

Graphene oxide size and oxidation degree govern its supramolecular interactions with siRNA

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Electronic Supplementary Information

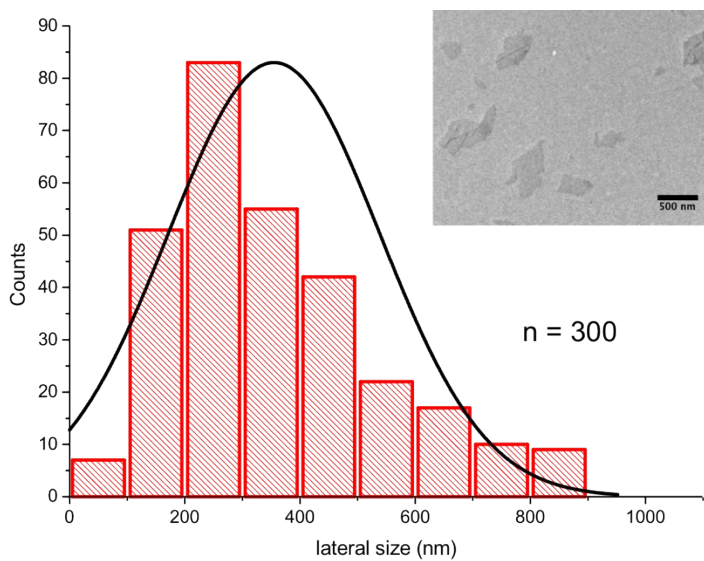


Figure S1. Lateral size distribution calculated by TEM, on the inset TEM image of GO_s.

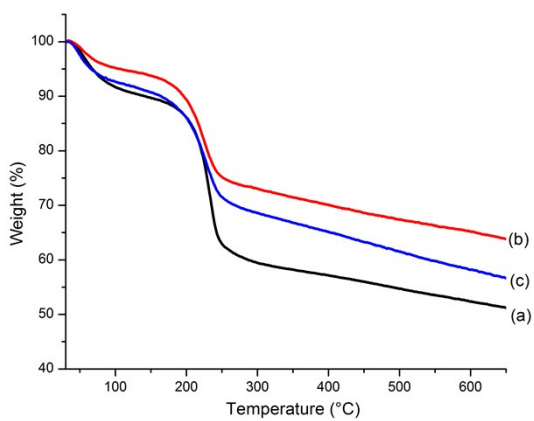


Figure S2. TGA spectra of the different graphene derivatives of GO_s. GO_s in black (a), rGO_s in red (b), and rGO_s-O₃ in blue (c).

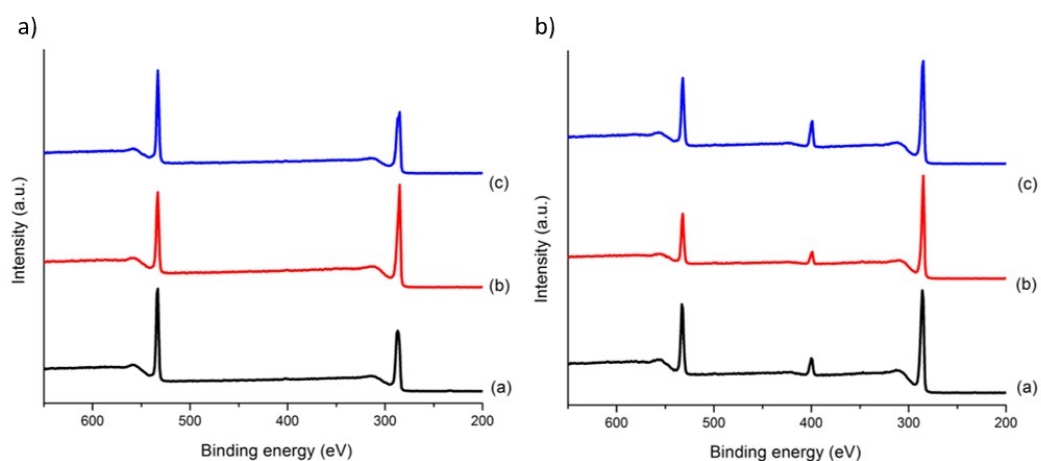


Figure S3. XPS spectra of different graphene derivatives of GO₅ and their corresponding PEI functionalization: a) GO₅ in black (a), rGO₅ in red (b), and rGO₅-O₃ in blue (c); b) GO₅-PEI in black (a), rGO₅-PEI in red (b), and rGO₅-O₃-PEI in blue (c) The C(1s) photoelectron binding energy was set at 284.5 ± 0.2 eV and used as reference for calibrating the other peak positions.

