

Supporting Information

A self-assembling peptidic platform for the delivery of oligonucleotides to the cell nucleus

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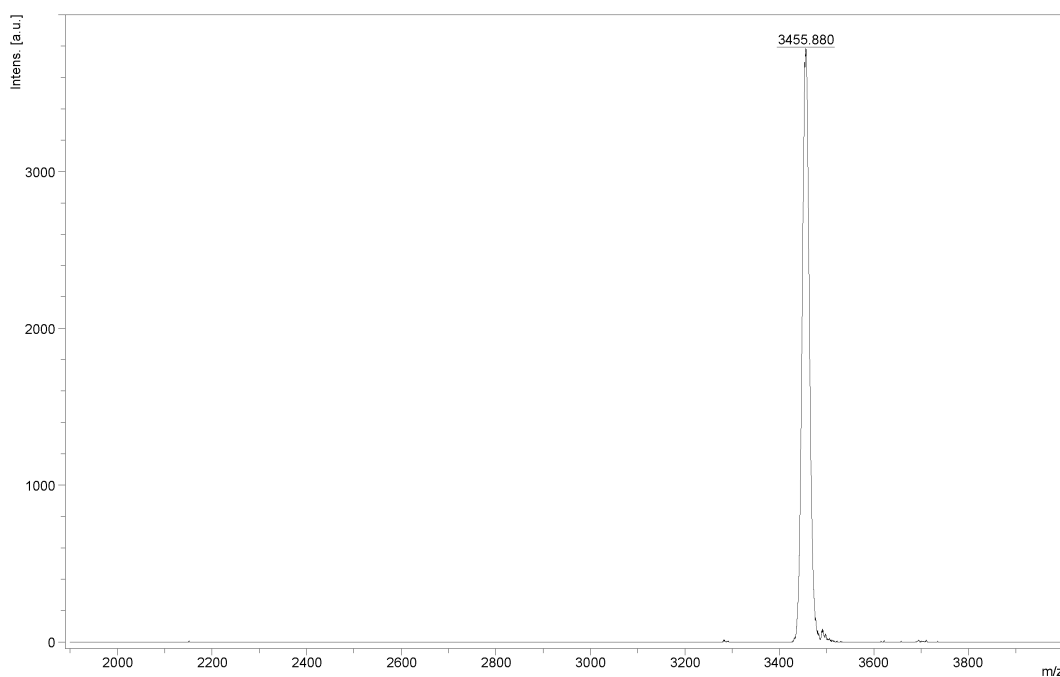


Figure S1. MALDI-TOF mass spectrum of NLS-peptide.

