## Supporting Information

## Manipulating the Membrane Penetration Mechanism of Helical Polypeptides via Aromatic Modification for Efficient Gene Delivery

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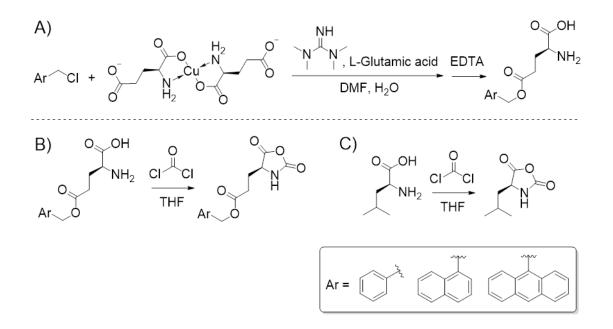
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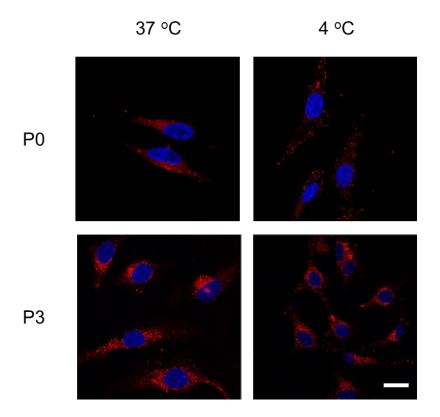


**Scheme S1.** Synthetic routes of aromatic glutamate (A), aromatic glutamate based NCA monomers (B), and the L-Leu-NCA monomer (C).

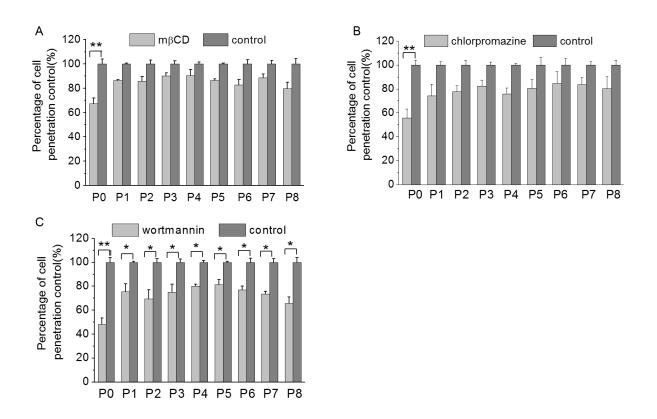
Name	$- [\theta]_{222} \times 10^{-3} (\text{cm}^2 \text{ deg dmol}^{-1})^{a)}$	Helical content (%) <sup>b)</sup>
PO	30.0	84.7
P1	27.6	78.3
P2	24.6	70.8
P3	27.2	77.4
P4	22.6	65.6
P5	31.4	88.3
P6	32.9	92.0
P7	27.5	78.3
P8	19.9	58.7

Table S1. Secondary conformational analysis of polypeptides

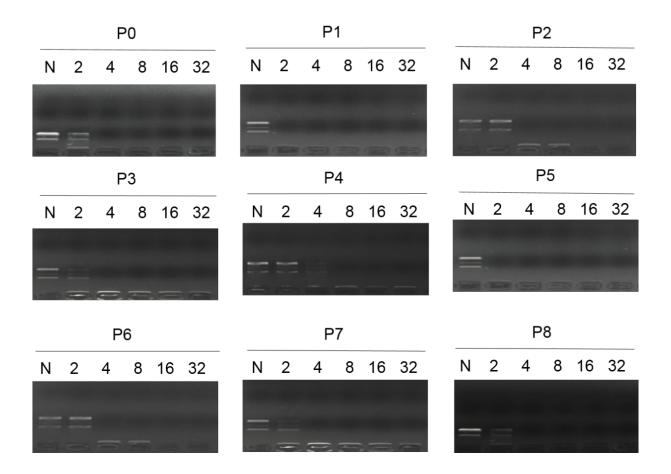
<sup>a)</sup> The mean residue ellipticity  $[\theta]$  was determined by following formula: Ellipticity ( $[\theta]$  in cm<sup>2</sup> deg dmol<sup>-1</sup>) = (millidegrees × mean residue weight) / (path length in mm × concentration of polypeptide in mg ml<sup>-1</sup>). <sup>b)</sup> The helical contents of the polypeptides were calculated by the following equation: helical content = ( $-[\theta_{222}] + 3000$ ) / 39000.



