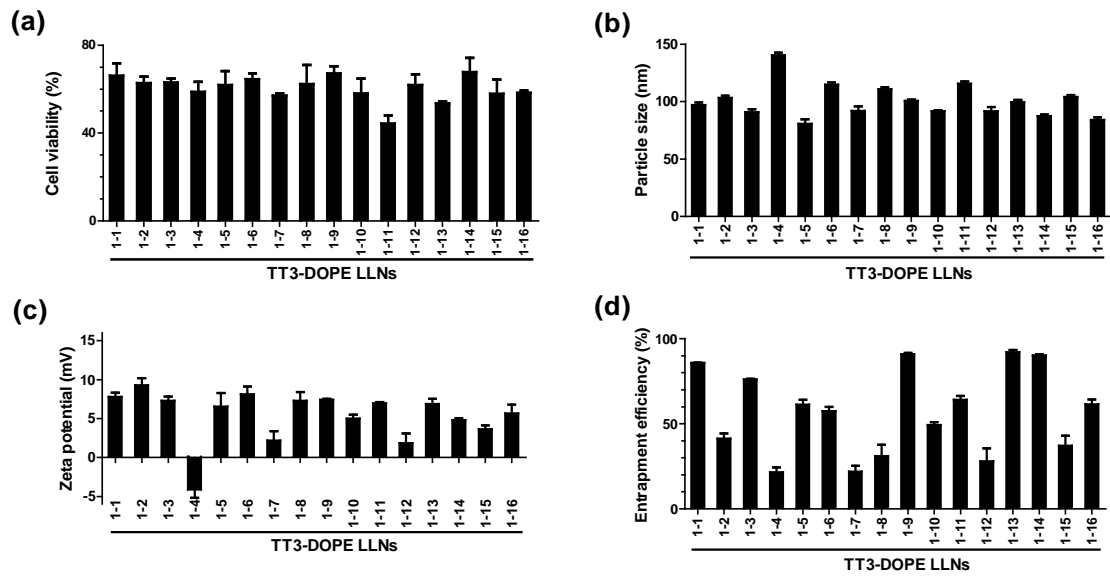
*N*<sup>1</sup>,*N*<sup>3</sup>,*N*<sup>5</sup>-tris(8-(didodecylamino)octyl)benzene-1,3,5-tricarboxamide (**TT8**): yield (quantitative). <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>, δ) 8.40 (3H, s), 6.86 (3H, br), 3.47 (6H, m), 2.82-2.52 (18H, m), 1.73-1.42 (30H, m), 1.42-1.17 (120H, m), 0.90 (18H, tri, *J* = 6.9 Hz). MS (*m/z*): [M + H]<sup>+</sup> calcd. for C<sub>105</sub>H<sub>205</sub>N<sub>6</sub>O<sub>3</sub>, 1599; found, 1599.



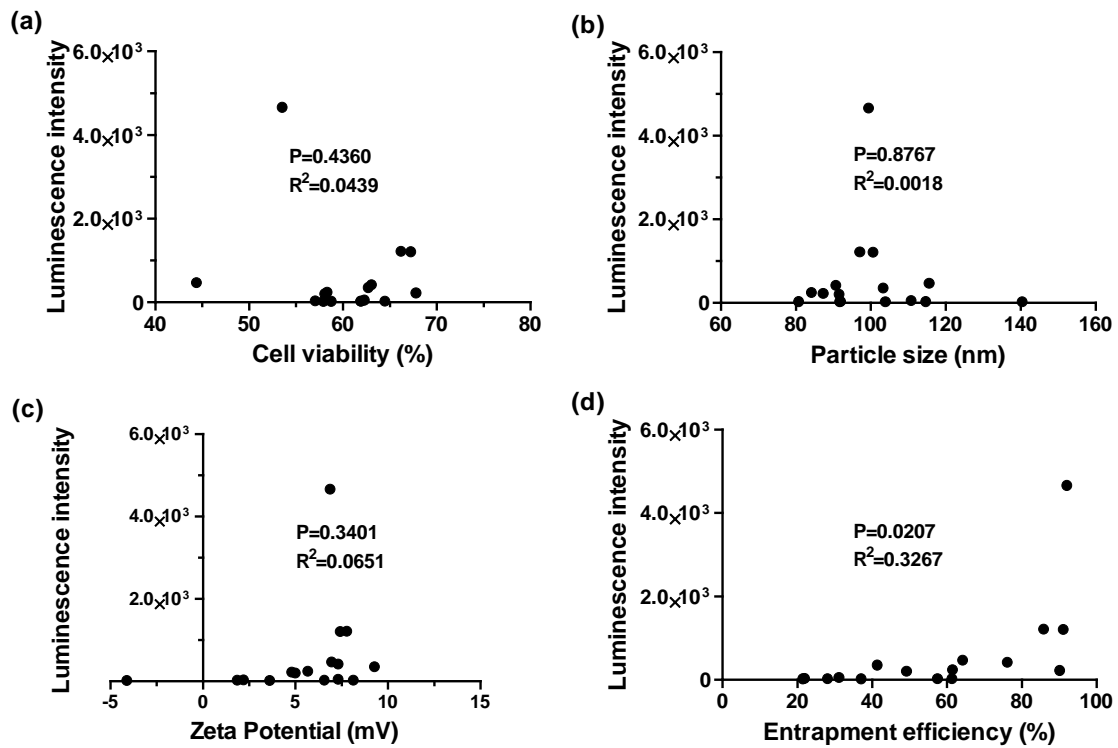
**Figure S1.** The synthetic route to compound **1**. Benzene-1,3,5-tricarbonyl trichloride (**2**) was reacted with Boc-protected diamine (**3**) to produce the intermediates (**4**). Deprotection of **4** gave compound **1**.