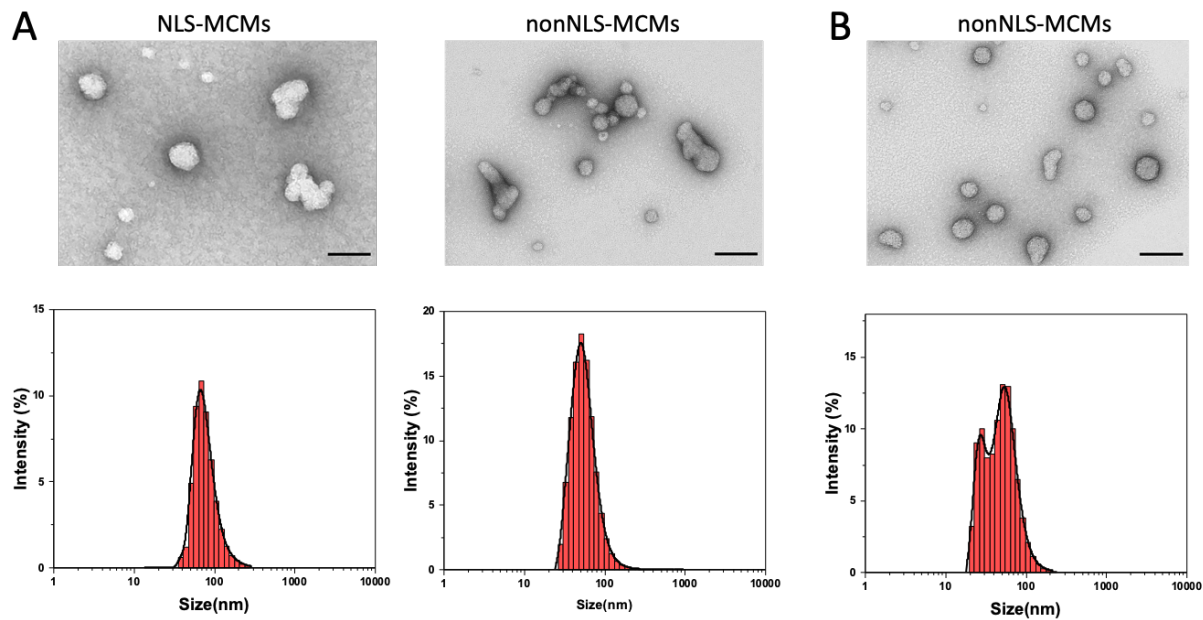


Figure S2. Ultrastructural morphology and size distribution of self-assembled NLS- and nonNLS-MCMs with different DNA payloads, A) scrambled, and B) G3139-GAP ASO. Scale bar, 100nm.

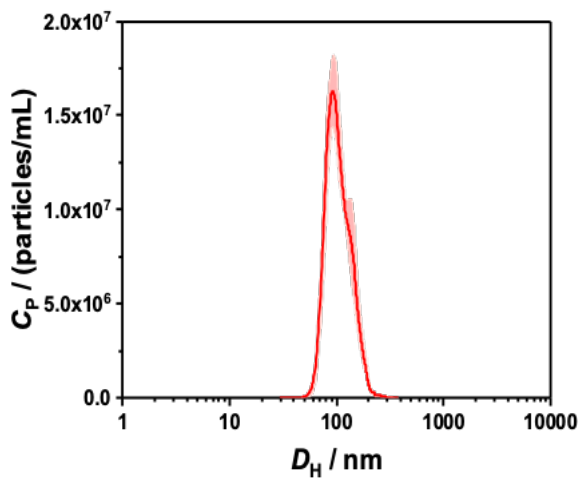


Figure S3. Hydrodynamic diameter of 22nt ssDNA-loaded NLS-MCMs determined by NTA.

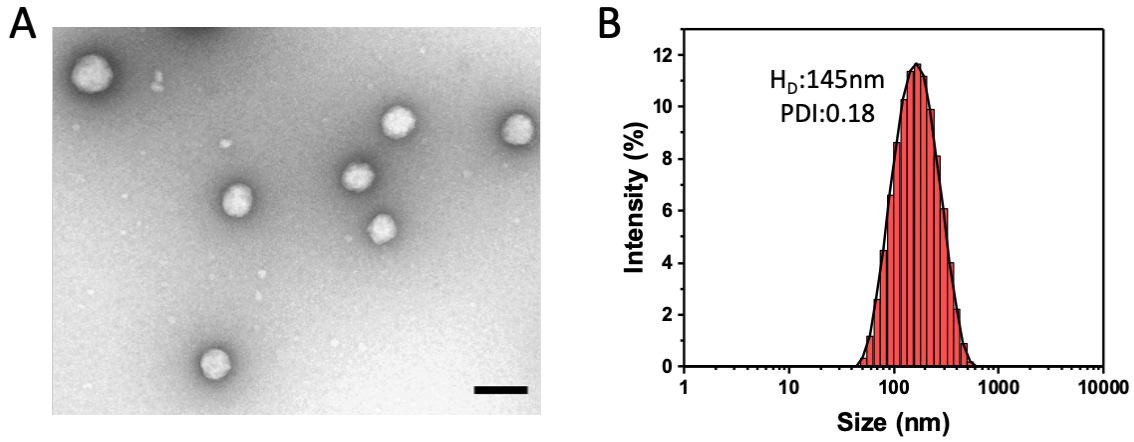


Figure S4. Characterization of self-assembled DNA-free NLS-MCMs. A) TEM, and B) DLS. Scale bar, 200nm.