**Fig. S1.** CLSM images of HeLa cells following incubation with RhB-**P0** and RhB-**P3** at 37 °C or 4 °C for 2 h. Cell nuclei were stained with DAPI. Bar represents 20  $\mu$ m.



**Fig. S2.** Cell penetration levels of RhB-labeled polypeptides in HeLa cells in the presence of various endocytosis inhibitors including m $\beta$ CD (A), chlorpromazine (B), and wortmannin (C) (n = 3).



**Fig. S3.** DNA condensation by polypeptides at various polypeptide/DNA weight ratios as evaluated by the gel retardation assay. N represents naked DNA.

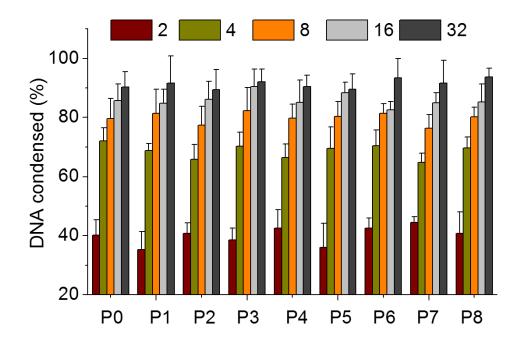


Fig. S4. DNA condensation by polypeptides at different polypeptide/DNA weight ratios as determined by the EB exclusion assay (n = 3).

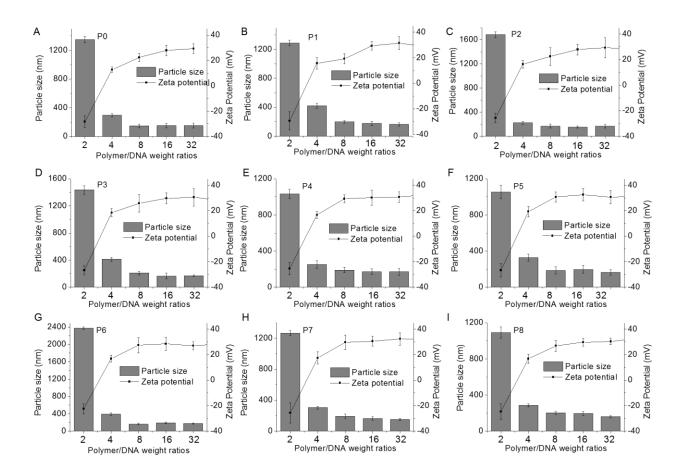


Fig. S5. Particle size and zeta potential of polypeptide/DNA complexes at different polypeptide/DNA weight ratios.

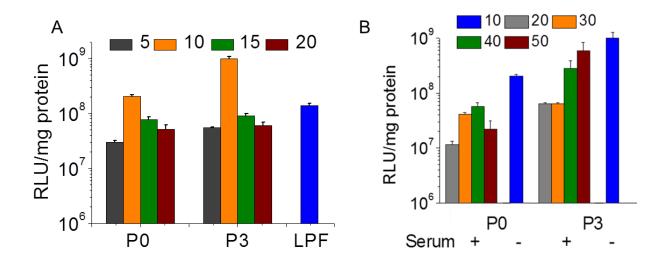
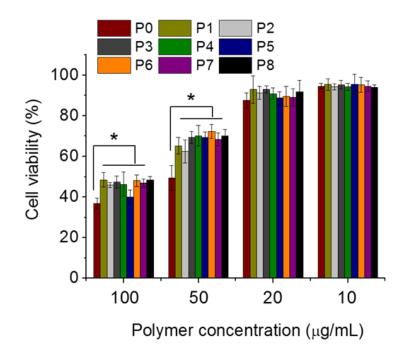
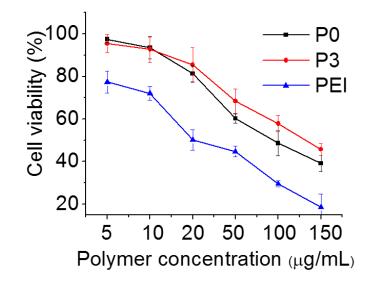


Fig. S6. Transfection efficiencies of P0 and P3 at various polypeptide/DNA weight ratios in B16F10 cells in the absence (A) or presence (B) of 10% serum (n = 3).