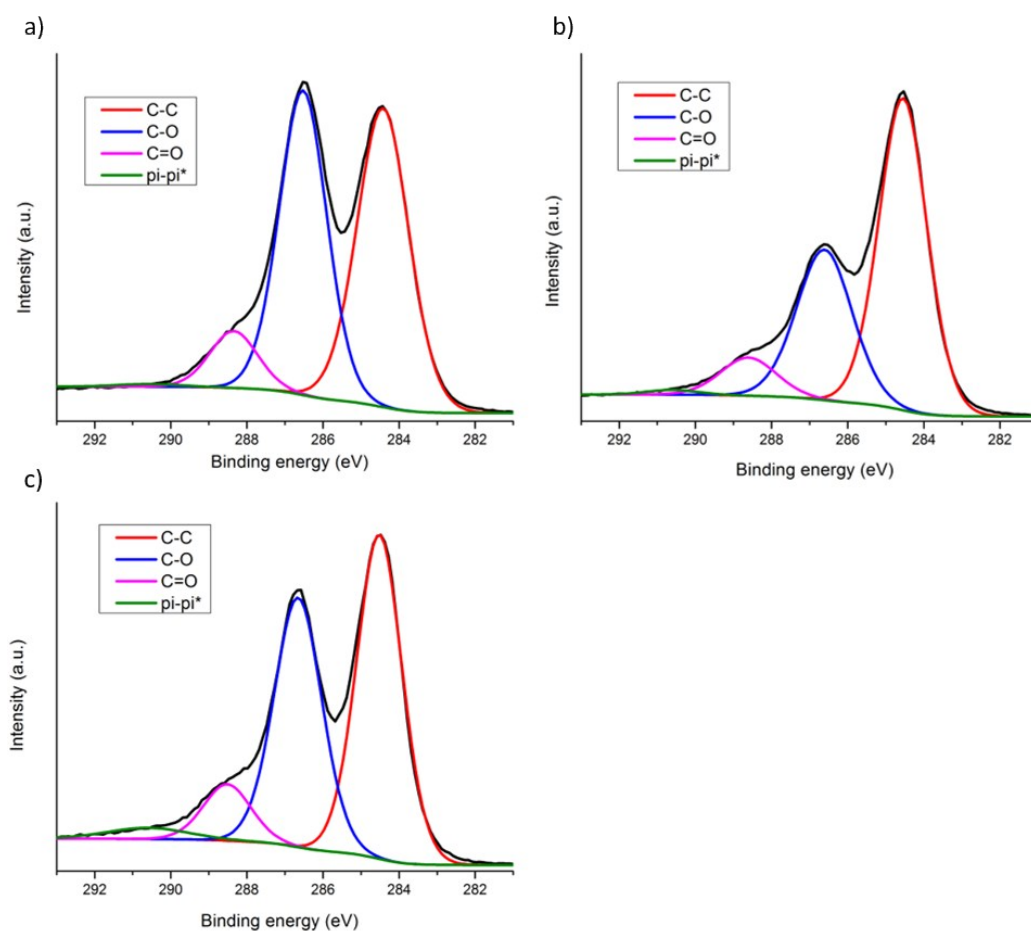


Figure S4. Deconvolution of the C(1s) peak for GO₅ (a), rGO₅ (b), and rGO₅-O₃ (c).