Table S1. Characterization of NLS- and nonNLS-MCMs loaded with different oligonucleotides, suspended in water, at pH 7.

MCM-NPs	PDI	$D_n(\text{nm})$	Zeta potential(mV)
Scrambled oligo loaded NLS-MCMs	0.19	76 ± 2	21 ± 7
G3139-GAP loaded nonNLS-MCMs	0.28	52 ± 14	10 ± 4
Scrambled oligo loaded nonNLS-MCMs	0.18	58 ± 8	8 ± 3

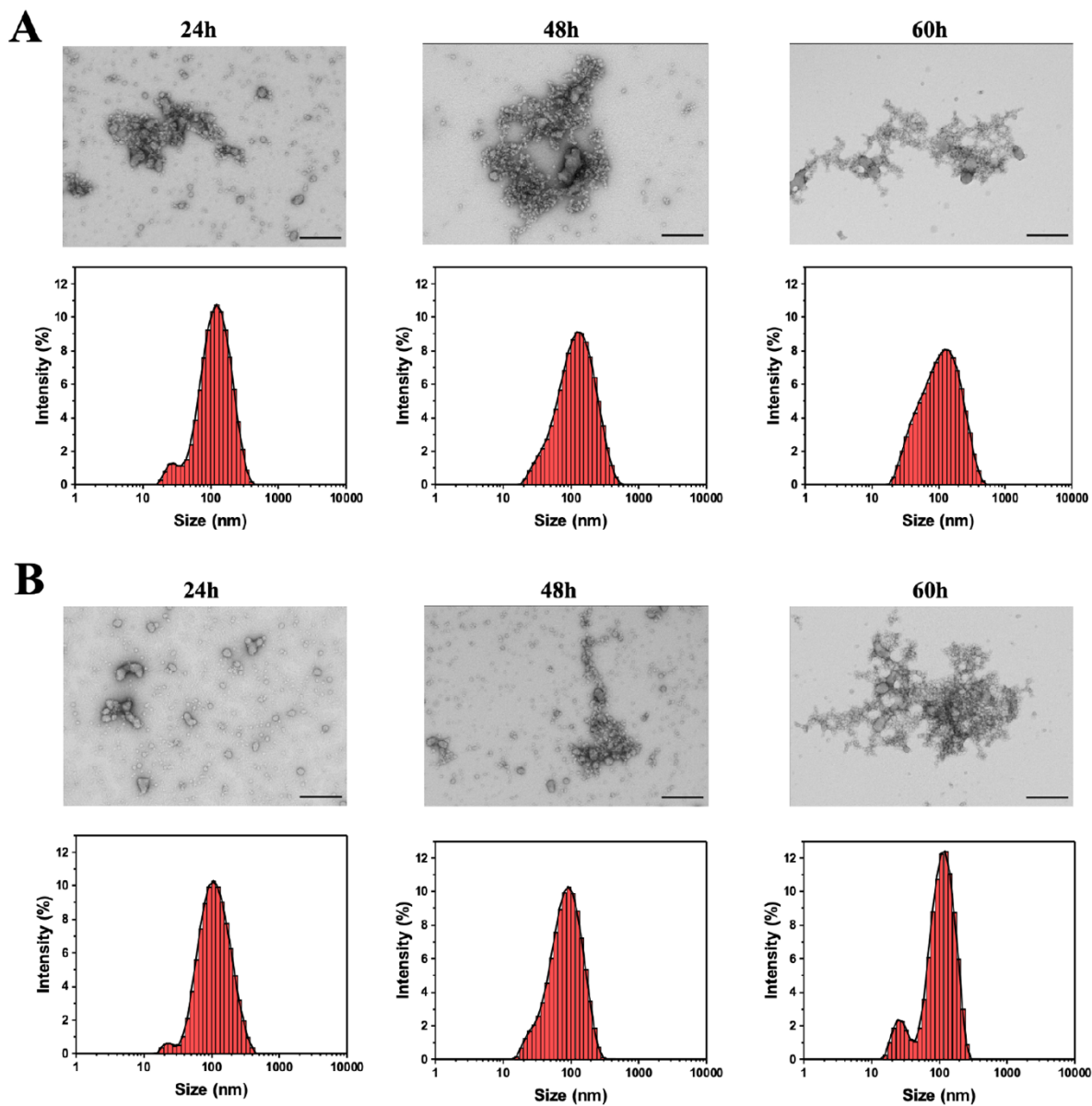


Figure S5. Thermo-responsiveness of NLS-MCMs. TEM and DLS analysis of NLS-MCMs A) with DNA, and B) without DNA after 24, 48 and 70 h incubation at 37 °C. Scale bars = 200 nm.

