

Supplementary information

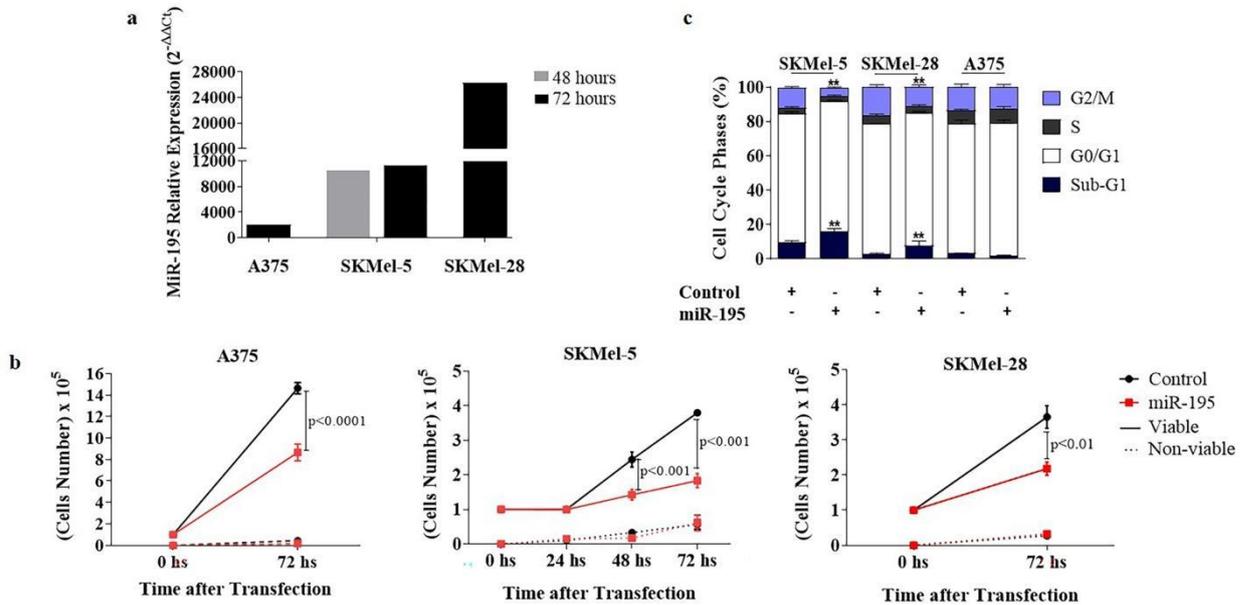


Figure S1. MiR-195-5p overexpression induces cytostatic and cytotoxic effect in human melanoma cells. **a.** miR-195-5p relative expression 48 and 72 h after transfection, compared to scramble cells. RNU48 was used as endogenous control. **b.** Cell number obtained by trypan blue count ($n = 5$). **c.** Cell cycle profile of propidium iodide (PI)-labeled cells ($n = 5$).