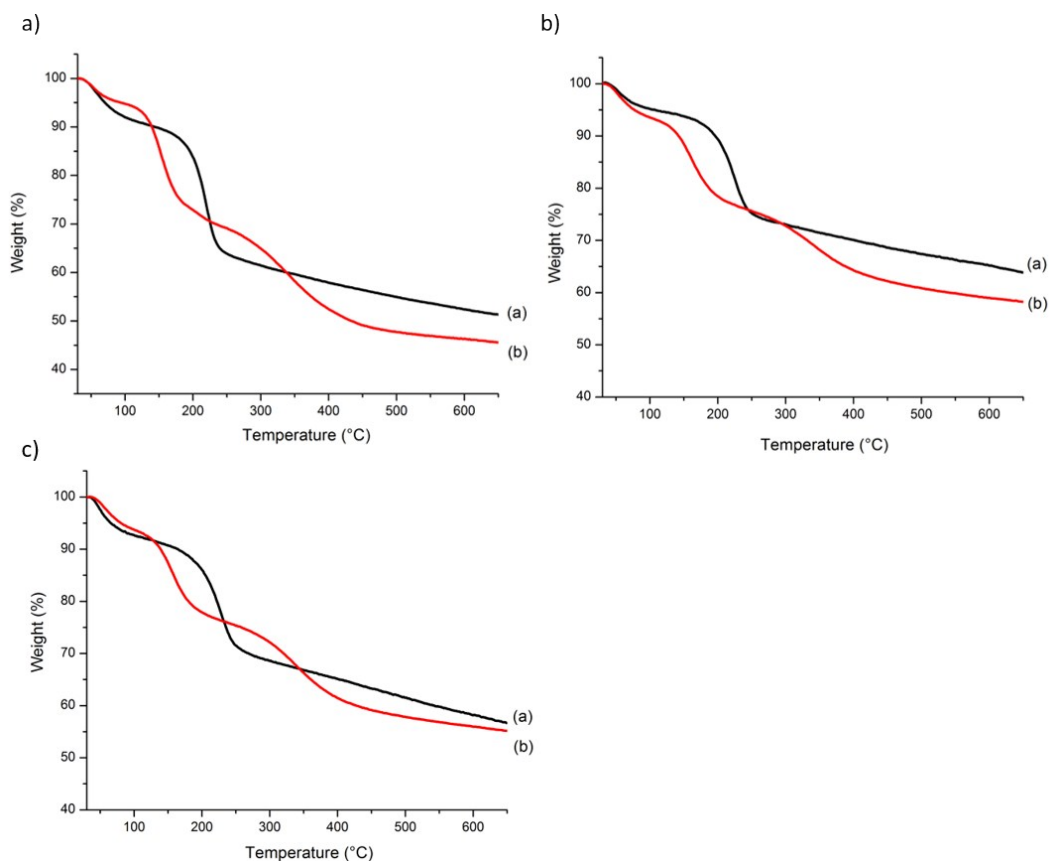


Figure S5. TGA spectra of PEI functionalized GO: a) GO_5 in black (a), $\text{GO}_5\text{-PEI}$ in red (b); b) rGO_5 in black (a), $\text{rGO}_5\text{-PEI}$ in red (b); c) $\text{rGO}_5\text{-O}_3$ in black (a), $\text{rGO}_5\text{-O}_3\text{-PEI}$ in red (b). GO displays a typical thermogram with degradation in three steps. The significant weight loss below 100 °C is ascribed to mainly the desorption water. PEI decomposition was recorded above 300 °C. We estimate a functionalization between 6 and 10% in weight of PEI for the GO_L , similar to that estimated for GO_5 [1].

