

Supplementary file 1

Saponin conjugation improves p-CRISPR delivery by polyethylenimine modified carbon dots

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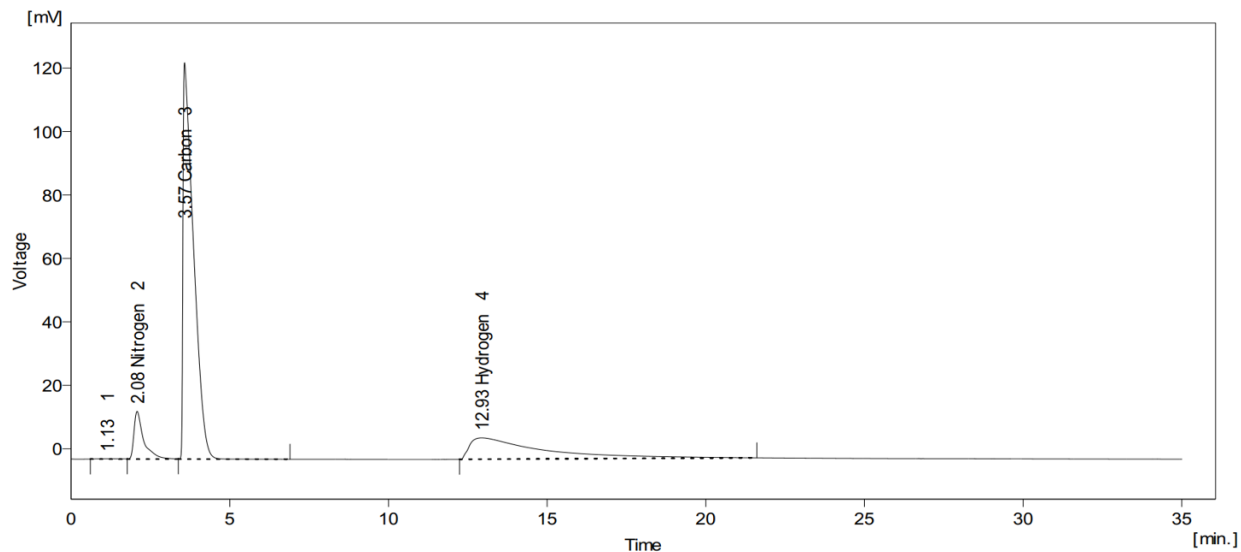


Fig S1: Elemental composition analysis of CD-PEI-Sap nanoparticles. This analysis confirmed the presence of carbon (30.86%), nitrogen (8.97%) and hydrogen (5.80%). The detectable nitrogen content originates from surface functionalization of the carbon dots with PEI 1.8kDa and saponin, and contributes to the positive surface charge of the nanoparticles, which explains the improved plasmid binding and enhanced transfection efficiency.

