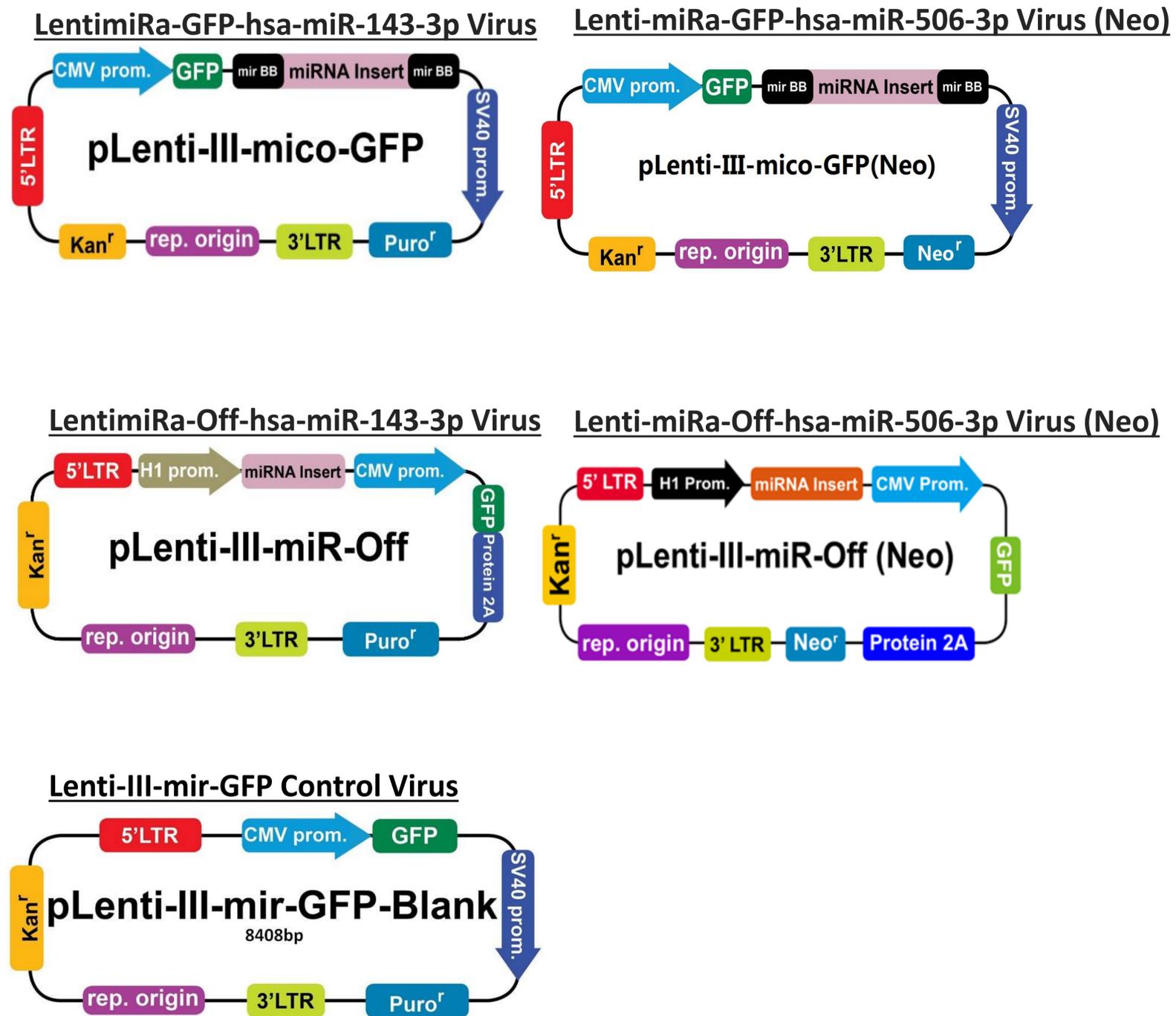


**Stable Dual miR-143 and miR-506 Upregulation Inhibits Proliferation and Cell Cycle Progression**

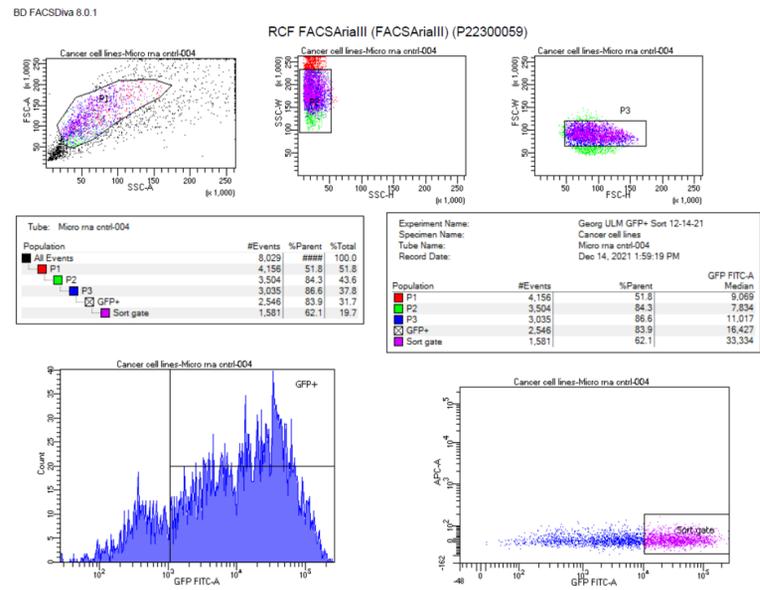
Archana Shrestha<sup>1</sup>, Behnaz Lahooti<sup>2</sup>, AKM Nawshad Hossian<sup>1</sup>, Mahboubeh Madadi<sup>3</sup>,  
Constantinos M. Mikelis<sup>2,4</sup>, George Mattheolabakis<sup>1,\*</sup>