**Fig. S7.** Cytotoxicity of polypeptide/DNA polyplexes following 24-h incubation in HeLa cells as determined by the MTT assay (n = 3).



**Fig. S8.** Cytotoxicity of **P0**, **P3**, and PEI (25 kDa) at various concentrations in HeLa cells following 24-h incubation as determined by the MTT assay (n = 3).

<sup>1</sup>H NMR spectra of new compounds.

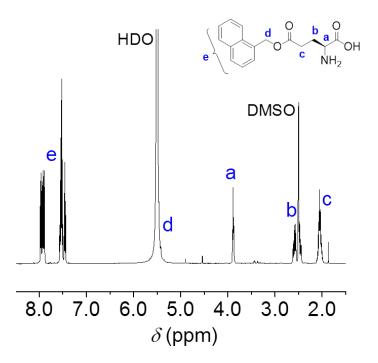


Fig. S9. <sup>1</sup>H NMR spectrum of Naph-L-Glu in DMSO-*d*<sub>6</sub>:DCl-D<sub>2</sub>O (9:1, v/v).

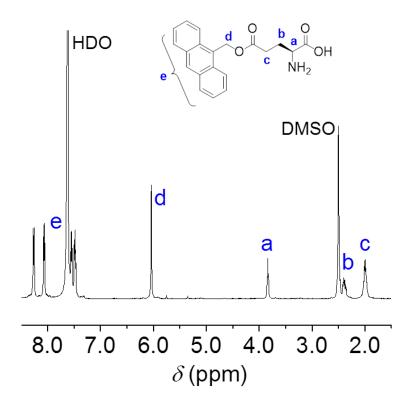


Fig. S10. <sup>1</sup>H NMR spectrum of Anth-L-Glu in DMSO-*d*<sub>6</sub>:DCl-D<sub>2</sub>O (9:1, v/v).

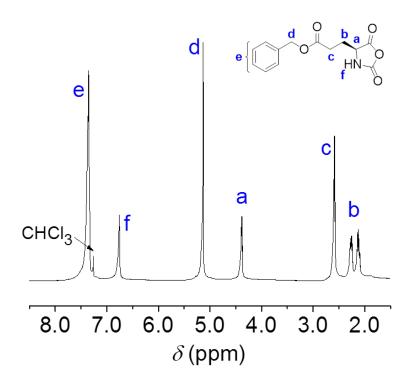


Fig. S11. <sup>1</sup>H NMR spectrum of B-L-Glu-NCA in CDCl<sub>3</sub>.

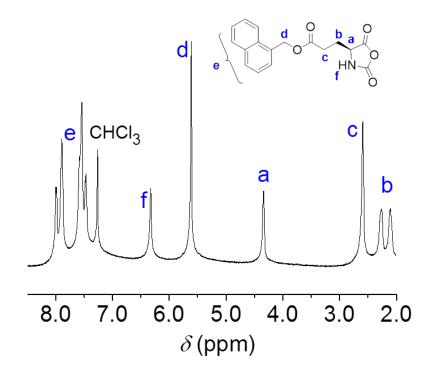


Fig. S12. <sup>1</sup>H NMR spectrum of Naph-<sub>L</sub>-Glu-NCA in CDCl<sub>3</sub>.

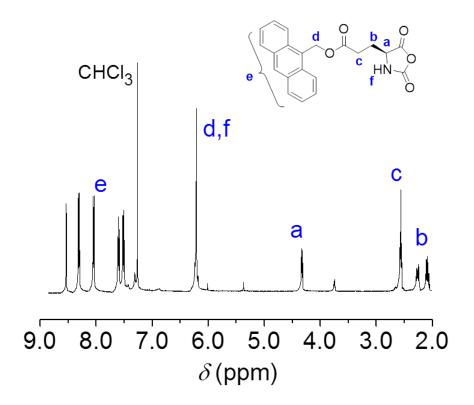


Fig. S13. <sup>1</sup>H NMR spectrum of Anth-L-Glu-NCA in CDCl<sub>3</sub>.