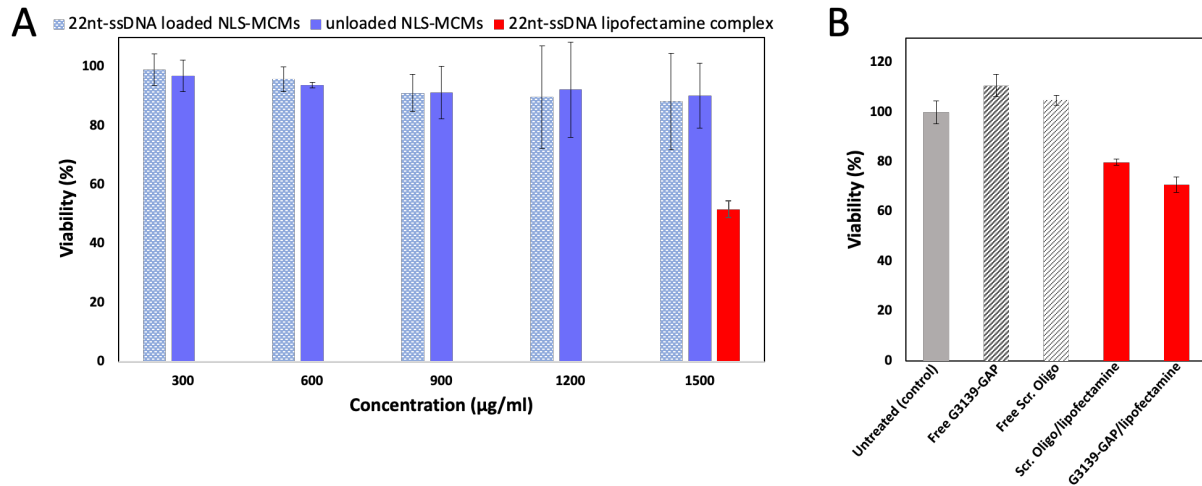


Figure S6. A) Effect of DNA-free and 22nt ssDNA-loaded NLS-MCMs on MCF-7 cell proliferation at different peptide concentrations after 24 h of exposure. Comparison at 1500µg/ml peptide concentration to lipofectamine complex with corresponding 22nt ssDNA concentration. B) lipofectamine-mediated transfection of scrambled oligonucleotides and G339-GAP ASO after 48 h of exposure.