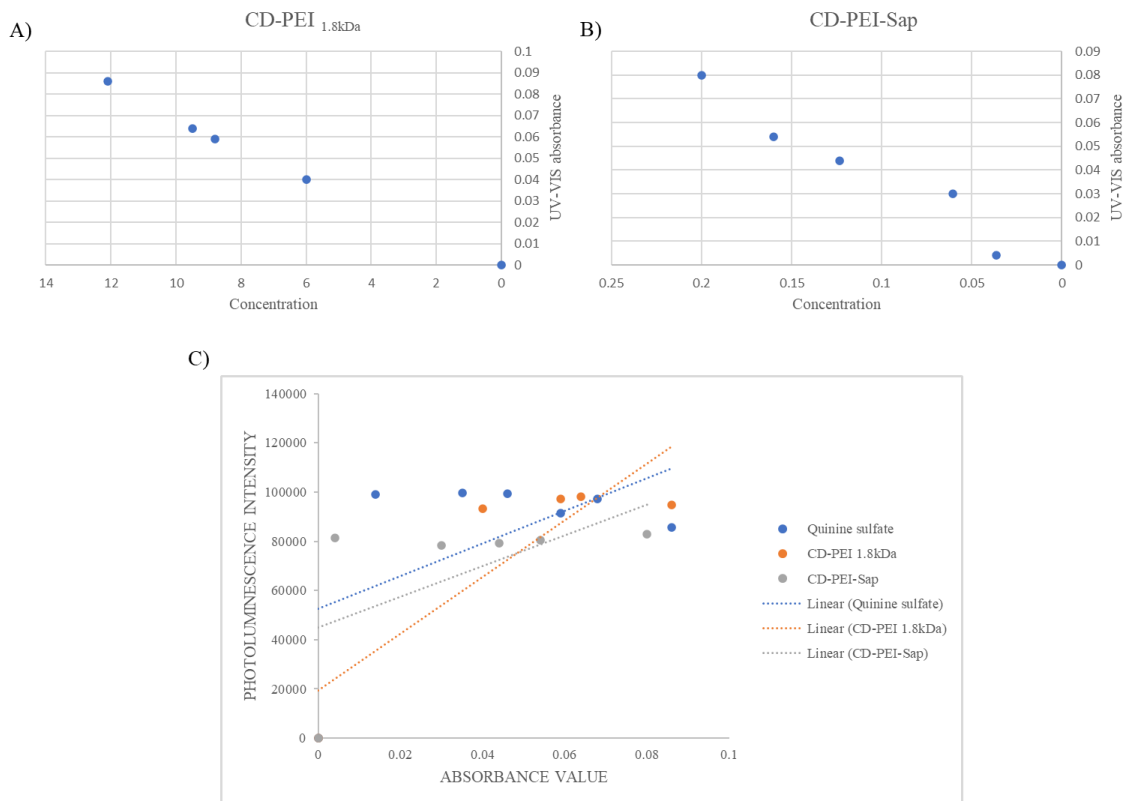
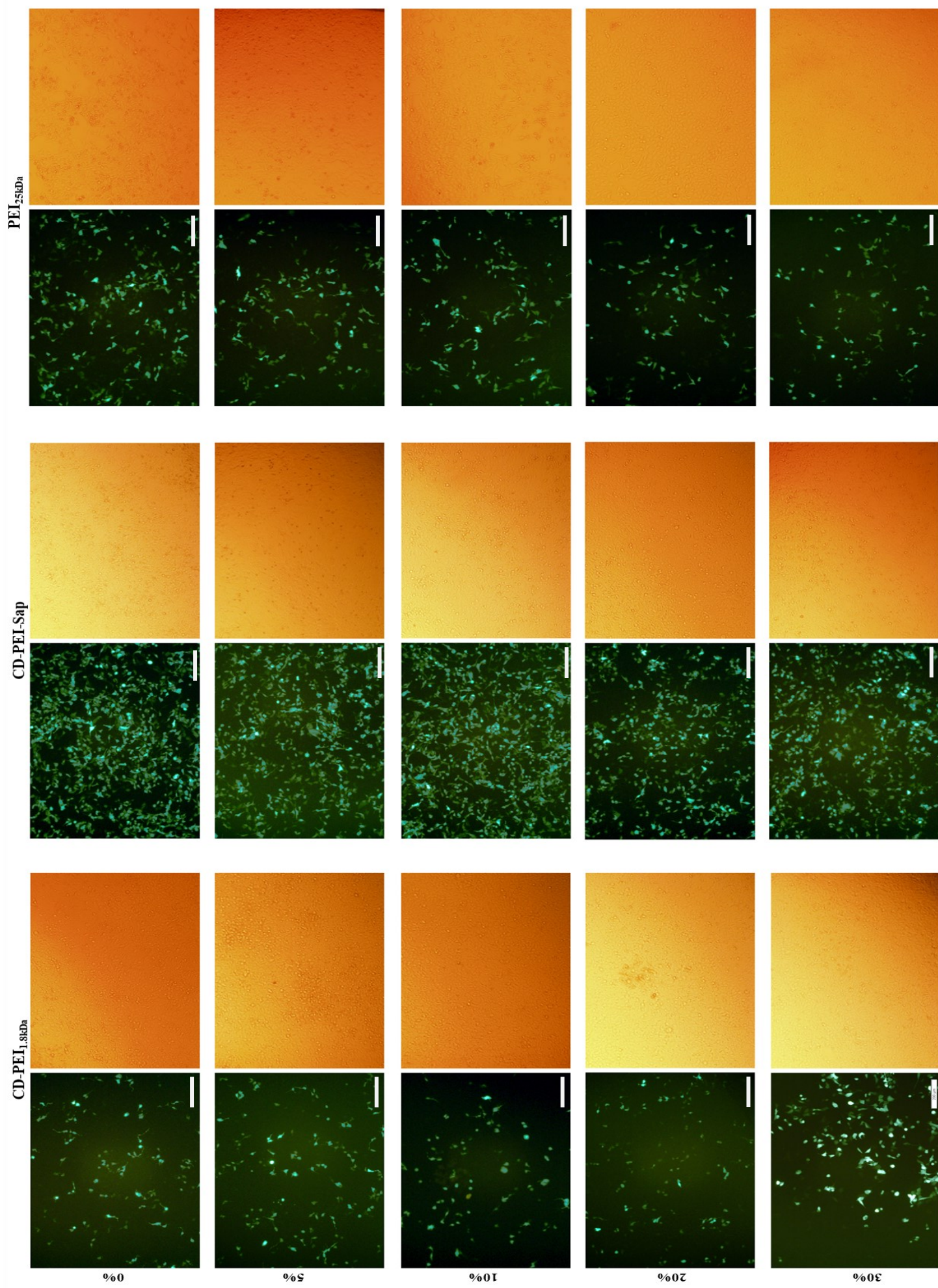