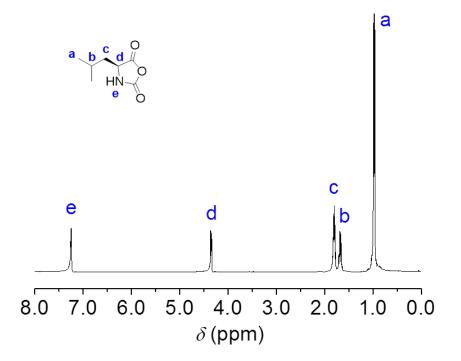
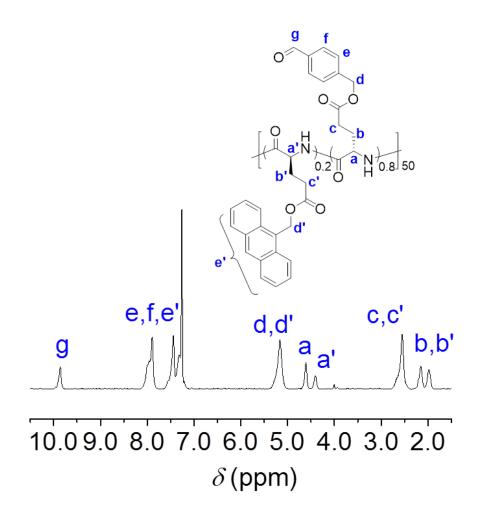


Fig. S14. <sup>1</sup>H NMR spectrum of L-Leu-NCA in CDCl<sub>3</sub>.



**Fig. S15.** Representative <sup>1</sup>H NMR spectrum of copolymer precursor for composition calculation (PALG-*r*-PABLG as an example, **P5** precursor) in CDCl<sub>3</sub>:TFA-*d* (85:15, v/v). The block composition was calculated by the integration ratio of the  $\alpha$ -protons in PALG residues (proton *a*) to the  $\alpha$ -protons in PABLG residues (proton *a*').

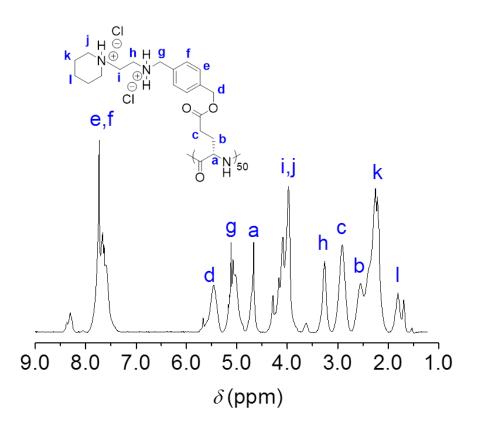


Fig. S16. <sup>1</sup>H NMR spectrum of P0 in TFA-*d*.

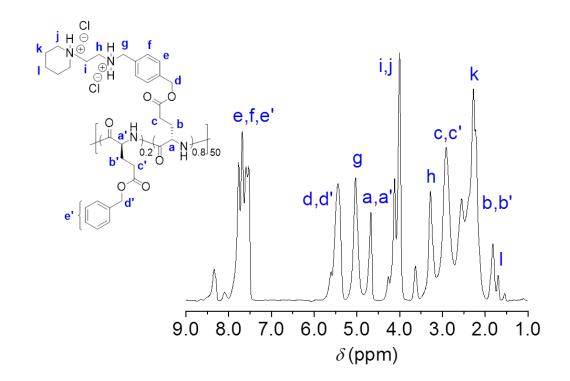


Fig. S17. <sup>1</sup>H NMR spectrum of P1 in TFA-*d*.

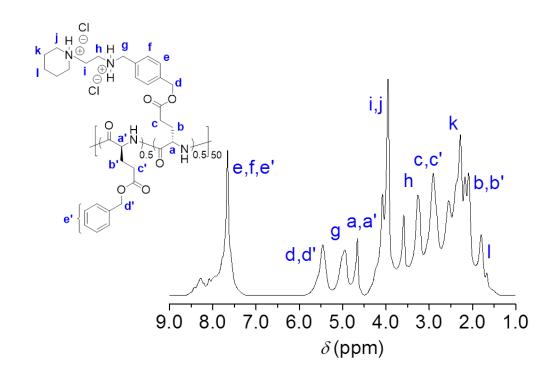


Fig. S18. <sup>1</sup>H NMR spectrum of P2 in TFA-d.

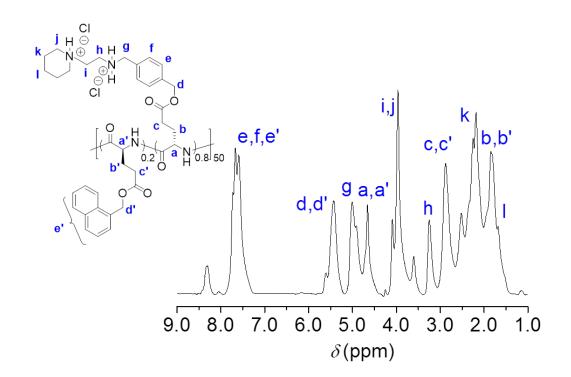


Fig. S19. <sup>1</sup>H NMR spectrum of P3 in TFA-*d*.