Supplementary Figure S1: Vector maps for lentiviruses used to induce miRNA deregulation to the A549 cells. Vector maps were provided by the vendor (ABMgood).

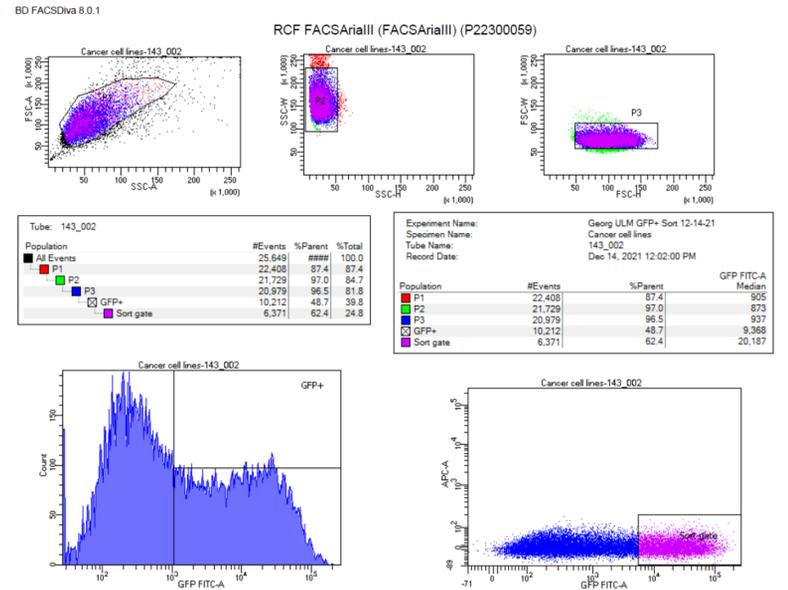


Supplementary Figure S2: Cell sorting of A549 cells transduced using lentiviral particles for miR deregulation. GFP signal was used for selecting cells.

