

Supporting Information

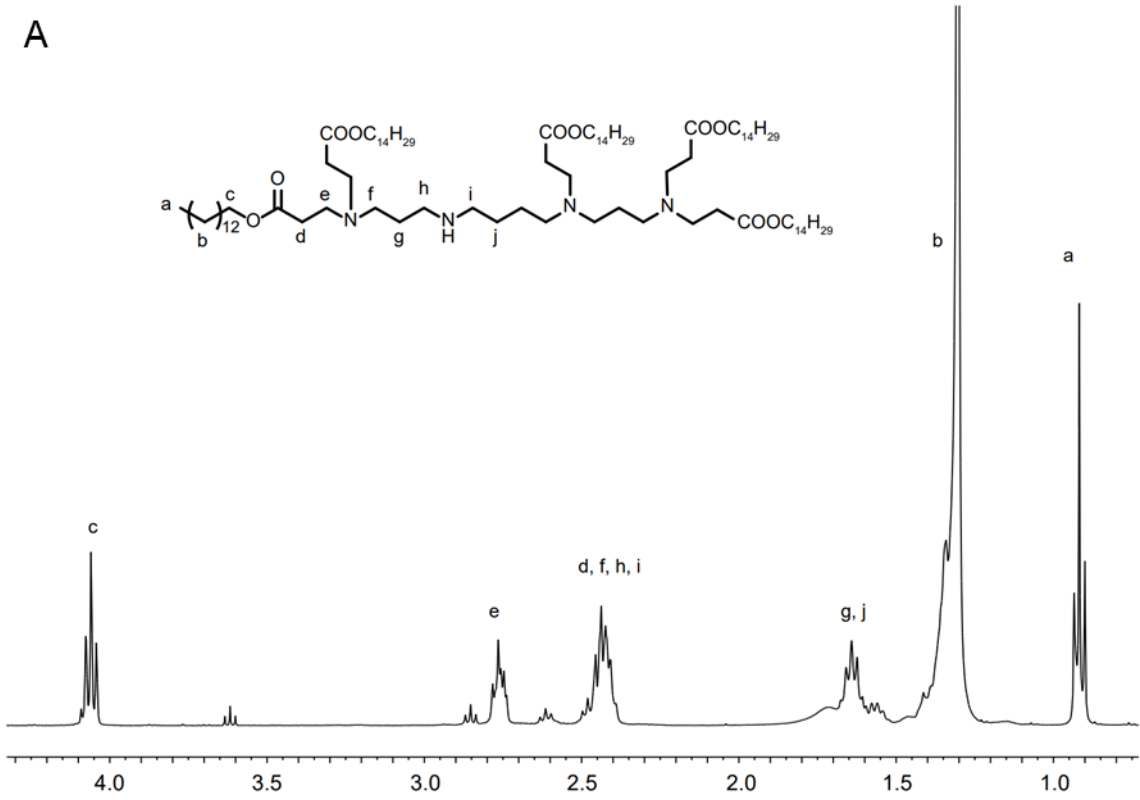
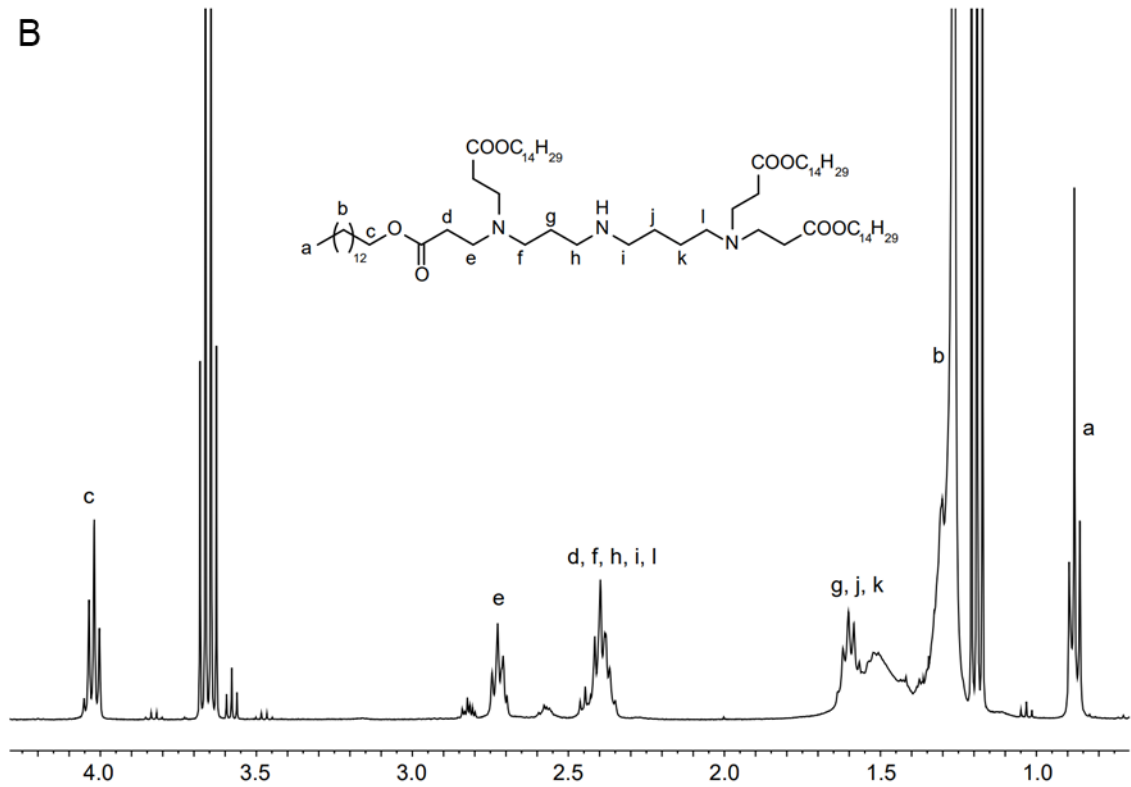
Boosting Ionizable Lipid Nanoparticle-mediated In Vivo mRNA Delivery through Optimization of Lipid Amine-Head Group