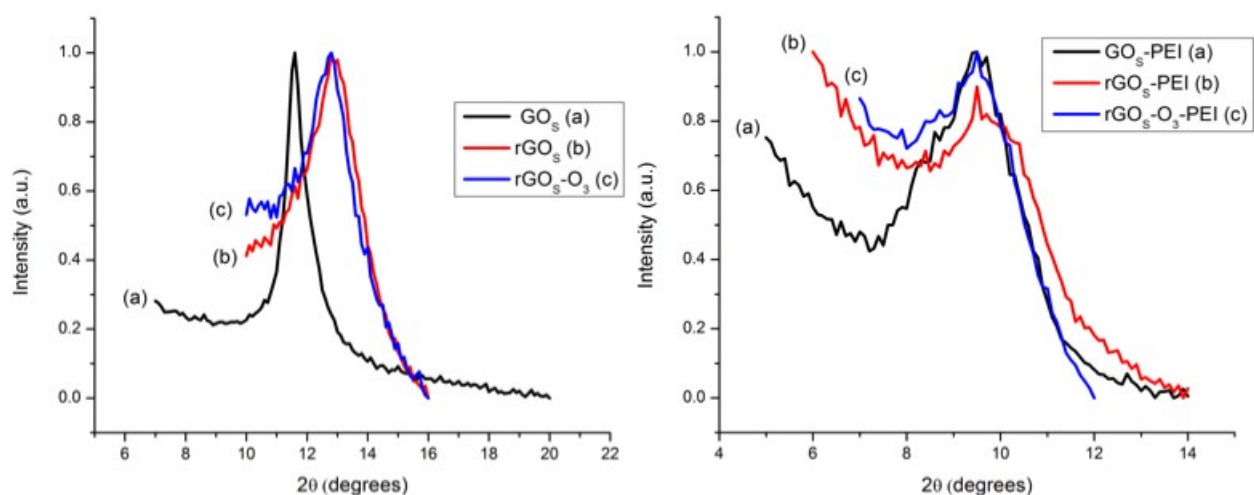


Figure S6. High resolution XRD spectra of the GO_5 series. On the left: a) GO_5 (black line), b) rGO_5 (red line), and (c) $\text{rGO}_5\text{-O}_3$ (blue line); on the right: a) $\text{GO}_5\text{-PEI}$ (black line), b) $\text{rGO}_5\text{-PEI}$ (red line), and (c) $\text{rGO}_5\text{-O}_3\text{-PEI}$ (blue line).