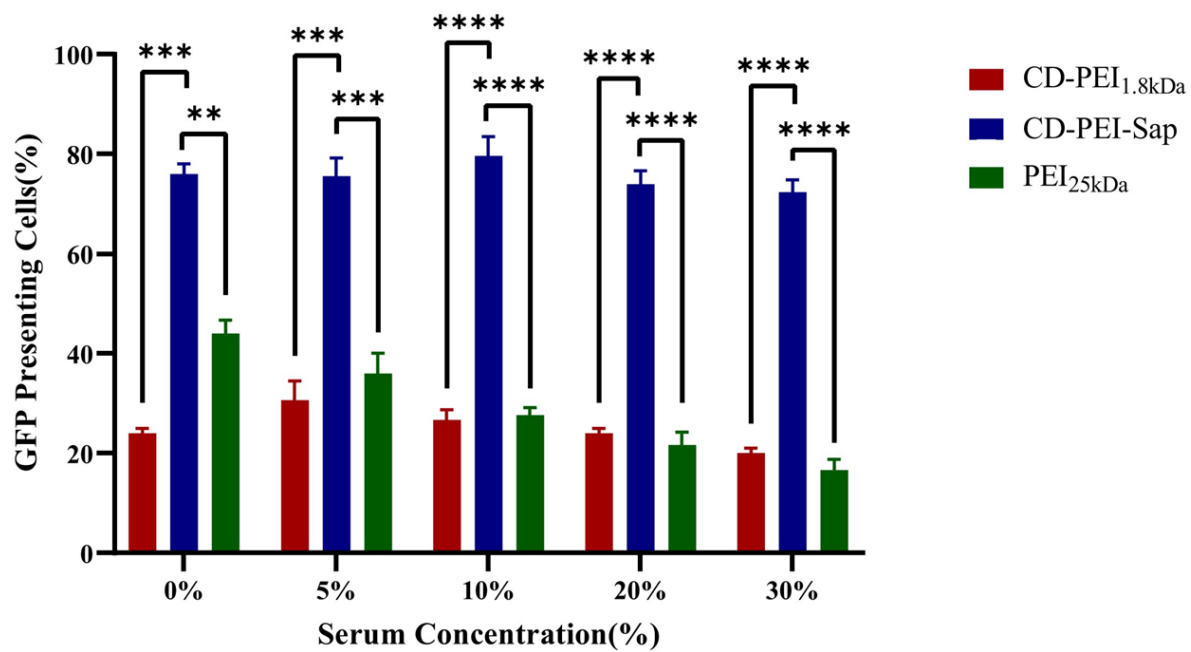