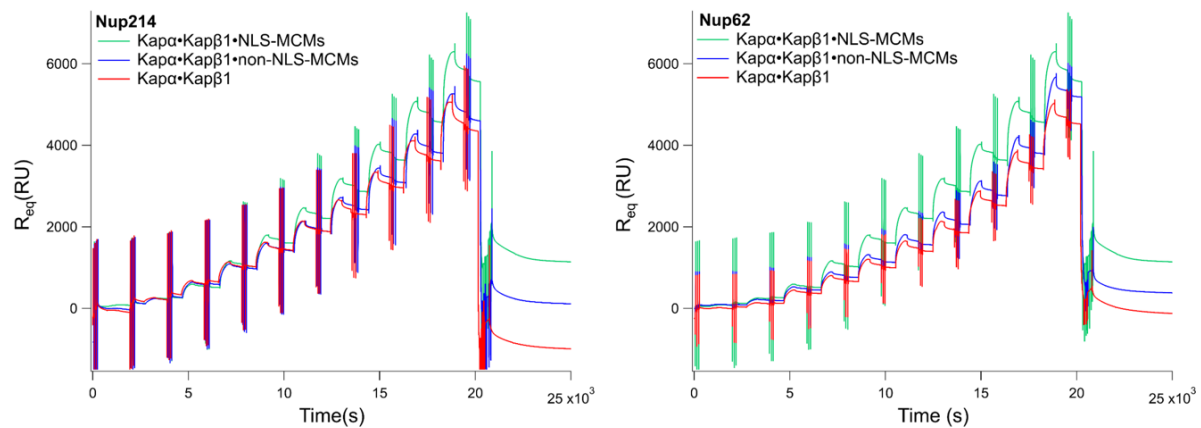


Figure S7. SPR sensograms resolve the binding interactions of Kap α •Kap β 1•NLS-MCMs, Kap α •Kap β 1•nonNLS-MCMs, and standalone Kap α •Kap β 1 to Nup214 (left) and Nup62 (right).

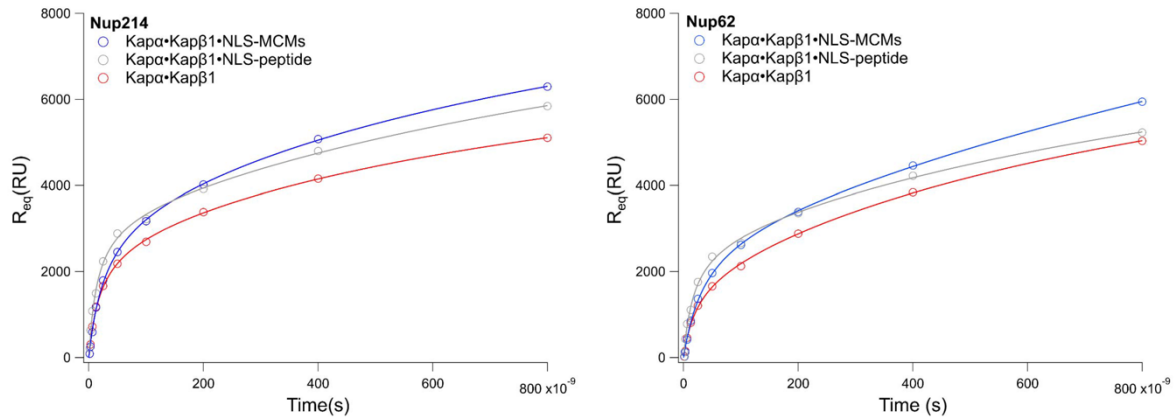


Figure S8. Langmuir isotherm fits (solid lines) for Kap α •Kap β 1•NLS-MCMs compared to free NLS-peptide. MCM nanocarriers elicit the highest FG Nup-binding response compared to free peptide as measured by SPR.

Table S2 Maximal SPR response signals and equilibrium dissociation constants of free peptide/karyopherin complexes binding to Nups.