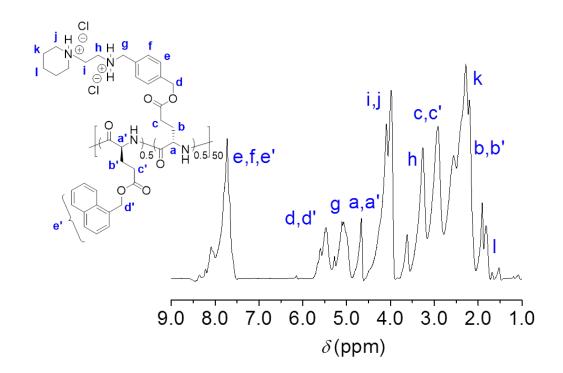


Fig. S20. <sup>1</sup>H NMR spectrum of P4 in TFA-d.

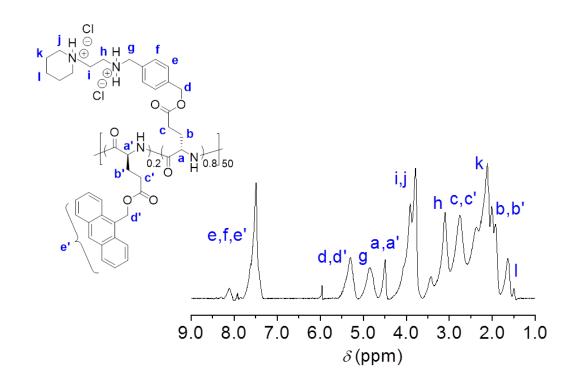


Fig. S21. <sup>1</sup>H NMR spectrum of P5 in TFA-*d*.

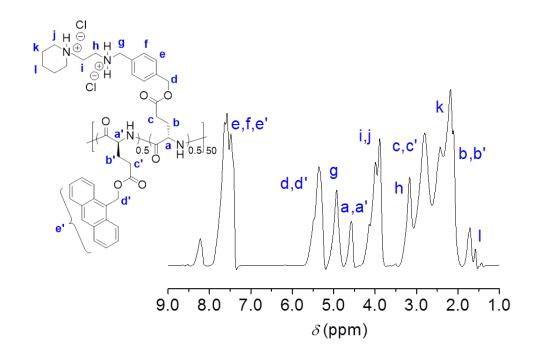


Fig. S22. <sup>1</sup>H NMR spectrum of P6 in TFA-*d*.

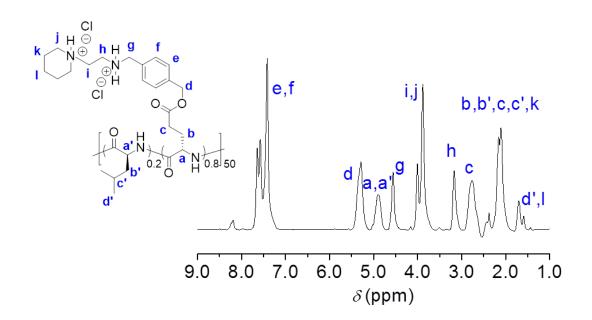


Fig. S23. <sup>1</sup>H NMR spectrum of P7 in TFA-d.

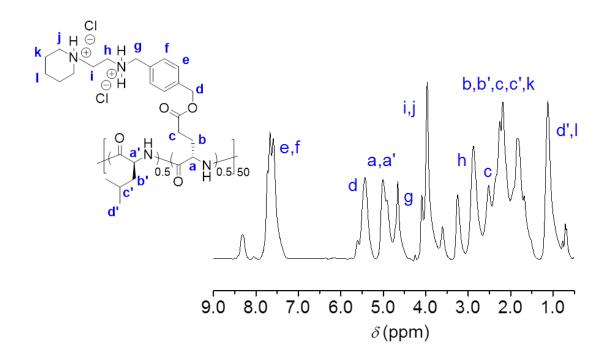


Fig. S24. <sup>1</sup>H NMR spectrum of P8 in TFA-d.