Fig S2: Optical characterization and photoluminescence spectroscopy of carbon dot-based nanoparticles A, B) UV-Visible spectra of CD-PEI_{1.8kDa} and CD-PEI-Sap is provided in various scatterplots because concentration scales were different. C) The linear graph compares the integrated photoluminescence intensity (vertical axis) and the absorbance value (horizontal axis) of CD-PEI_{1.8kDa}, CD-PEI-Sap with a reference (quinine sulfate). PL spectra were recorded over a range of emission wavelengths using a fixed excitation wavelength of 340 nm. Quinine sulfate was used as a reference standard for comparison. The CD-PEI-Sap nanoparticles exhibited slightly lower PL intensity compared to CD-PEI_{1.8 kDa}, which is attributed to fluorescence quenching effects induced by saponin incorporation in the carbon dot structure.

A)



B)



C)

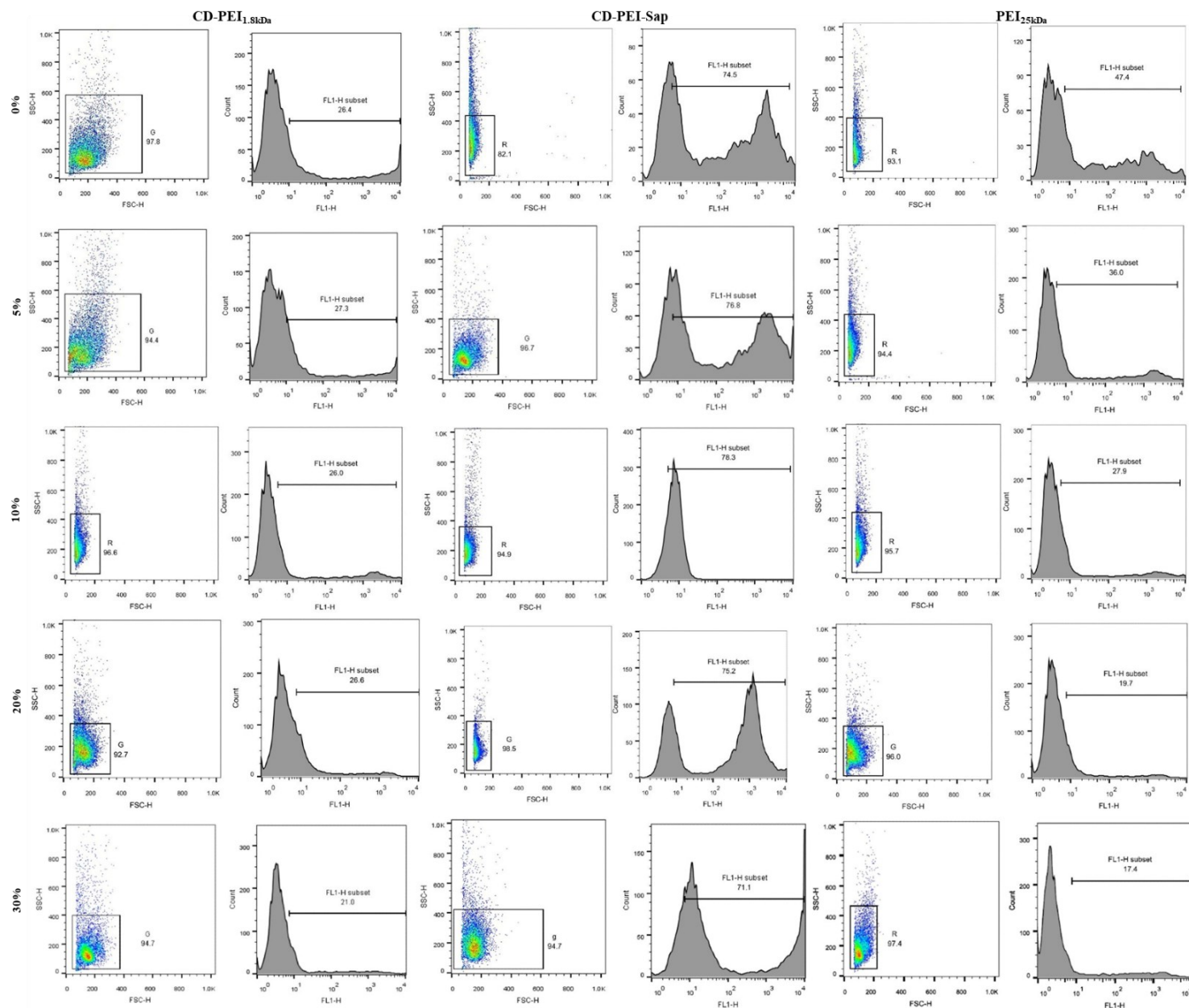
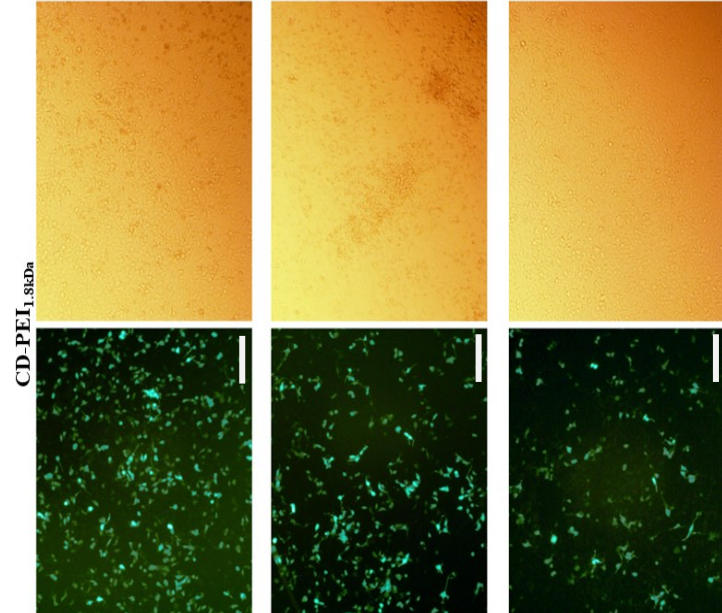
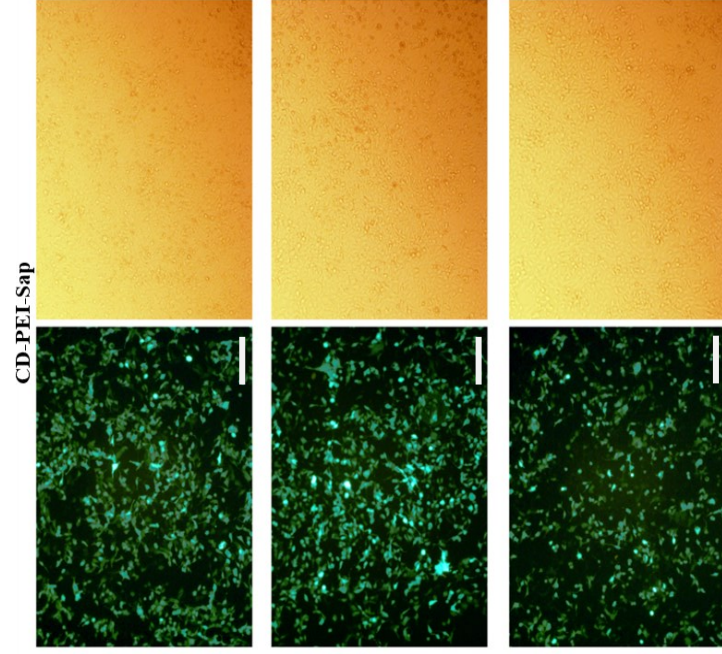
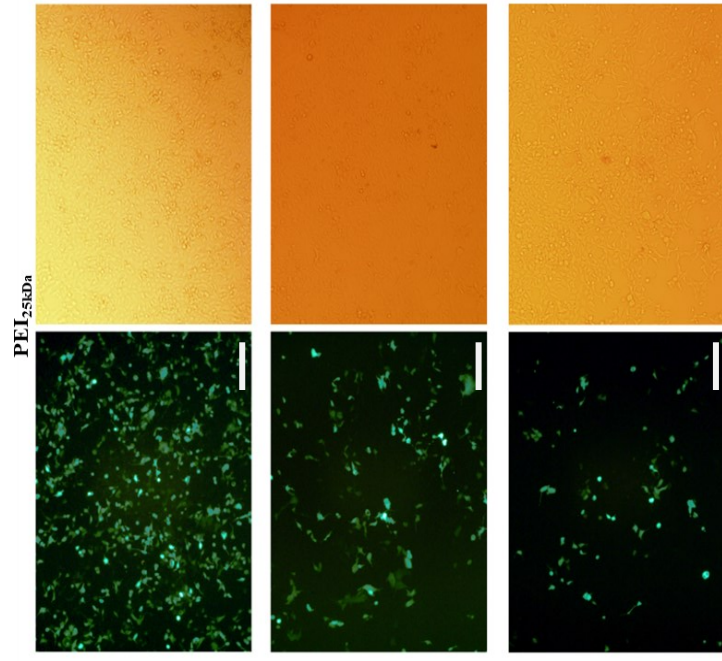