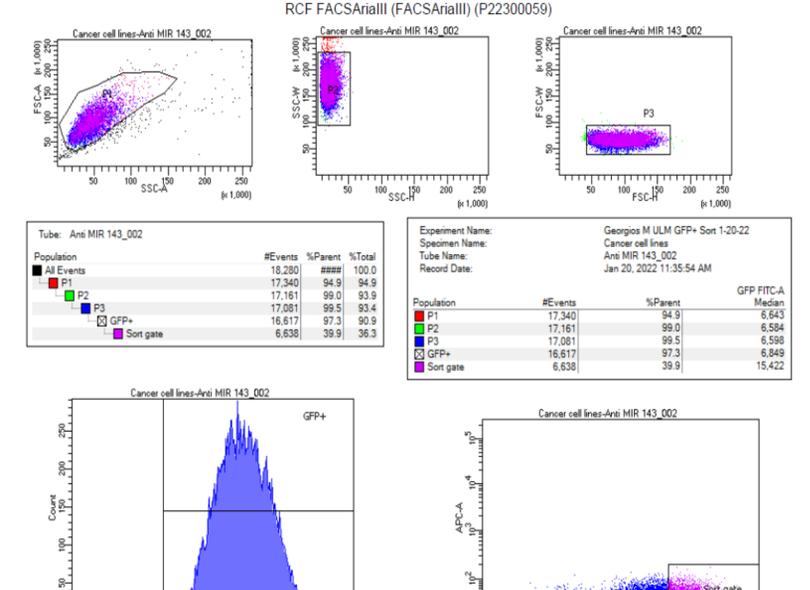
Sorting analysis of A549-miR-control cells