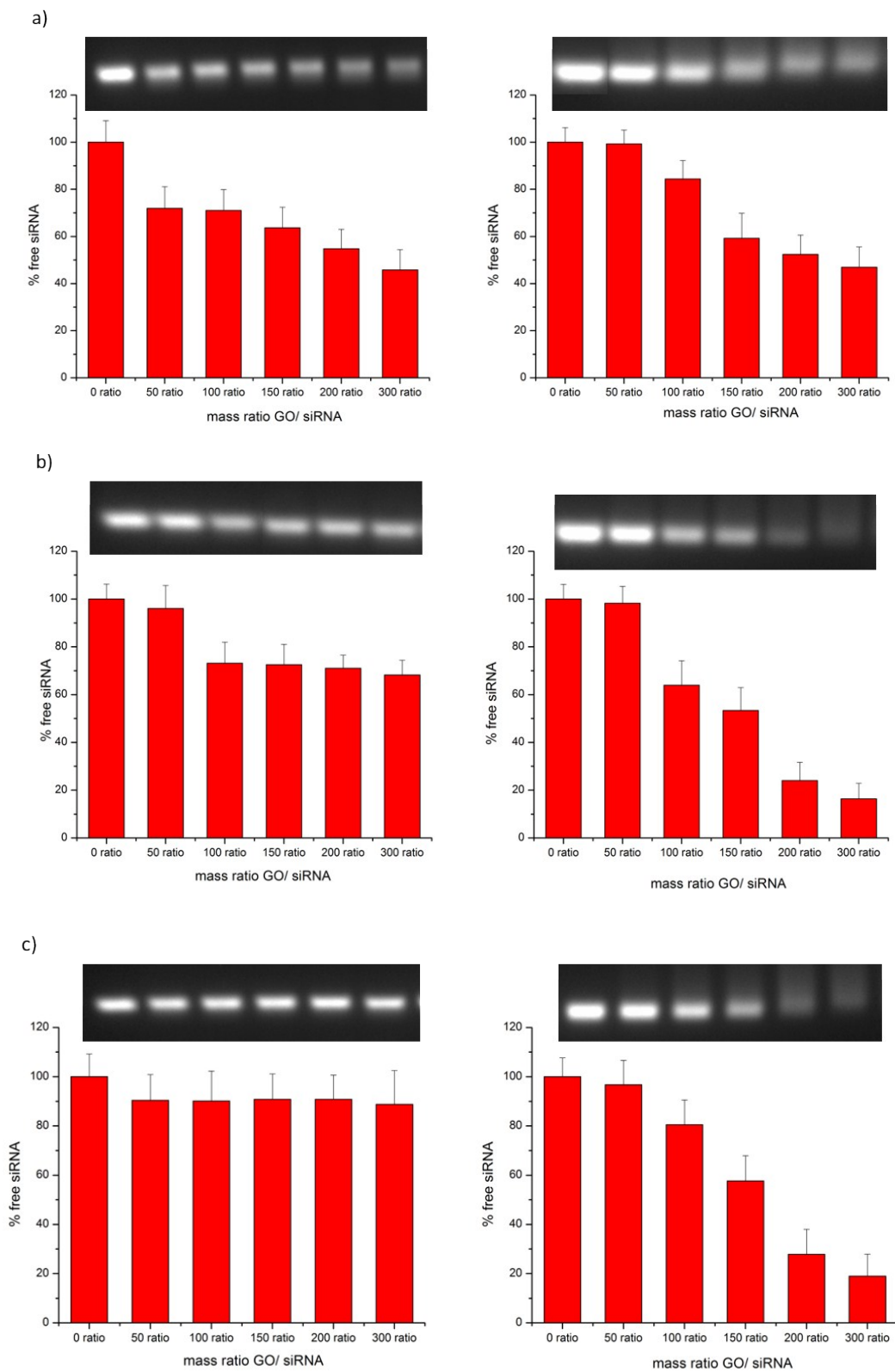


Figure S7. Complexation of different graphene derivatives of GO₅ (first column) and their corresponding PEI functionalized (second column): row a) GO₅, row b) rGO₅, row c) rGO₅-O₃. Top: image of the electrophoresis gel; bottom: histograms showing the free siRNA signal at different GO/siRNA mass ratios.

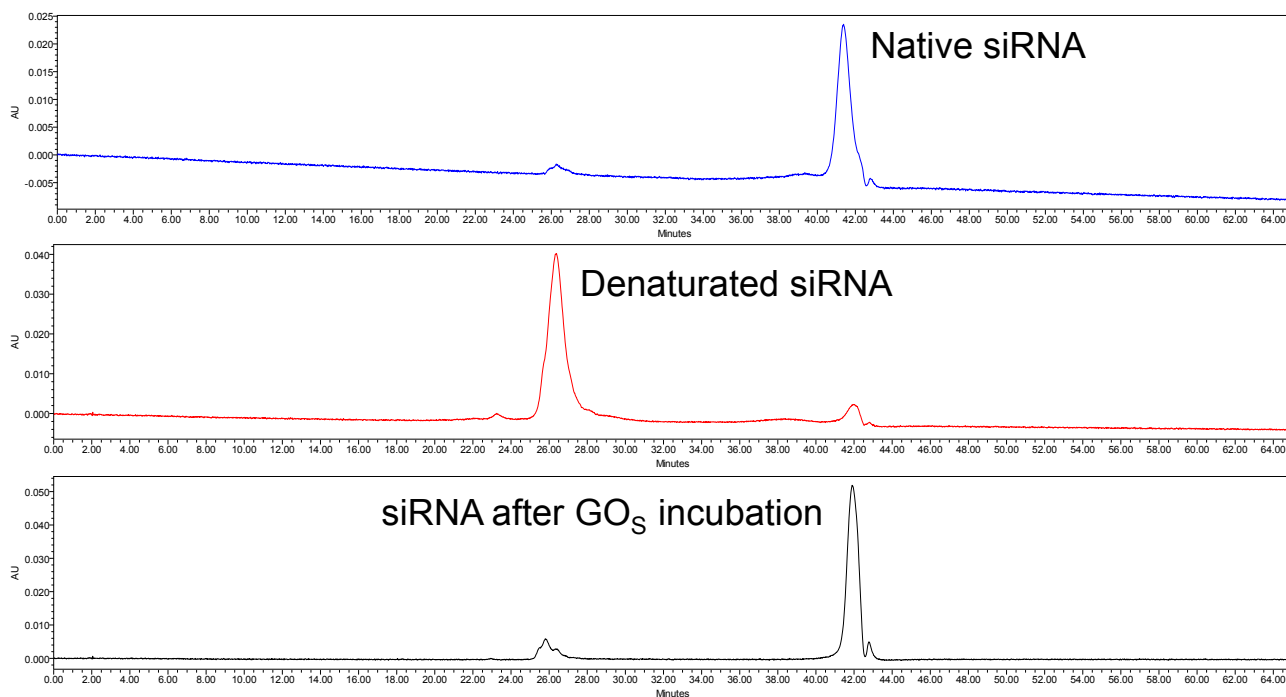


Figure S8. HPLC analysis of Alexa Fluor® 647 labelled siRNA: blue curve corresponds to native siRNA, red curve correspond to the siRNA denatured by formaldehyde, and black curve corresponds to siRNA after incubation with GO_S.