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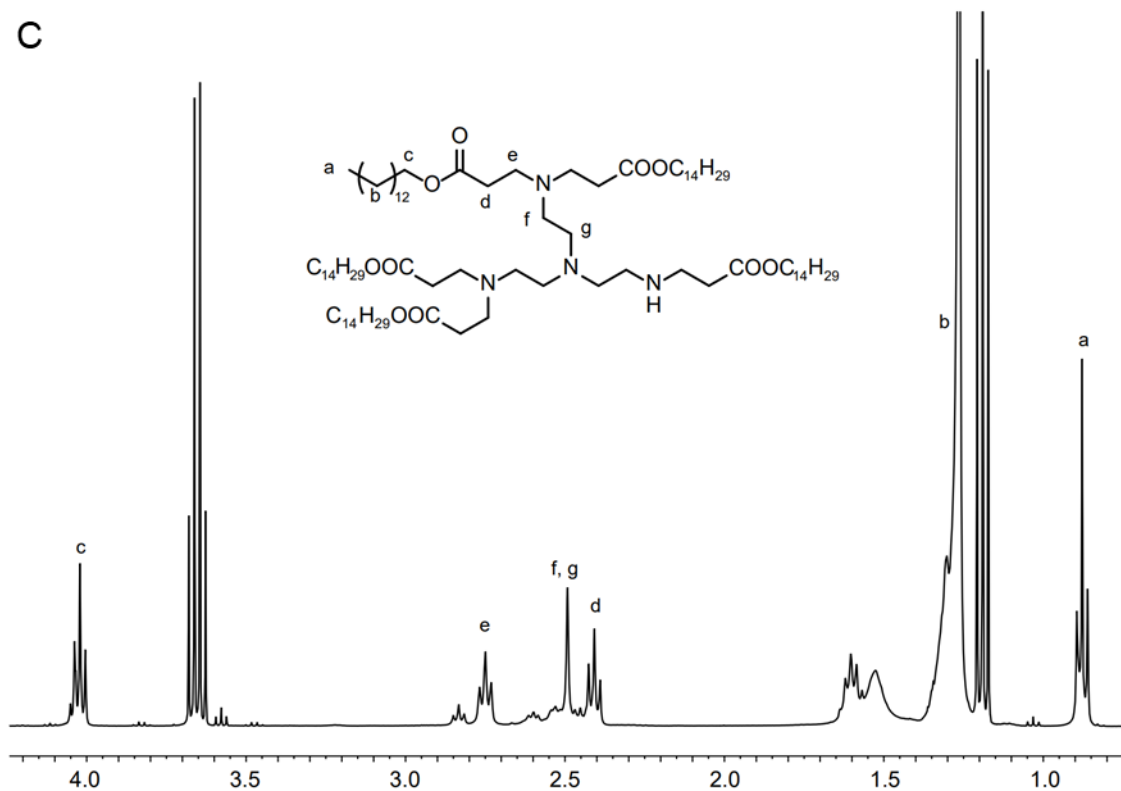
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Table S1. Primers used for RT-qPCR.

