Supporting Information

RNA Delivery Using a Graphene Oxide-Polyethyleneimine Hybrid Inhibiting Myotube Differentiation

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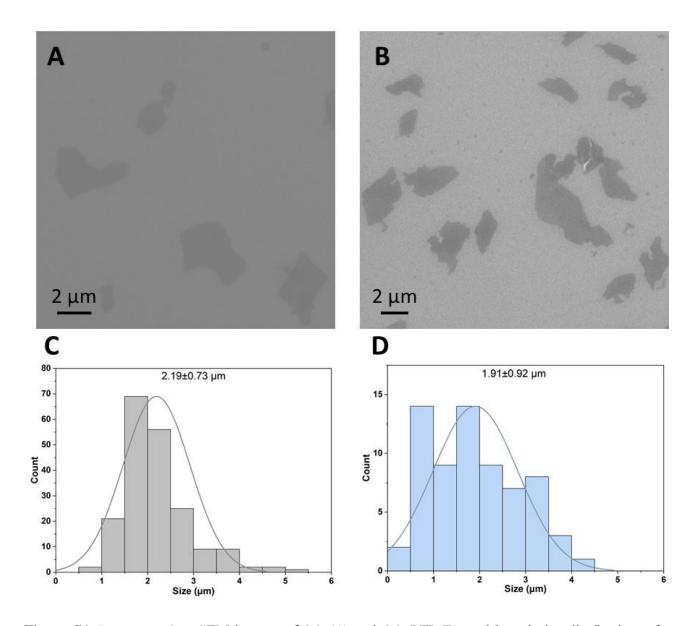


Figure S1. Representative SEM images of GO (A) and GO-PEI (B), and lateral size distribution of GO (C) and GO-PEI (D), measuring 67 and 198 sheets respectively.

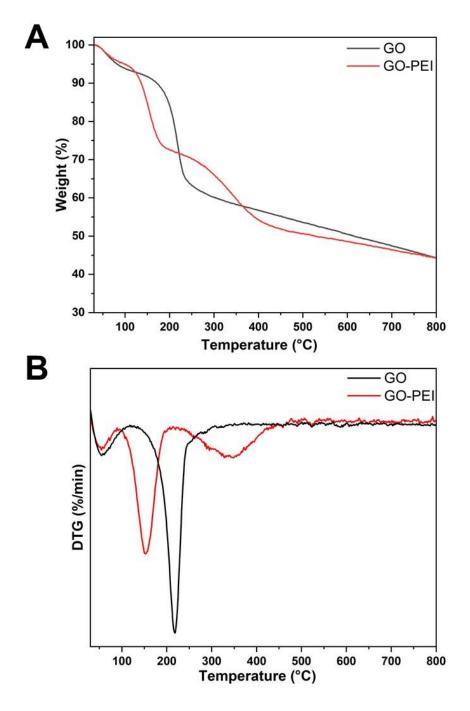


Figure S2. TGA (A) and DTG (B) curves of GO and GO-PEI.

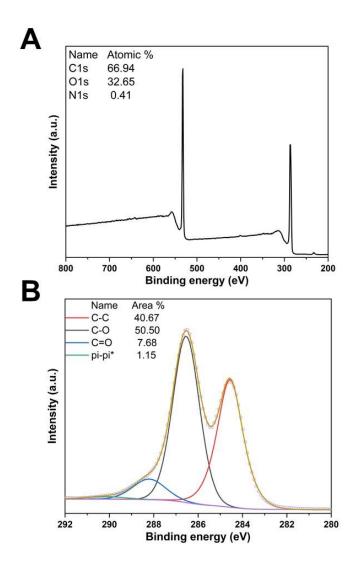


Figure S3. XPS survey spectrum of GO (A), and high-resolution C1s spectrum of GO (B).

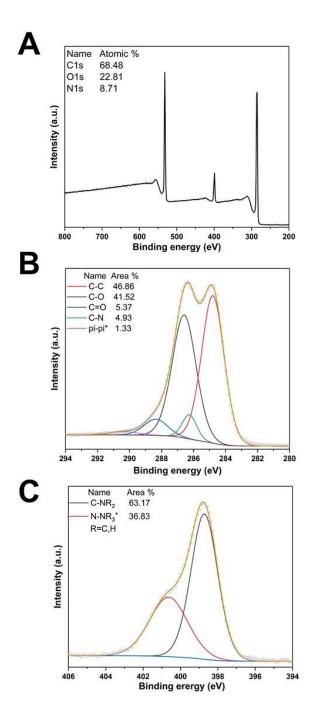


Figure S4. XPS survey spectrum of GO-PEI (**A**), high-resolution C 1s spectrum of GO-PEI (**B**), and high-resolution N 1s spectrum of GO-PEI (**C**).