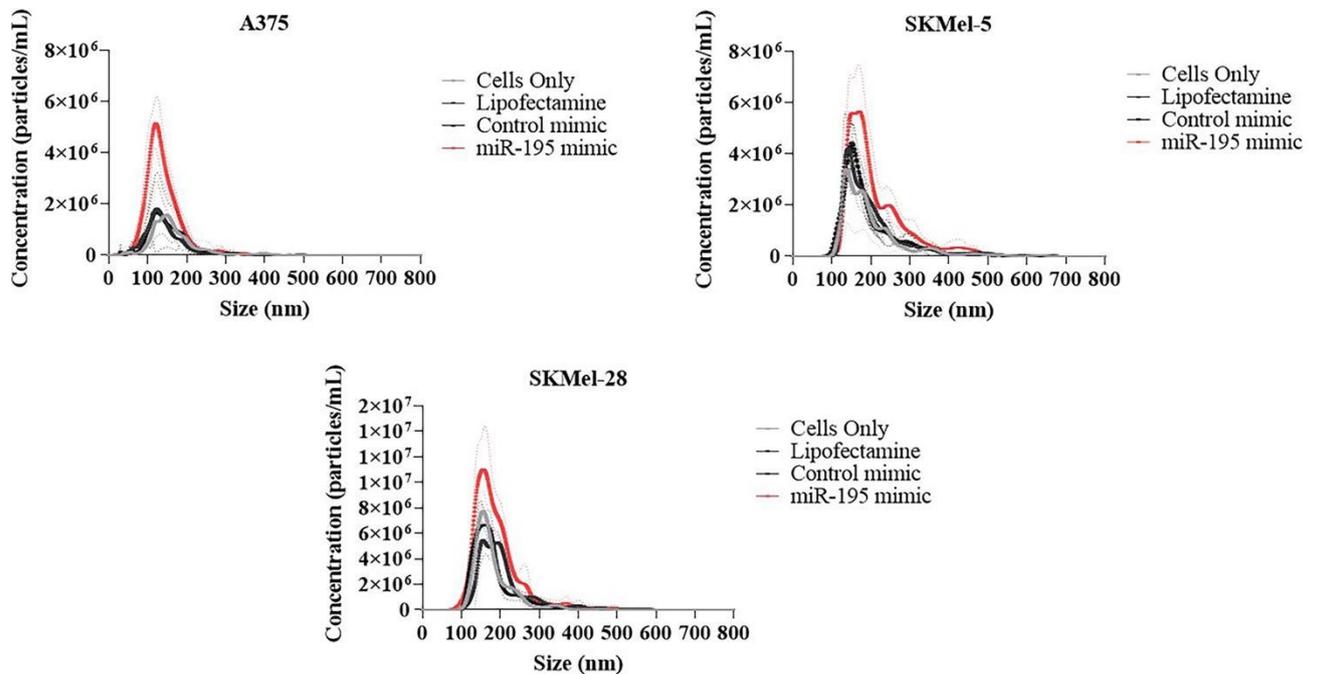


Figure S2. Vesiculation profile of control, lipofectamine and transfected cells confirming that induction of small EVs release is associated with miR-195-5p transfection.