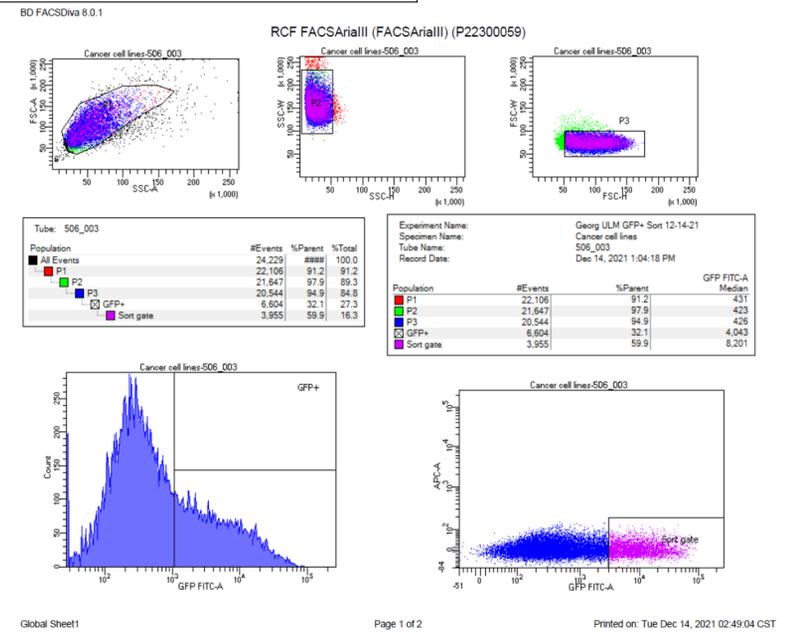
Sorting analysis of A549-miR-143 cells



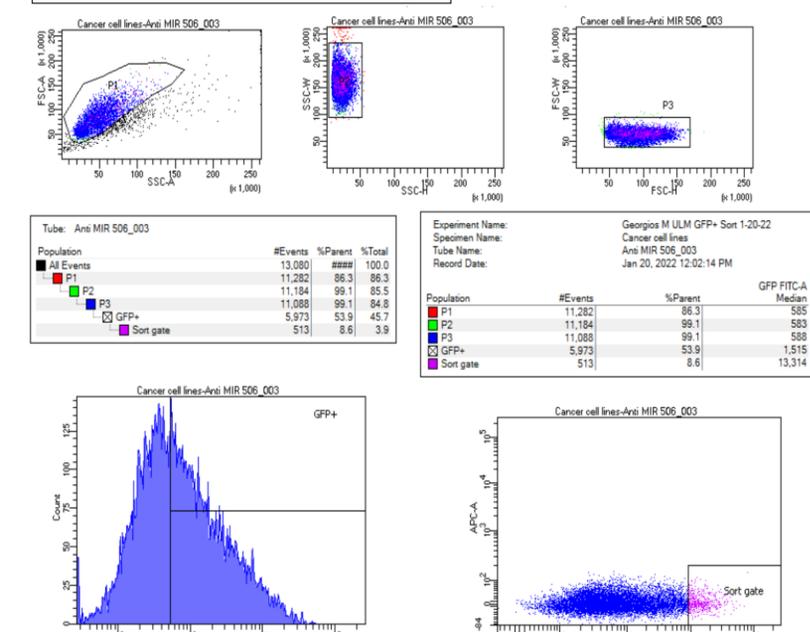
Sorting analysis of A549-anti-miR-143 cells