		R_{max1} (RU)	R_{max2} (RU)	K_{D1} (nM)	K_{D2} (nM)
Kap α •Kap β 1 / NLS (free peptide)	Nup214	3282.8±286	7809.3±3870	14.6±2.5	1574.6±1340
	Nup62	2706±283	7122±2700	17.7±3.4	1396.3±990
Kap α •Kap β 1 / nonNLS (free peptide)	Nup214	2814.9±384	5748.8±1690	14.1±3.6	878.8±637
	Nup62	1804.6±226	4764.2±485	14.5±3.2	637.2±216
Kap α •Kap β 1	Nup214	2672±387	5711.4±1850	19.4±4.8	1028.1±779
	Nup62	1895±229	7495.2±1420	22.4±5.7	1077.3±477

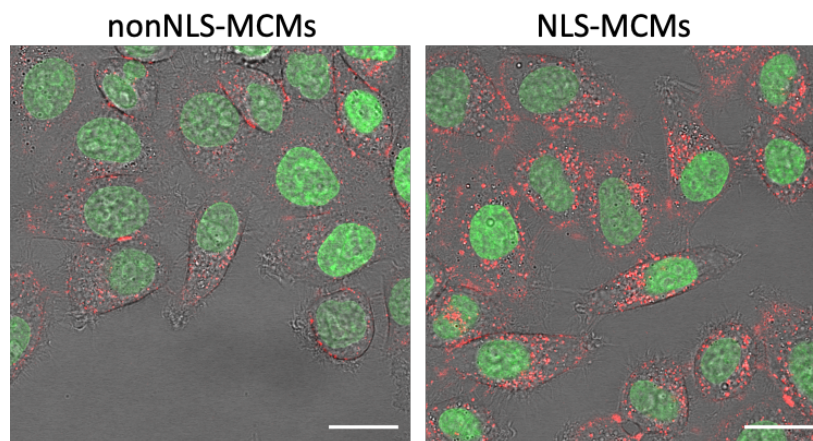


Figure S9 CLSM merged images of H2B-GFP expressing HeLa cells recorded after 5 h incubation with DNA-loaded NLS- and nonNLS-MCMs. Scale bars, 20 μ m.