**Figure S2.** The cell viability (a), particle size (b), zeta potential (c), and entrapment efficiency (d) of LLNs.

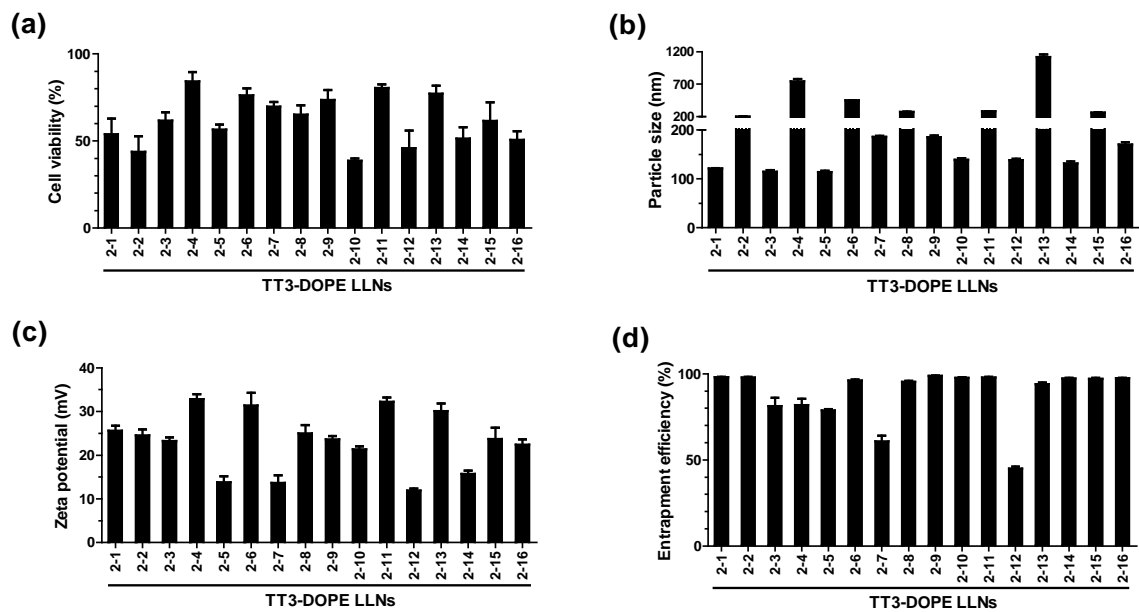


**Figure S3.** The cell viability (a), particle size (b), zeta potential (c), and entrapment efficiency (d) of TT3-DOPE LLNs formulated through the first orthogonal array.