Fig S3: CD-PEI-Sap transfection yield at increasing concentrations of serum. (A) Representative fluorescence microscopy images showing the expression of GFP in HEK 293T cells transfected with CD-PEI_{1.8kDa}, CD-PEI-Sap, PEI_{25kDa} at their optimal weight ratios in various concentrations of FBS (0%, 10%, 20% and 30%). (B) Quantitative analysis of transfection efficiency provided as the percentage of GFP-positive cells following transfection with vector/p-CRISPR (3 kb) complexes under identical serum conditions. Transfections were performed using 1 μ g of plasmid DNA and evaluated 48 h post-transfection. (Scale bar = 100 μ m) CD-PEI-Sap maintained high transfection efficiency even at elevated serum concentrations, indicating boosted serum stability compared to PEI_{25kDa}, whose performance decreased with rising serum levels. C) Average flow cytometry data for different serum concentrations.



B)

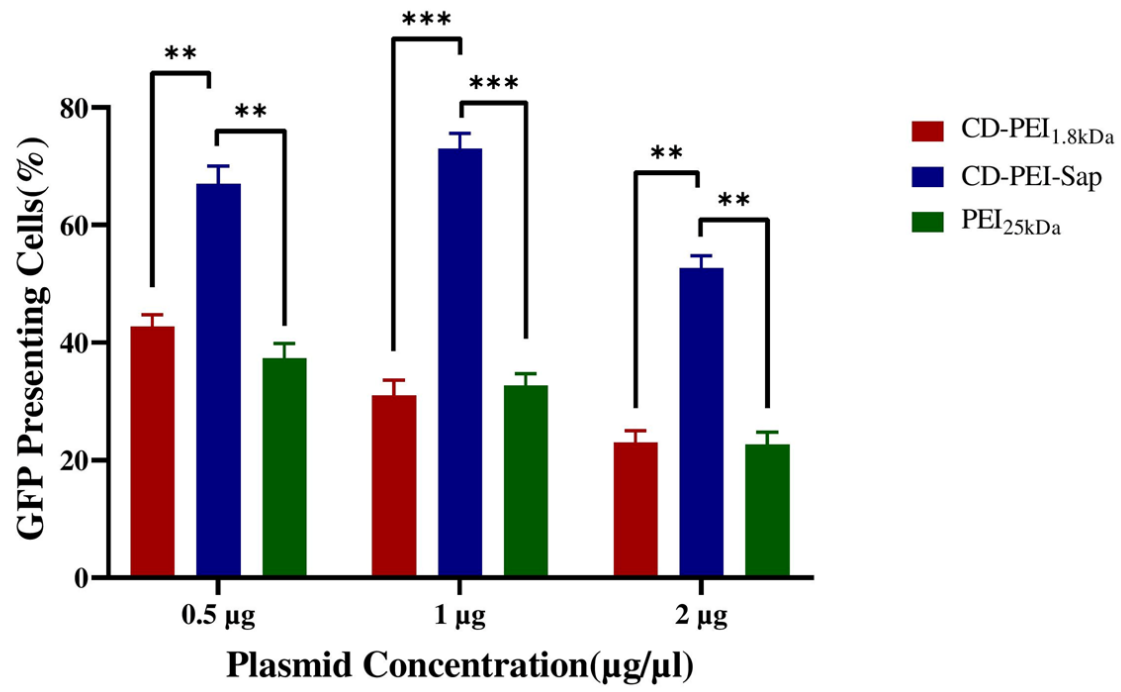
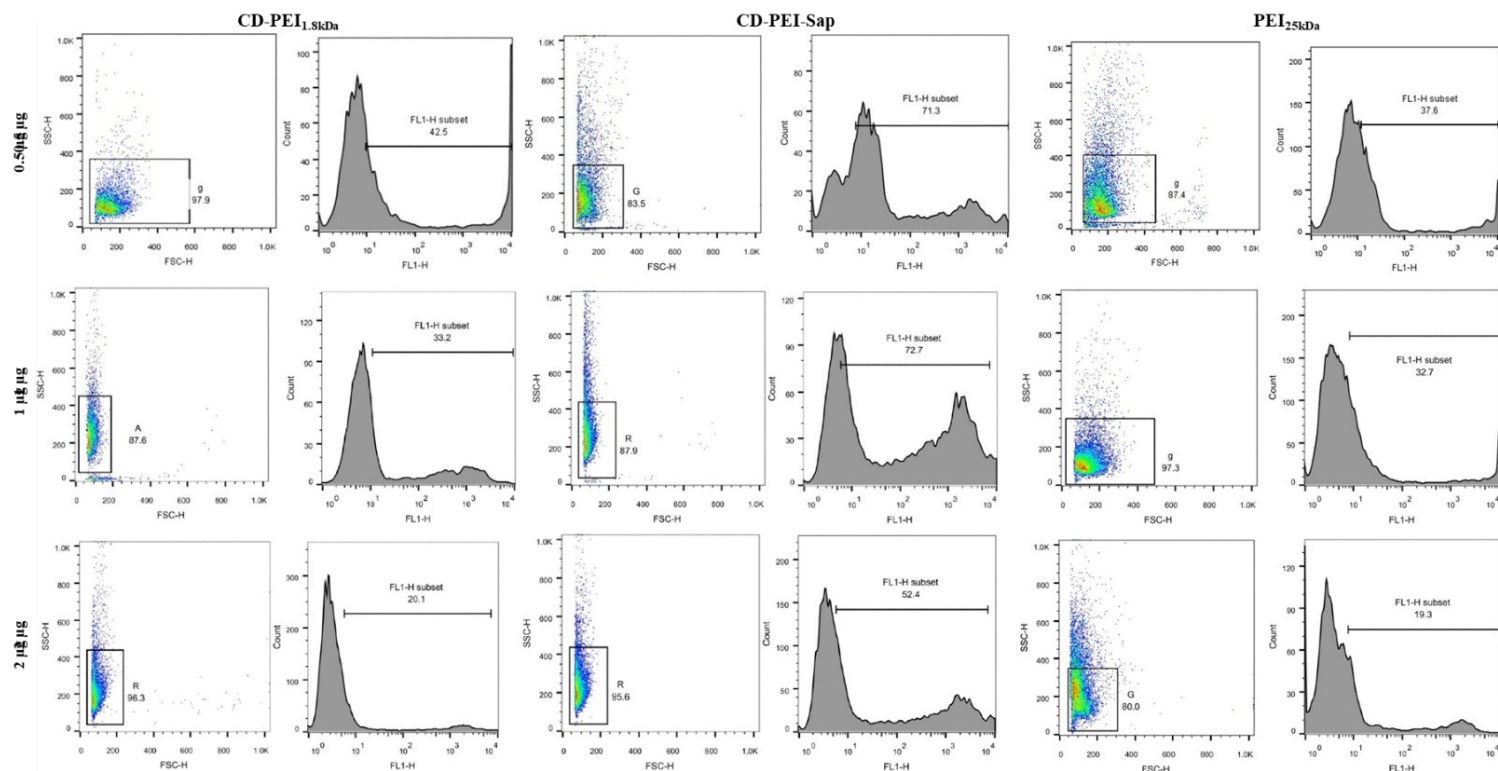
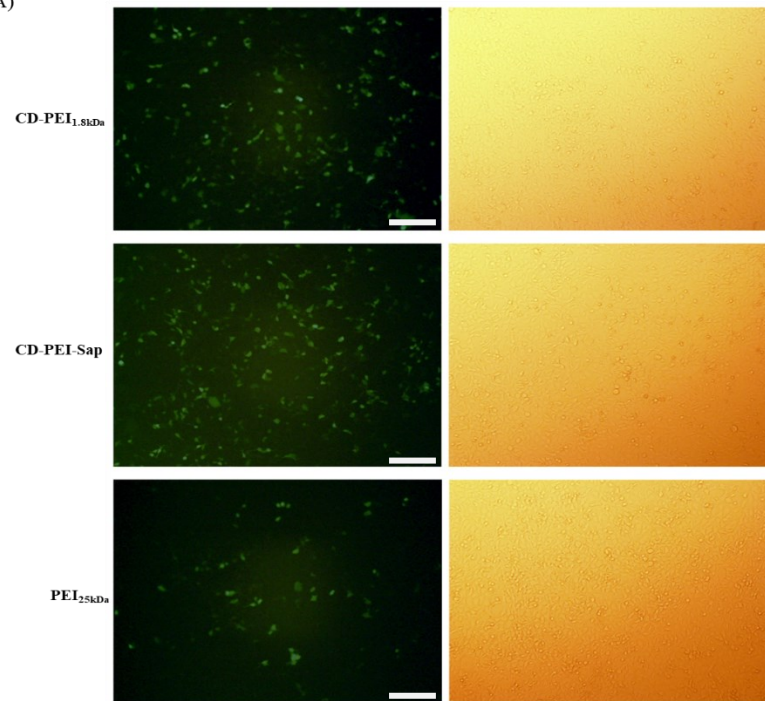