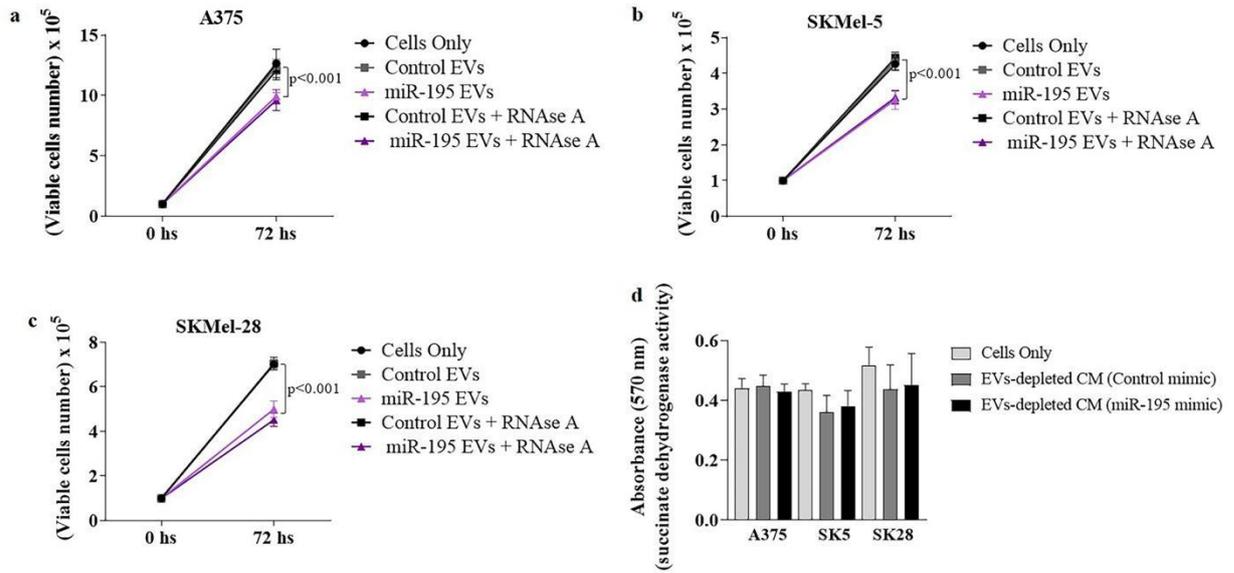


Figure S3. Cytostatic effect in *naïve* cells is exerted through EVs-mediated cargo transfer. **a-c.** Number of viable cells obtained by trypan blue count after EVs treatment with RNase A (n = 3). **d:** Cell viability, measured by MTT assay, after incubation with EVs-depleted medium for 72 h (n = 8).

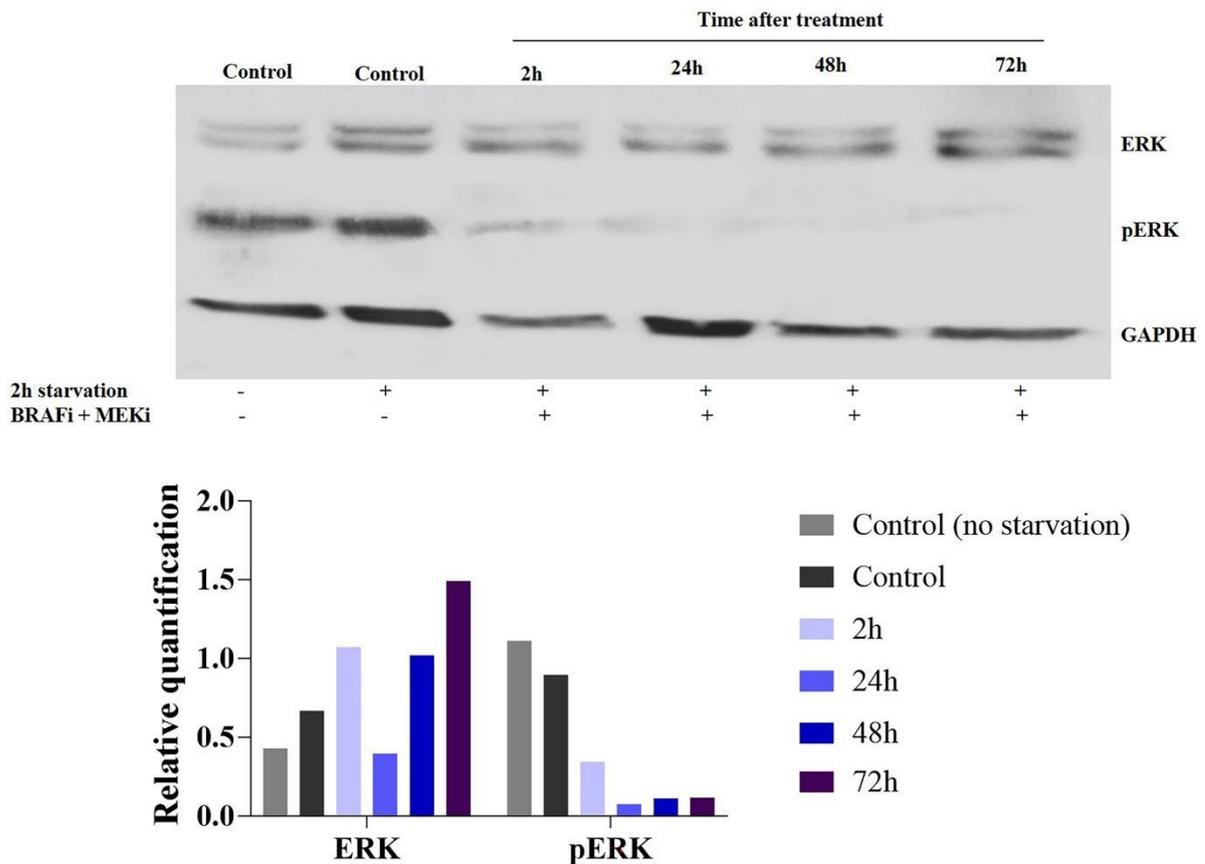


Figure S4. Inhibition of ERK phosphorylation after MAPKi treatment.