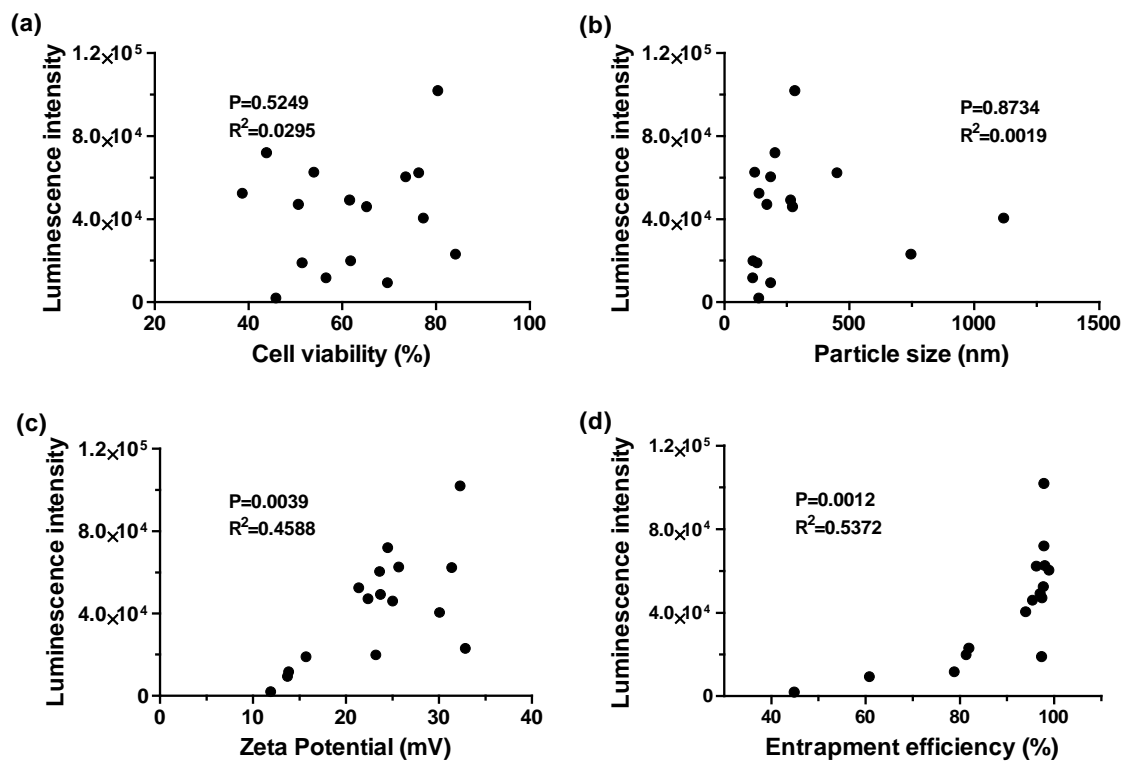


**Figure S4.** The correlation between transfection efficiency of TT3-DOPE LLNs formulated through the first orthogonal array and corresponding cell viability, particle size, zeta potential, and entrapment efficiency.

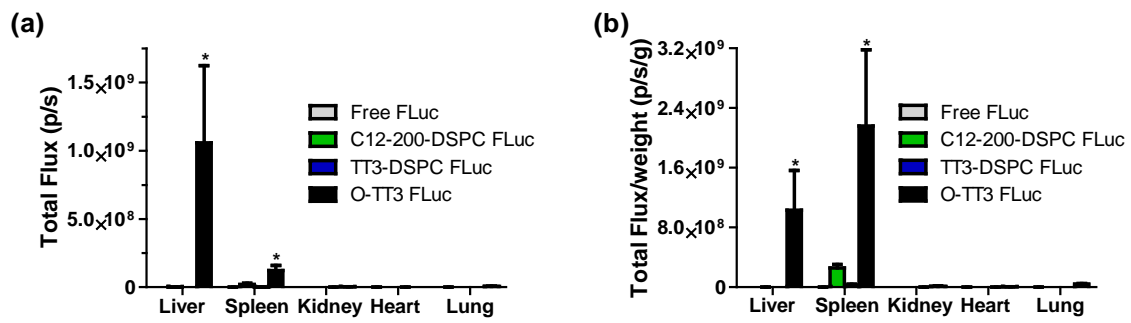




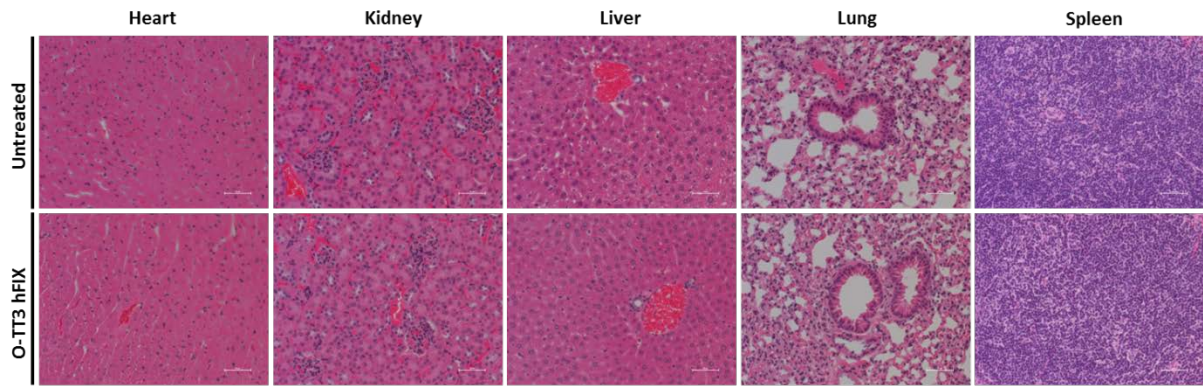
**Figure S5.** The cell viability (a), particle size (b), zeta potential (c), and entrapment efficiency (d) of TT3-DOPE LLNs formulated through the second round of orthogonal array.