Fig S4: The transfection efficiency of CD-PEI-Sap was evaluated at low doses of p-CRISPR (3kb). (A) The transfection efficiency of CD-PEI_{1.8kDa}, CD-PEI-Sap and PEI_{25kDa} at the optimized weight ratio was

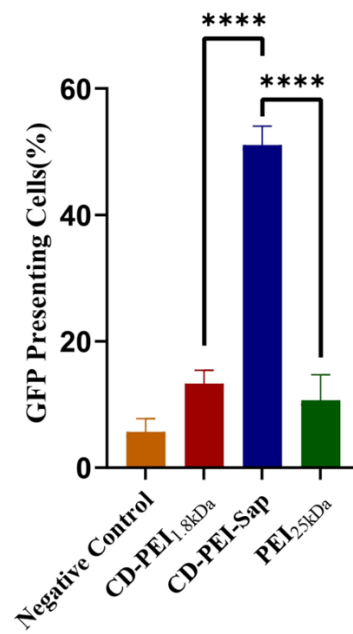


evaluated using fluorescence microscopy images. (B) Quantitative analysis of transfection efficiency expressed as the percentage of GFP-positive cells following transfection with carrier/p-CRISPR (3 kb) complexes at diverse plasmid doses (0.5, 1, and 2 µg). Transfections were evaluated 48 h post-incubation. (Scale bar = 100 µm) CD-PEI-Sap revealed high transfection efficiency even at the lowest plasmid dose (0.5 µg), outperforming PEI_{25kDa} and demonstrating efficient gene delivery at reduced DNA concentrations. C) Average flow cytometry information for low doses p-CRISPR.