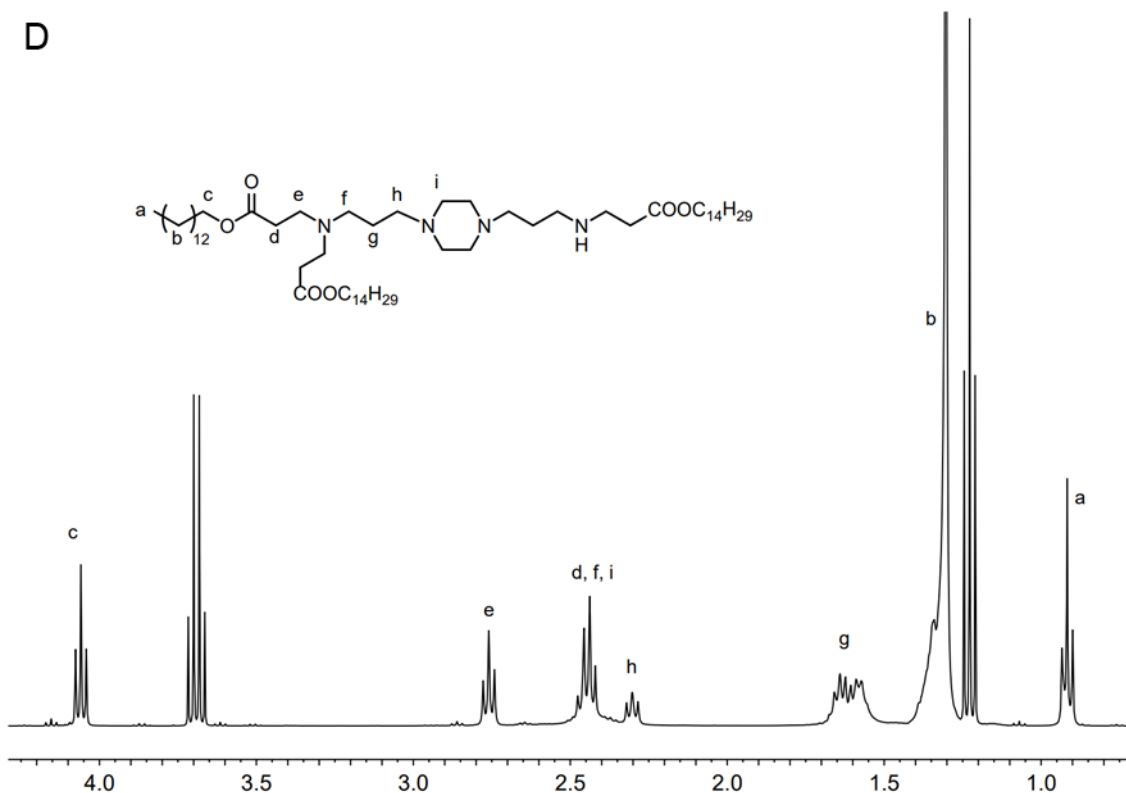
Gene	Forward (5'-3')	Reverse (5'-3')
GAPDH	GACGGCCGCATCTTCTTGTG	GCGCCCAATACGGCCAAATC
EPO	TACCTGCTGGAGGCCAA	CACGGTGATGTTCTCGTTCA