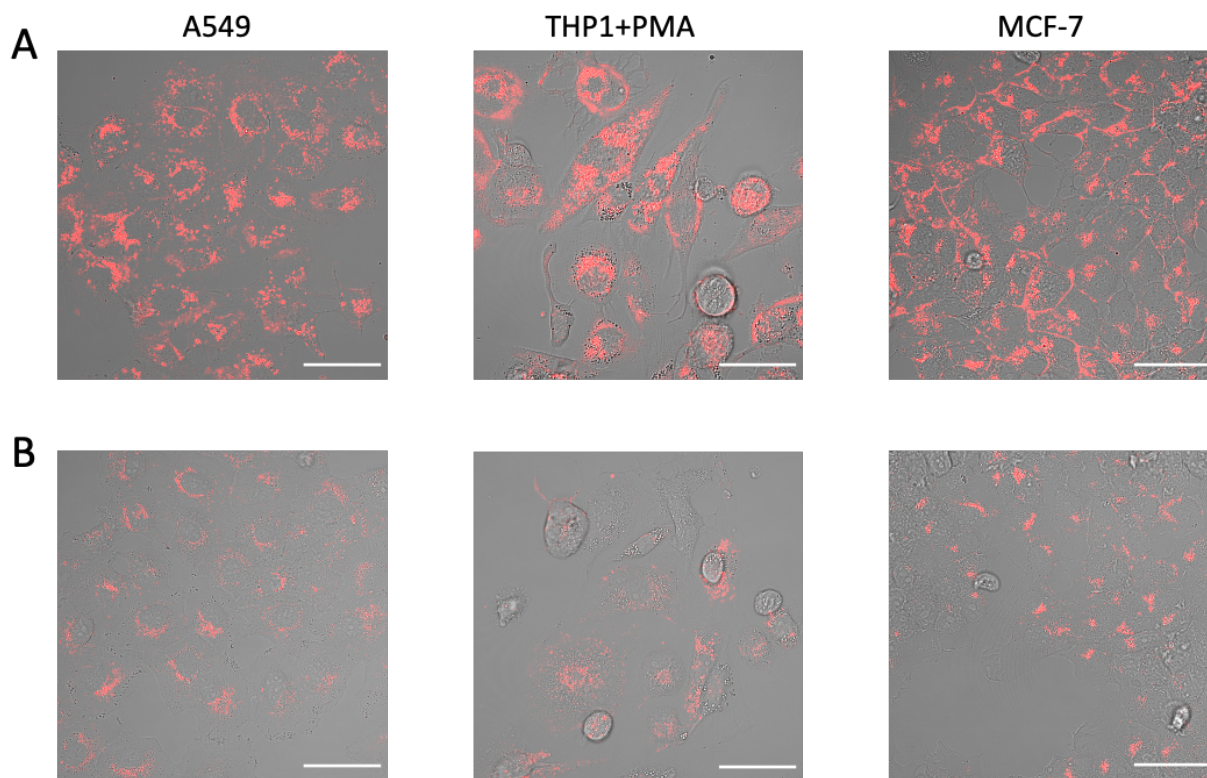


Figure S10. CLSM merged images of A549, THP1-stimulated with PMA and MCF-7 cells recorded after 24 h incubation with G3139-GAP-loaded A) NLS-MCMs, and B) nonNLS-MCMs. Scale bars = 50 μ m.

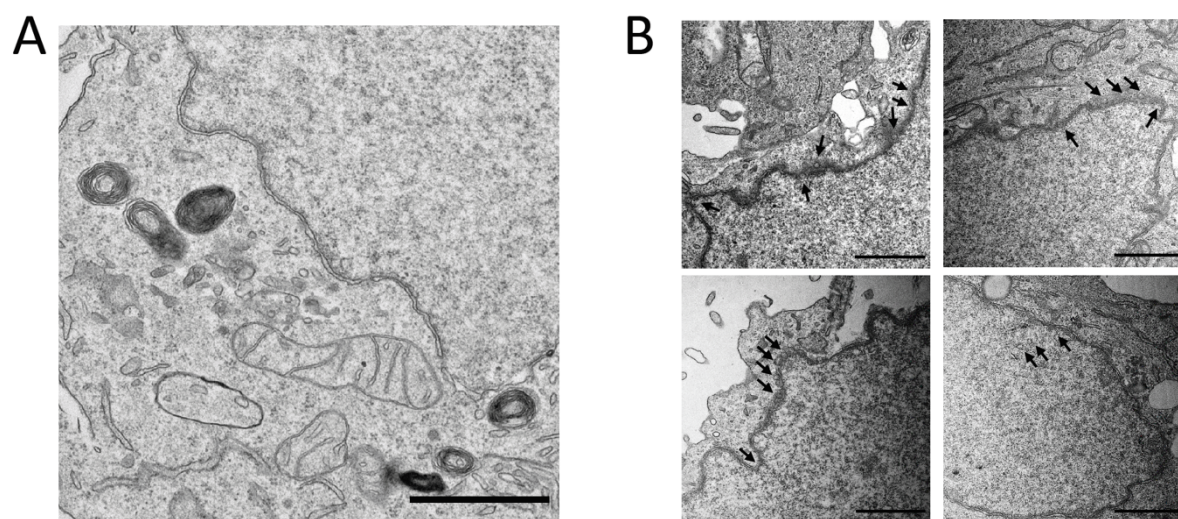


Figure S11. Ultrastructural analysis of A) untreated cells (control), B) cells treated with DNA-loaded NLS-MCMs, Scale bars, 1 μ m.