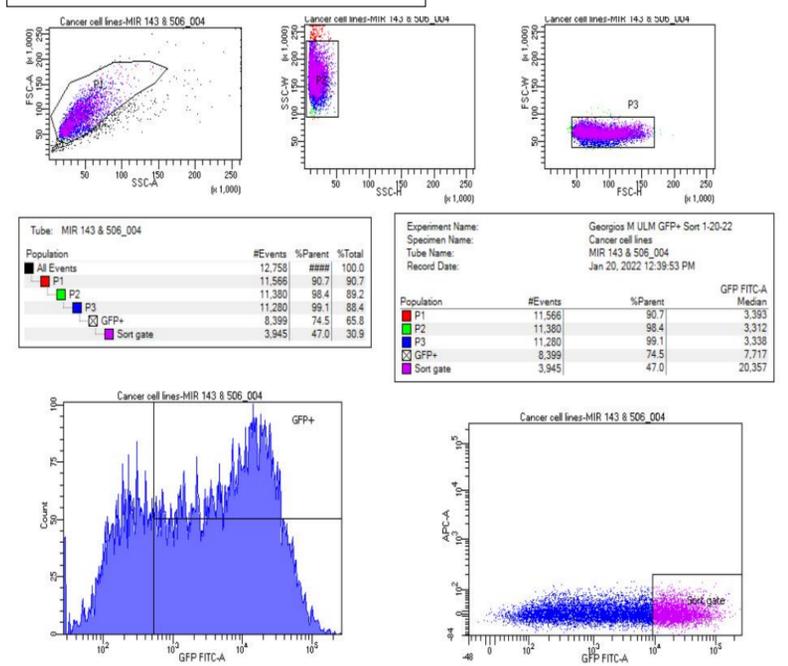
Sorting analysis of A549-miR-506 cells



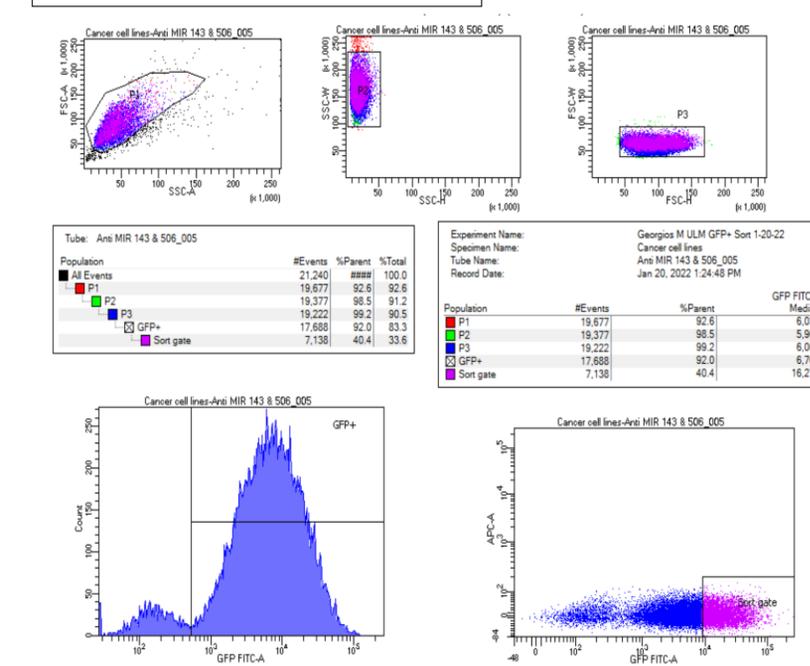
Sorting analysis of A549-anti-miR-506 cells



Sorting analysis of A549-miR-143/506 cells



Sorting analysis of A549-anti-miR-143/506 cells



Supplementary Figure S3: Cell authentication of A549 transduced cells for the different miR deregulations indicated 100% match to the parent A549 cell line.

% Match	ATCC® Cat. No.	Designation	D5S818	D13S317	D7S820	D16S539	vWA	TH01	AMEL	TPOX	CSF1PO
100	STRC2013	A549(m-GFP2)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12
100	CCL-185	A-549; Lung Carcinoma; Human (Homo sapiens)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12

% Match	ATCC® Cat. No.	Designation	D5S818	D13S317	D7S820	D16S539	vWA	TH01	AMEL	TPOX	CSF1PO
100	STRC2014	A549(ANTI_m-143)	11	11	8,11	11,12,13	14	8,9.3	X,Y	8,11	10,12
100	CCL-185	A-549; Lung Carcinoma; Human (Homo sapiens)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12

% Match	ATCC® Cat. No.	Designation	D5S818	D13S317	D7S820	D16S539	vWA	TH01	AMEL	TPOX	CSF1PO
100	STRC2015	A549(m-143)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12
100	CCL-185	A-549; Lung Carcinoma; Human (Homo sapiens)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12