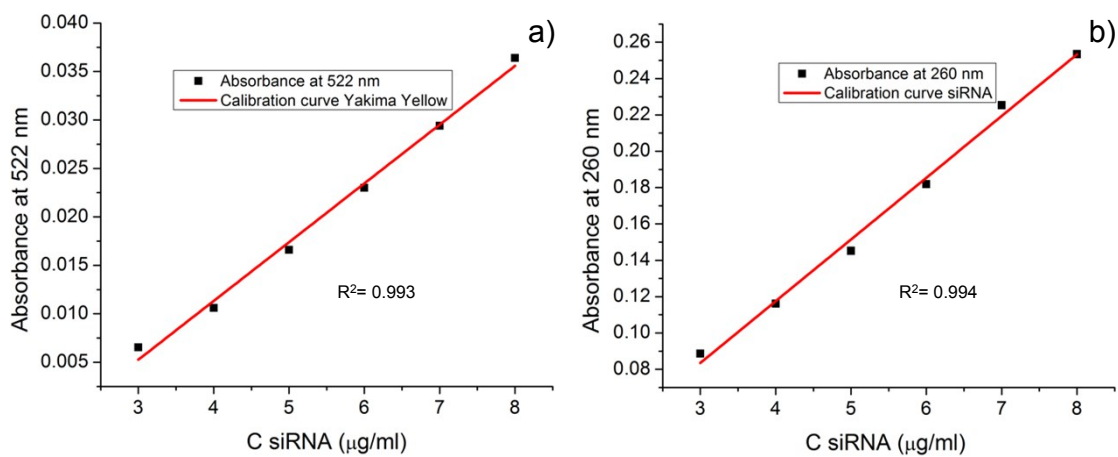


Figure S9. a) Calibration curve of Yakima Yellow signal at 522 nm; b) calibration curve of siRNA signal a 260 nm.