A**B**

C



D



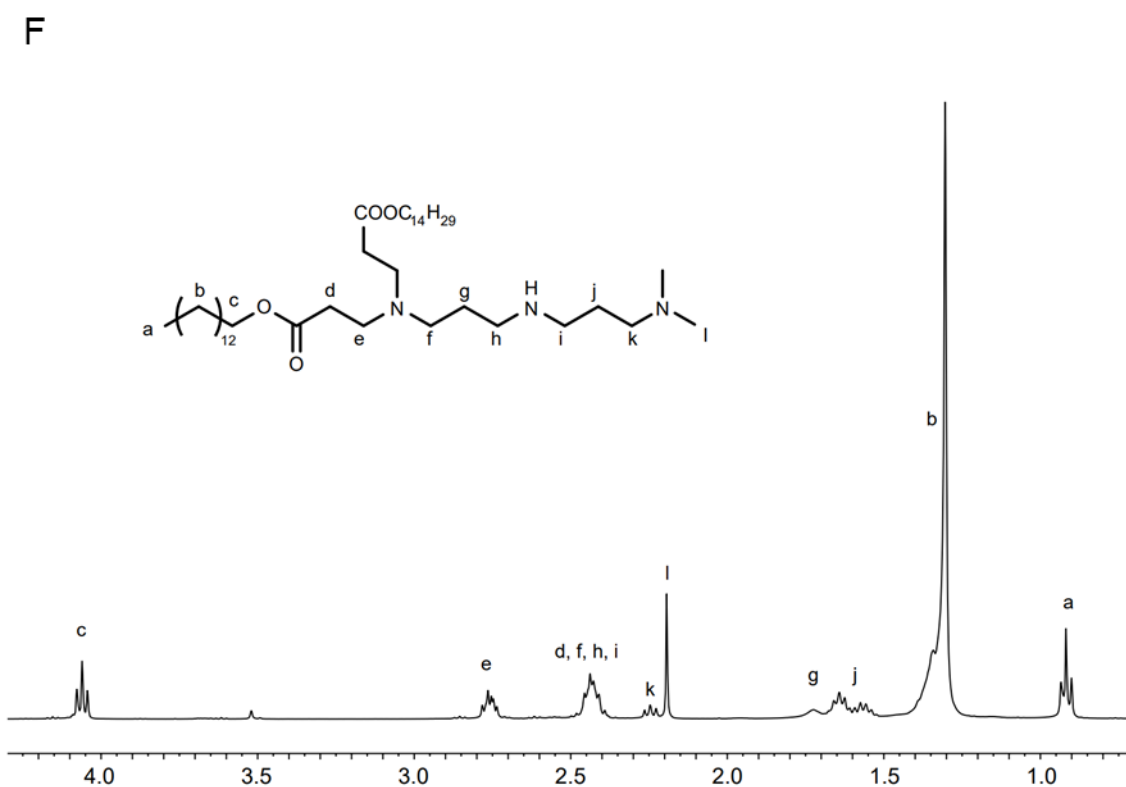
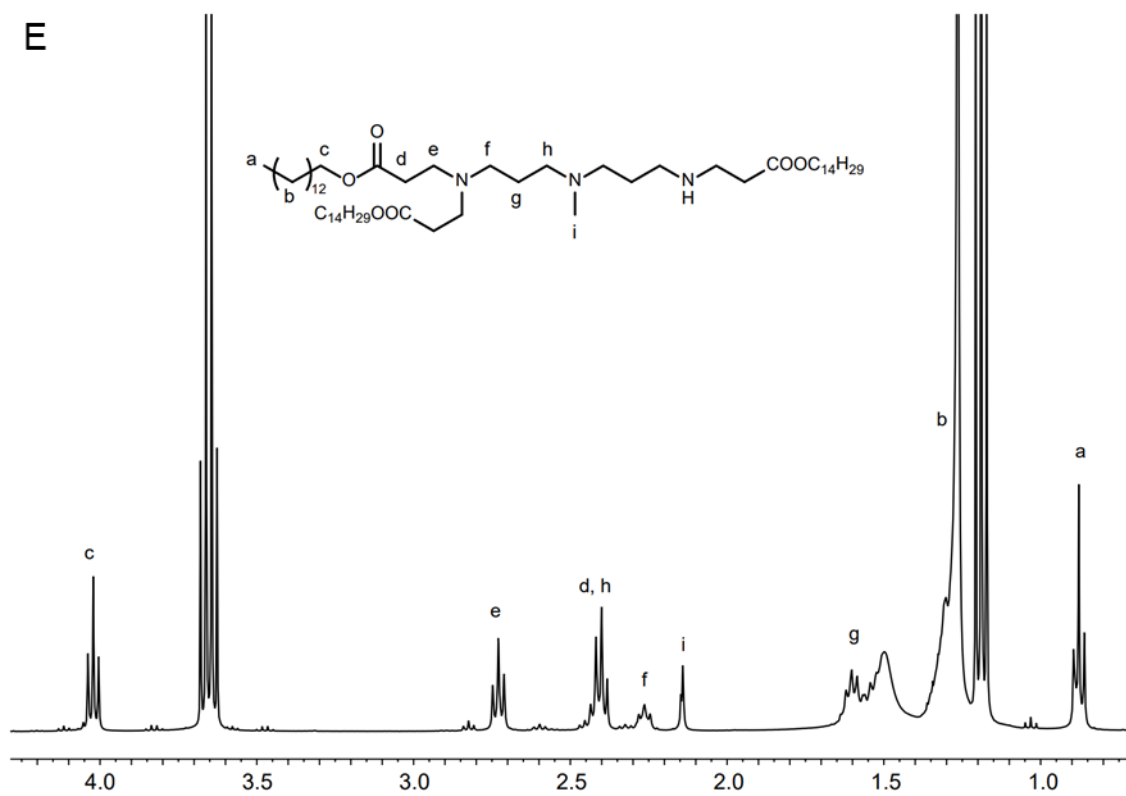