A)



B)



C)

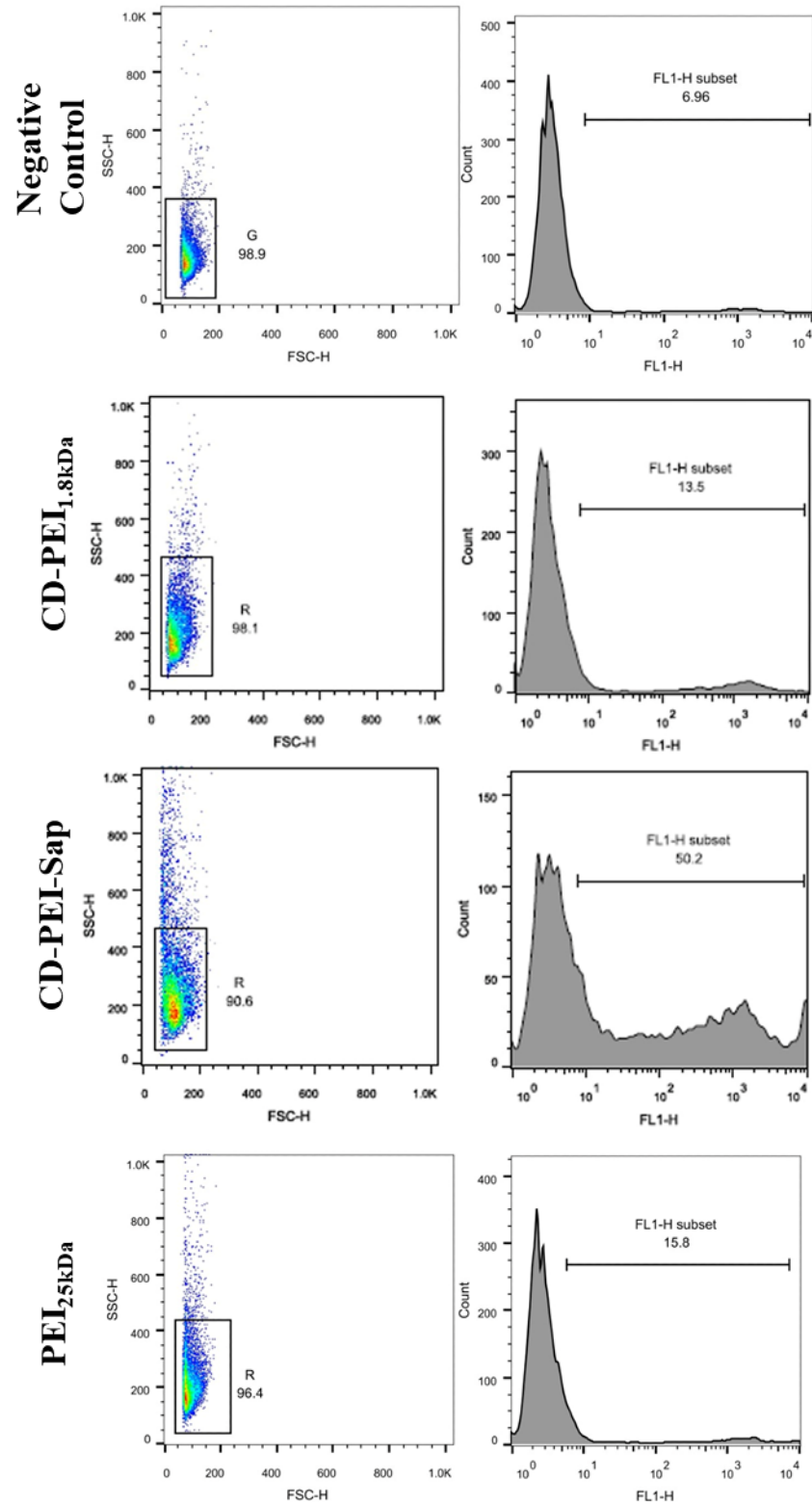


Fig S5: Delivery of large p-CRISPR plasmid (9 kb) by CD-PEI-Sap nanocarriers in HEK 293T cells. A) Representative fluorescence microscopy images showing GFP expression after transfection of p-CRISPR (9 kb) into HEK 293T cells 48 h after transfection with a large p-CRISPR plasmid (9 kb) delivered by CD-PEI-Sap nanoparticles. (Scale bar = 100 μ m) B) Quantitative assessment of transfection efficiency determined by flow cytometry following delivery of the p-CRISPR (9 kb) plasmid using CD-PEI-Sap at its adjusted carrier-to-plasmid weight ratio (80:1). Scale bar = 100 μ m. Despite the large plasmid size, CD-PEI-Sap achieved notable gene delivery, demonstrating its capability to transport high-molecular-weight genetic cargos. C) Average flow cytometry data of p-CRISPR plasmid (9kb) in HEK 293T cells.