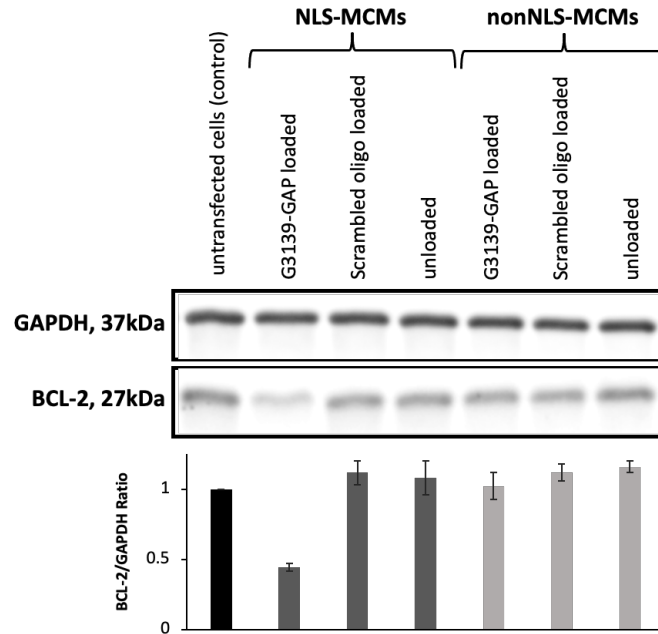


Figure S12. Western blot analysis of BCL-2 proteins in 6 μ g of MCF-7 total cell extract after 24 h. GAPDH was used as loading control. Quantitative densitometry of the immunoblots are mean \pm s.d. of the relative intensity of the bands from three independent assays; nonNLS-MCM-treated (dark grey bars), NLS-MCM-treated (dark grey bars), untreated (black bar).

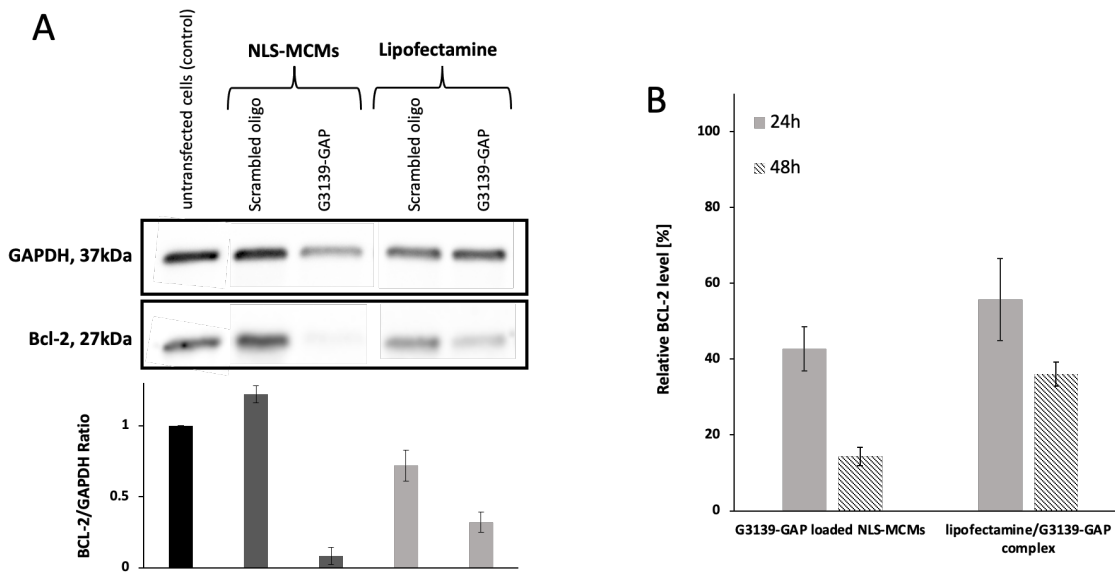


Figure S13. Therapeutic effects of cells transfected with lipofectamine/G3139-GAP complex. A) Western blot analysis of BCL-2 proteins in 6 μ g of MCF-7 total cell extract after 48 h. GAPDH was used as loading control. B)

Quantitative densitometry of the immunoblots. Lipofectamine-treated (light grey bars), NLS-MCM-treated (dark grey bars), untreated (black bar). C) BCL-2 expression levels in cells transfected with G3139-GAP relative to scrambled oligo delivered with NLS-MCMs or lipofectamine complexes. Data are mean \pm s.d. of the relative intensity of the bands from three independent assays.