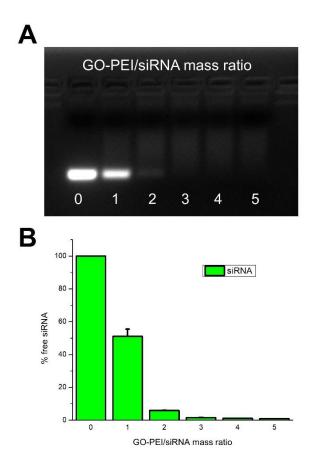


Figure S5. Complexation of GO-PEI with siRNA. (**A**) Image of agarose gel electrophoresis of GO-PEI/siRNA. (**B**), Histograms showing the free siRNA signal intensity at different GO-PEI/siRNA mass ratios.

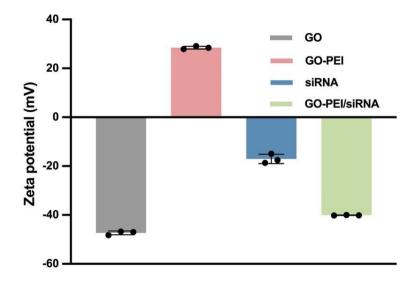


Figure S6. Zeta potential of GO, GO-PEI, siRNA and GO-PEI/siRNA (1:1).

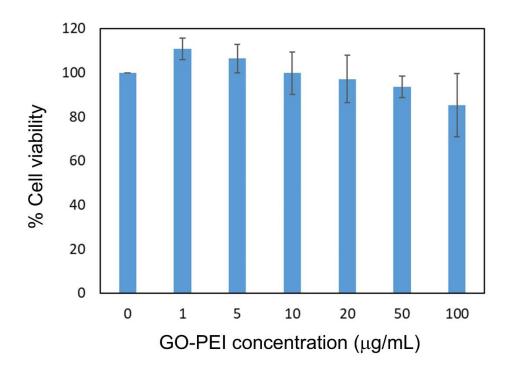


Figure S7. C2C12 cell viability analysis using AlamarBlue assay at different GO-PEI concentration. Error bars are standard deviation (S.D.) of independent experiments (N=3 \sim 5). Significant difference with the averages was analyzed by one-way ANOVA (*P<0.05). The Dunnett method compares each group with the control group (0 μ g/mL). No significant difference between all other groups with the control was found.

Table \$1. dsRNA sequences used on the siRNA experiments.

dsRNA code	RNA sequences (sense strand only)	CDS position						
RNAi positive								
1	AGGUGUGUAAGAGGAAGUC	264-282						
2	ACCAUGCCCAACUGAGAUU	713-731						
	Scrambled negative control							
S1	GUUAAGGCGGUGAAUGAGA							
S2	GAAUCAGUUCCGAUCCACA							

Table S2. Primer sequences for the qPCR experiments.

Myogenin				
Forward	CATCCAGTACATTGAGCGCCTA			
Reverse	GAGCAAATGATCTCCTGGGTTG			
Beta-actin (house keeper-gene)				
Forward	TTGCTGACAGGATGCAGAAG			
Reverse	se GTACTTGCGCTCAGGAGGAG			

Table S3. Comparison of our and previous GO-PEI hybrids as RNA transfection materials.

Type of RNA	Type of GO functionalization	Average size (nm)	ζ-potential (mV)	Cell types or animals	Purpose of the transfection	Ref.*
mRNA	Non-covalent interaction	205	+26.7	Adipose tissue-derived fibroblasts	Induction of pluripotent stem cells	1 (22)
mRNA	Non-covalent interaction	220	-	DC2.4, RAW264.7, B16-OVA cells/mice	Release of ovalbumin encoding mRNA	2 (23)
siRNA	Amide bond formation	~200	+55.5	HeLa cells	Anticancer drug transfection	3 (24)
siRNA	Amide bond formation	~200	+27.4	Human blast carcinoma	Suppression of breast cancer	4 (25)
miRNA	Amide bond formation	174	-	HuCCT1/mice	Inhibition of intrahepatic cholangial carcinoma	5 (26)
siRNA	Epoxide ring opening reaction	1910	+28.5	C2C12	Myogenic silencing	This work

^{*}The numbers in parentheses indicate the number of the reference in the main manuscript.

References

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