Western blot showing protein levels of total ERK and phospho-ERK after MAPKi treatment. Cells were previously starved for 2 h (cultured without FBS).

Table S1. Primers sequences used for RT-qPCR.

Gene	Forward	Reverse
<i>β-ACTINA</i>	5'-AGAAAATCTGGCACCACA-3'	5'-AGAGGCGTACAGGGATAGCA-3'
<i>BCL2-L1</i>	5'-ACAGCAGCAGTTTGGATGC-3'	5'-GGTGATGTGGAGCCTGGGATG-3'
<i>CDC42</i>	5' - GCCCGTGACCTGAAGGCTGTCA - 3'	5' - TGCTTTTAGTATGATGCCGACACCA - 3'
<i>GAPDH</i>	5'-TGCACACCAACTGCTTAGC-3'	5'-GGTGGACTGTGGTCATGAG-3'
<i>HPRT</i>	5'-CCCTGGCGTCGTGATTAGT-3'	5'-TCTCGAGCAAGAGGTTTCAGT-3'
<i>RAB27A</i>	5' - AGAGGAGGAAGCCATAGCAC - 3'	5' - CATGACCATTTGATCGCACCAC - 3'
<i>RAB27B</i>	5' - GGAAGTGGCTGACAAATATGG - 3'	5' - CAGTATCAGGGATTTGTGTCTT - 3'
<i>RAB31</i>	5' - ATCTTTGGGCTGGGTTTG - 3'	5' - ATGGGCTCATTAGTGGGTAG - 3'

Table S2. List of genes analyzed by microfluidic RT-qPCR.

<i>ABCB1</i>	<i>COL6A2</i>	<i>MMP12</i>	<i>TGFB1</i>
<i>ABCC1</i>	<i>CXCR4</i>	<i>MMP9</i>	<i>TRAF6</i>
<i>ACTB</i>	<i>E2F1</i>	<i>MYC</i>	<i>TWIST1</i>
<i>ACVR1B</i>	<i>EGFR</i>	<i>NANOG</i>	<i>TWIST2</i>
<i>AKT1</i>	<i>EGR1</i>	<i>NFE2L2</i>	<i>VEGFA</i>
<i>ALDH1A1</i>	<i>EIF2AK3</i>	<i>NFE2L3</i>	<i>VIM</i>
<i>ATF4</i>	<i>EPHA2</i>	<i>NODAL</i>	<i>WNT3A</i>
<i>ATF6</i>	<i>ERN1</i>	<i>NOTCH1</i>	<i>WNT5A</i>
<i>ATM</i>	<i>FGF13</i>	<i>NOTCH4</i>	<i>YAP1</i>
<i>AXIN2</i>	<i>FGF2</i>	<i>PDCD1</i>	<i>YWHAZ</i>
<i>B2M</i>	<i>FGF2R</i>	<i>PHB</i>	<i>ZEB1</i>
<i>BAD</i>	<i>FLOT2</i>	<i>POU5F1</i>	<i>ZEB2</i>
<i>BAX</i>	<i>FOS</i>	<i>PTEN</i>	
<i>BBC3</i>	<i>GAPDH</i>	<i>RB1</i>	
<i>BCL2</i>	<i>GUSB</i>	<i>RHOA</i>	
<i>BCL2L1</i>	<i>HIF1A</i>	<i>RIPK2</i>	
<i>BIRC5</i>	<i>HMGB1</i>	<i>RIPK3</i>	
<i>CASP8</i>	<i>HPRT1</i>	<i>RPLP0</i>	
<i>CCND1</i>	<i>IGF1R</i>	<i>SNAI1</i>	
<i>CD274</i>	<i>JUN</i>	<i>SOX2</i>	
<i>CD44</i>	<i>KRAS</i>	<i>SOX7</i>	
<i>CDH1</i>	<i>LAMC2</i>	<i>STAT1</i>	
<i>CDH2</i>	<i>LATS1</i>	<i>STAT3</i>	
<i>CDH5</i>	<i>LATS2</i>	<i>STK3</i>	
<i>CDKN1A</i>	<i>MCL1</i>	<i>STK4</i>	
<i>CDKN2A</i>	<i>MDM2</i>	<i>TAZ</i>	
<i>CDKN3</i>	<i>MITF</i>	<i>TBK1</i>	
<i>CFLAR</i>	<i>MMP1</i>	<i>TFRC</i>	