Figure S1. A representative ¹H NMR spectrum of 114 (A), 214 (B), 314 (C), 414 (D), 514 (E), 614 (F) ionizable lipids in CD₂Cl₂.

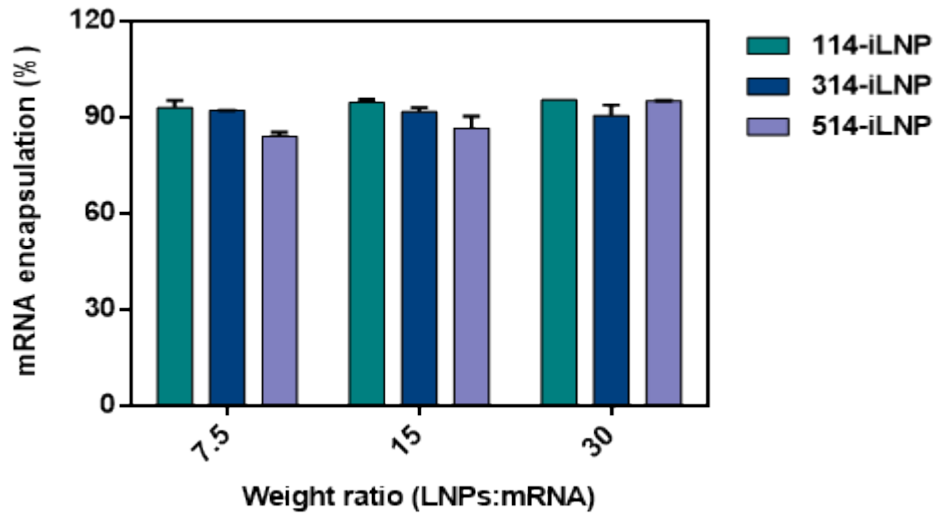


Figure S2. The mRNA encapsulation determined by RiboGreen assay at different weight ratio of iLNP:mRNA (n=2).