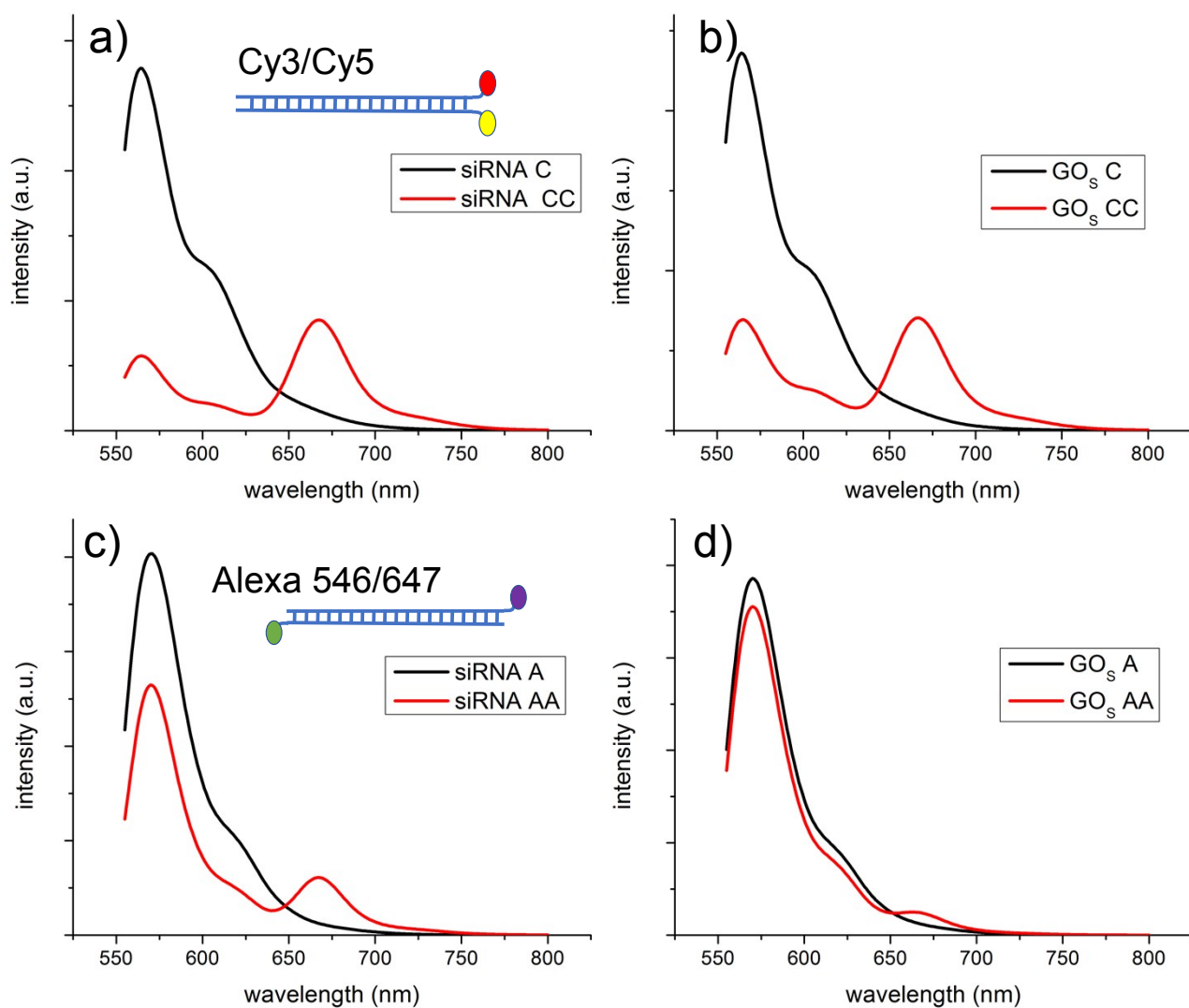


Figure S10 Emission spectra of a) 3'-Cy3 (black line) and 3'-5'-Cy3/Cy5 (red line) labelled untouched siRNA and b) after incubation with GO_s. c) Emission spectra of 5'-Alexa Fluor[®]546 (black line) and 5'-5'-Alexa Fluor[®]546/ Alexa Fluor[®]647 (red line) labelled untouched siRNA and d) after incubation with GO_s. Mass ratio siRNA/GO corresponds to 1:20. Excitation wavelegth 523 nm.

Table S1. N atomic % of PEI functionalized GO_s calculated from the XPS peak N1s at 400 eV.

	Small GO		
	Starting	Hydrothermal	Re-epoxidation
PEI-functionalized	GO _s -PEI	rGO _s -PEI	rGO _s -O ₃ -PEI
N atomic %	7.4 ± 0.1	5.9 ± 0.1	10.1 ± 0.1

NOTE: The values for large GO_L are reported in our previous publication [1].

Table S2. [002] Calculated distance from XRD using Bragg law, $\lambda = 1.541\text{\AA}$.

Sample	XRD max. (2 θ)	Distance (\AA)
GO _s	11.6	7.6
rGO _s	12.9	6.8
rGO _s -O ₃	12.8	6.9
GO _s -PEI	9.5	9.3
rGO _s -PEI	9.5	9.3
rGO _s O ₃ -PEI	9.5	9.3

Equations (1) and (2):

$$E = 1 - \frac{A_D I_{DA}}{A_{DA} I_D} \quad (1)$$

$$E = \frac{R_0^6}{R^6 + R_0^6} \quad (2)$$

A_D and A_{DA} are the UV absorbance at the excitation wavelength of the donor (535 nm) alone and of the donor in the presence of the acceptor, respectively, while I_D and I_{DA} are the fluorescence intensities of the solution of the donor alone and the donor in the presence of the acceptor at the maximum (570 nm for Alexa Fluor[®]546, and 565 nm for Cy[®]3). The FRET distances (R) have been calculated following equation (2), where R is the distance between the donor and the acceptor, and R_0 is the Förster radius distance.

References

[1] N. D. Q. Chau, G. Reina, J. Raya, I. A. Vacchi, C. Ménard-Moyon, Y. Nishina and A. Bianco, *Carbon*, 2017, **122**, 643–652.