**Figure S6.** The correlation between transfection efficiency of TT3-DOPE LLNs formulated through the second round of orthogonal array and their corresponding cell viability, particle size, zeta potential, and entrapment efficiency.



**Figure S7.** Bioluminescence signal (6 hours after administration in C57BL/6 mice). Free FLuc mRNA served as a negative control. Total bioluminescence signal (a) and normalized bioluminescence signal with tissue weight (b). Statistically significant difference was observed in the groups (\*:  $P < 0.05$ ; O-TT3 FLuc vs C12-200-DSPC FLuc; O-TT3 FLuc vs TT3-DSPC FLuc).



**Figure S8.** Histological analysis of major organs (heart, kidney, liver, lung, and spleen) after treatment of **O-TT3** hFIX. Untreated groups served as negative controls. No significant alteration of histology was observed in the **O-TT3** LLNs treated groups compared to the control groups.

**Table S1.** Orthogonal array table  $L_{16}(4^4)$  (a) and  $K_n$  values (b) for the first round of **TT3**-DOPE LLNs.

(a) Orthogonal array table  $L_{16}(4^4)$

TT3-DOPE LLNs	Formulation components (mole ratio)				Relative luminescence intensity
	TT3	DOPE	Cholesterol	DMG-PEG <sub>2000</sub>	
1-1	30	2.5	38.5	3	1216
1-2	40	10	18.5	1.5	351
1-3	50	10	38.5	6	418
1-4	60	2.5	18.5	0.75	20
1-5	30	5	18.5	6	28
1-6	40	1.25	38.5	0.75	28
1-7	50	1.25	18.5	3	34
1-8	60	5	38.5	1.5	51
1-9	30	1.25	48.5	1.5	1207
1-10	40	5	28.5	3	203
1-11	50	5	48.5	0.75	469
1-12	60	1.25	28.5	6	30
1-13	30	10	28.5	0.75	4663
1-14	40	2.5	48.5	6	220
1-15	50	2.5	28.5	1.5	20
1-16	60	10	48.5	3	243

(b)  $K_n$  values

	TT3	DOPE	Cholesterol	DMG-PEG <sub>2000</sub>
$K_1^*$	<b>1779</b>	325	108	<b>1295</b>
$K_2^*$	201	369	<b>1229</b>	407
$K_3^*$	235	188	428	424
$K_4^*$	86	<b>1419</b>	535	174
$\Delta K^{**}$	1693	1231	1121	1121

$$K_n^* = \Sigma RLI_n/4$$

$$\Delta K^{**} = K_{\max} - K_{\min}$$

The impact of the four components on mRNA delivery ( $\Delta K$ ): **TT3** > DOPE > Cholesterol = DMG-PEG<sub>2000</sub>.

**Table S2.** Orthogonal array table  $L_{16}(4^4)$  (a) and  $K_n$  values (b) for the second round of **TT3**-DOPE LLNs.

(a) Orthogonal array table  $L_{16}(4^4)$

TT3-DOPE LLNs	Formulation components (mole ratio)				Relative luminescence intensity
	TT3	DOPE	Cholesterol	DMG-PEG <sub>2000</sub>	
2-1	10	20	35	0.15	62542
2-2	20	40	25	0.03	71935
2-3	30	40	35	0.75	19869
2-4	40	20	25	0	23096
2-5	10	30	25	0.75	11687
2-6	20	10	35	0	62216
2-7	30	10	25	0.15	9394
2-8	40	30	35	0.03	45962
2-9	10	10	40	0.03	60333
2-10	20	30	30	0.15	52468
2-11	30	30	40	0	101863
2-12	40	10	30	0.75	1917
2-13	10	40	30	0	40410
2-14	20	20	40	0.75	18962
2-15	30	20	30	0.03	49102
2-16	40	40	40	0.15	47049

(b)  $K_n$  values

	TT3	DOPE	Cholesterol	DMG-PEG <sub>2000</sub>
$K_1^*$	43743	33465	29028	<b>56896</b>
$K_2^*$	<b>51395</b>	38426	35974	56833
$K_3^*$	45057	<b>52995</b>	47647	42863
$K_4^*$	29506	44816	<b>57052</b>	13109
$\Delta K^{**}$	21889	19529	18025	43788

$$K_n^* = \Sigma RLI_n/4$$

$$\Delta K^{**} = K_{\max} - K_{\min}$$

The most efficient formulation in the second round of optimization was validated with a designated code of **Hi-TT3** LLNs (formulation ratio is **TT3**/DOPE/Chol/DMG-PEG<sub>2000</sub>=20/30/40/0).

## References

1. Matsuura, K.; Murasato, K.; Kimizuka, N. *J. Am. Chem. Soc.* **2005**, *127*, 10148-10149.