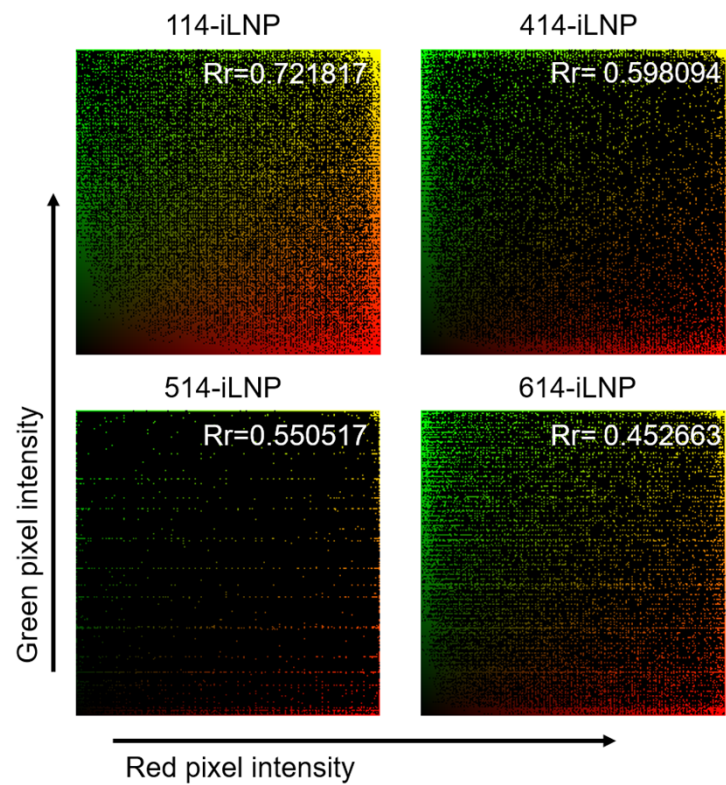


Figure S3. Colocalization analysis of endosomal escaping from iLNP/mRNA by Pearson's correlation.

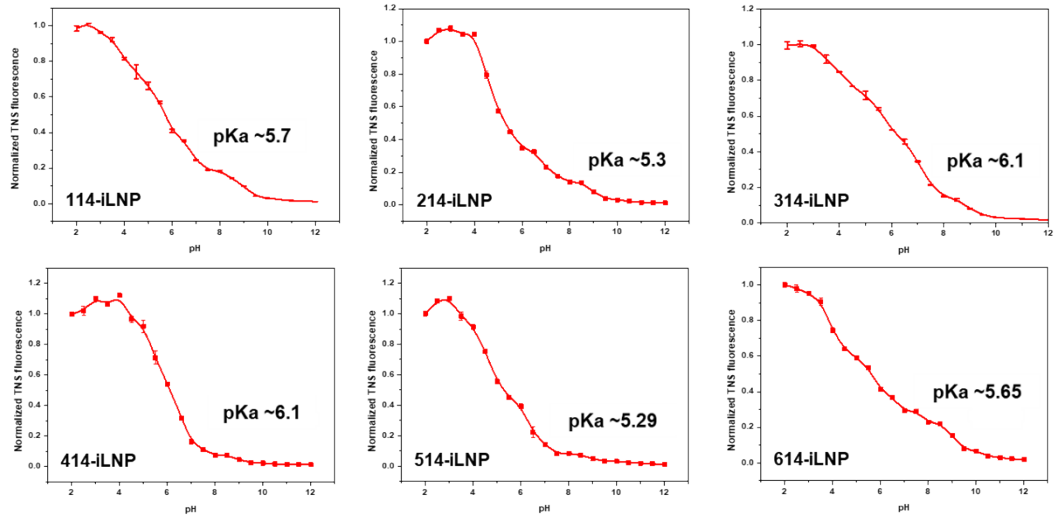


Figure S4. The pKa of iLNPs determined by TNS assay (n=2).