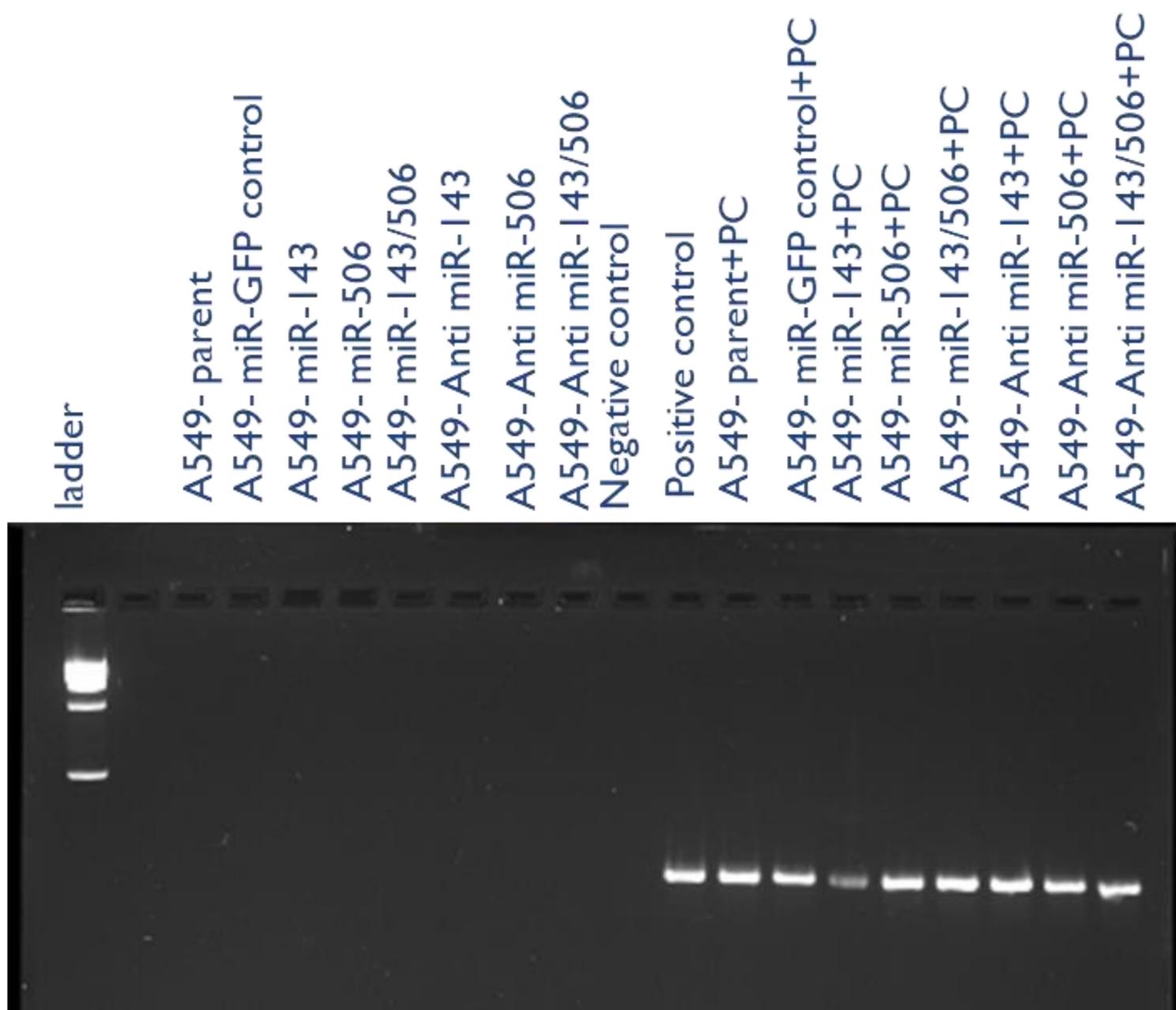
% Match	ATCC® Cat. No.	Designation	D5S818	D13S317	D7S820	D16S539	vWA	TH01	AMEL	TPOX	CSF1PO
100	STRC2016	A549(m-506)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12
100	CCL-185	A-549; Lung Carcinoma; Human (Homo sapiens)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12

% Match	ATCC® Cat. No.	Designation	D5S818	D13S317	D7S820	D16S539	vWA	TH01	AMEL	TPOX	CSF1PO
100	STRC2017	A549 (ANTI_m-506)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12
100	CCL-185	A-549; Lung Carcinoma; Human (Homo sapiens)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12

% Match	ATCC® Cat. No.	Designation	D5S818	D13S317	D7S820	D16S539	vWA	TH01	AMEL	TPOX	CSF1PO
100	STRC2018	A549 (m-143/506)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12
100	CCL-185	A-549; Lung Carcinoma; Human (Homo sapiens)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12

% Match	ATCC® Cat. No.	Designation	D5S818	D13S317	D7S820	D16S539	vWA	TH01	AMEL	TPOX	CSF1PO
100	STRC2019	A549 (ANTI_m-143/506)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12
100	CCL-185	A-549; Lung Carcinoma; Human (Homo sapiens)	11	11	8,11	11,12	14	8,9.3	X,Y	8,11	10,12

Supplementary Figure S4: Mycoplasma analysis of transduced A549 cells with the various miR deregulations. PC: positive control



Supplementary Figure S5: Representative cell cycle analyses images for the different miR deregulated A549 cells using ModFit LT 5.0.