Table S3. Antibodies used for WB.

Antibody	Dilution (titer)	Brand
Anti-Calnexin	1:500	Abcam – ab58504
Anti-CD63	1:700	Thermo Fischer Scientific - PA592370
Anti-CD9	1:500	Thermo Fischer Scientific - PA5-85955
Anti-BCL2-L1	1:1000	Thermo Fischer Scientific - MA5-15142
Anti-gERK1/2	1:10000	Sigma – M5670
Anti-pERK1/2	1:2000	Sigma – M8159
Anti-ACTIN	1:2000	Sigma – A5060
Anti-GAPDH	1:1000	Sigma – G8795
Anti-rabbit IgG Peroxidase	1:7000	Sigma – A9169
Anti-mouse IgM Peroxidase	1:4000	Sigma – M-6274

Table S4. Predicted targets and binding sites.

Gene	Predicted pairing
<i>BCL2-L1</i> (946-952) miR-202-3p	5'...CCCCAGGGUCU <u>UCCCUACCUCAG</u> ...3' 3' AAGGGUACGGGAU <u>AUGGAGA</u> 5'
<i>BCL2-L1</i> (2529-2536) miR-195-5p	5'...GAAUAUCCAAUCCUG <u>UGCUGC</u> A...3' 3' CGGUUAUAAAGAC <u>ACGACGAU</u> 5'
<i>YAP1</i> (162-168) miR-195-5p	5'...CUCU <u>UCCUUGUCCAUUGCUGC</u> ...3' 3' CGGUUAUAAAGAC <u>ACGACGAU</u> 5'
<i>VEGFA</i> (292-299) miR-195-5p	5'...CCAUUUUUUUUUCU <u>UGCUGC</u> A...3' 3' CGGUUAUAAAGAC <u>ACGACGAU</u> 5'
<i>CCND1</i> (1961-1967) miR-195-5p	5'...CCAUUUUUCUUAUUGC--- <u>GCUGC</u> UAC...3' 3' CGGUUAUAAAGACA <u>CGACGAU</u> 5'
<i>CCND1</i> (2033-2040) miR-195-5p	5' ...CUCU <u>UUCACAUUGUU-UGCUGC</u> A...3' 3' CGGUUAUAAAGAC <u>ACGACGAU</u> 5'
<i>CCND1</i> (2213-2220) miR-202-3p	5' ...GGU <u>UGCUGUUUCACA</u> AU <u>ACCUC</u> A...3' 3' AAGGGUACGGGAU <u>AUGGAGA</u> 5'
<i>CCND1</i> (2873-2879) miR-202-3p	5'...AACACGGCUCACGCU <u>UACCUC</u> AA...3' 3' AAGGGUACGGGAU <u>AUGGAGA</u> 5'
<i>CCND1</i> (2730-2750) miR-152-3p	5'...CCGCACGATTTCA <u>TTGAACA</u> ...3' 3' UCAGUGCAUGACAG <u>AACUUGG</u> 5'