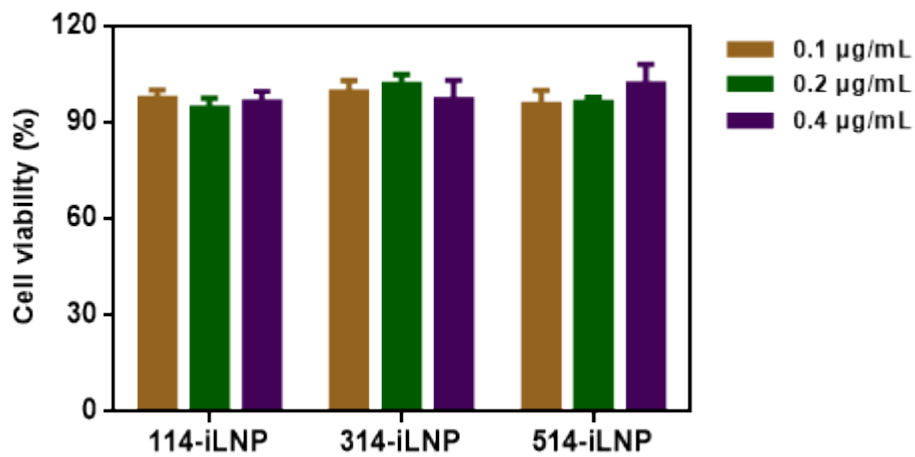


Figure S5. Cell viability of HeLa cells treated with top performance iLNP/mRNA complexes at mRNA concentration of 0.1, 0.2 or 0.4 µg/mL RNA for 24 h (n=3).

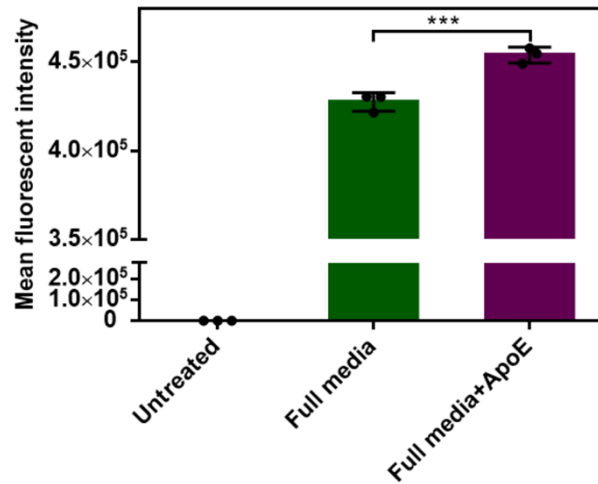


Figure S6. Quantification of cellular uptake in HeLa cells following treatment of cells with 114-iLNP in the absence or presence of Apolipoprotein (ApoE, 1 $\mu\text{g}/\text{mL}$) (n=3). Statistical analysis was performed using One-way ANOVA with Tukey test (GraphPad software Prism, Version 6.01) and statistical difference was defined as ***p < 0.001.

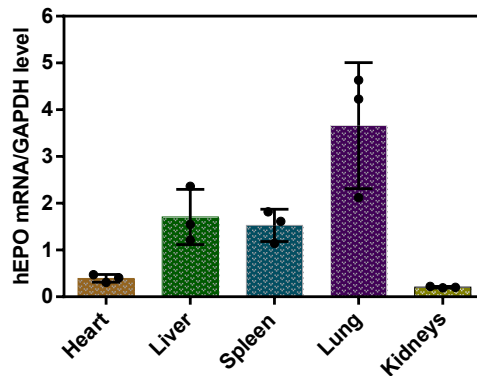


Figure S7. Quantification of hEPO mRNA in major organs by RT-qPCR including heart, liver, spleen, lung and kidneys after 6h i.v. injection of 114-iLNP/mhEPO at mRNA dose of 0.5 mg/kg (n=3). GAPDH was utilized as house-keeping gene during RT-qPCR analysis.