

Table S1. Nomenclature of Extracellular Vesicles and Nanoparticles

Name	Category	EV class ^a	Size	Typical sedimentation	Protein markers	Biogenesis	Comments	Principal References
Classical exosome	Exosome	Small EV	40 - 150 nm	100,000 - 200,000 × g ^c	CD63, (CD81, CD9)	MVE	CD63 generally most accepted, CD81 and CD9 likely less specific but present. CD63 strongly labels MVEs ^h	Kowal et al 2016, van Niel et al 2018, this work
Non-classical exosome	Exosome	Small EV	40 - 150 nm	100,000 - 200,000 × g	? (CD63-, CD81- and CD9-negative)	MVE	Hypothetical CD63/CD81/CD9-negative exosome. Subpopulations of ILVs and MVEs may exist ⁱ	van Niel et al 2011, Edgar et al 2014, this work
Classical microvesicle	Microvesicle	Large EV	~ 150 - 1000 nm ^b	10,000 - 20,000 × g	Annexin A1, (ARF6, Annexin A2?)	Plasma membrane	This work identifies Annexin A1 as a marker of classical microvesicles	Muralidharan-Charivan et al 2009, this work
Large oncosome	Microvesicle	Large EV	1 - 10 µm	10,000 × g	Annexin A1, (ARF6, Annexin A2?)	Plasma membrane	This work identifies Annexin A1 as a likely marker of large oncosomes	Minciacci et al 2015, Minciacci et al 2017, this work
ARRM	Microvesicle	Small EV	~ 40 - 100 nm	100,000 - 200,000 × g	ARRDC1, (TSG101) ^f	Plasma membrane	Small EV-sized budding microvesicles that depend on ARRDC1 and TSG101 for biogenesis	Nabhan et al 2012, Wang and Lu 2017, this work
Apoptotic body	Apoptotic EV	Large EV	1 - 5 µm	1,000 - 10,000 × g	Annexin V (PS exposure)	Apoptosis	Generally accepted size range for apoptotic bodies	Hristov et al 2004, Crescitelli et al 2013,
Apoptotic vesicle	Apoptotic EV	Small to large EV	~ 100 - 1000 nm	10,000 - 200,000 × g ^d	Annexin V (PS exposure)	Apoptosis	Apoptosis-related vesicle, this work demonstrates classical exosomes do not express Annexin V ^j	Hristov et al 2004, Schiller et al 2008, this work
Autophagic EV	Autophagic EV	Small to large EV	40 - 1000 nm ^e	10,000 - 200,000 × g ^e	LC3B-PE, p62	Autophagy	Does not express CD63, CD81 and CD9, extracellular release is induced by inhibition of autophagic flux	Hessvik et al 2016, this work
Exomere	Nanoparticle	Non-EV	~ 35 - 50 nm	100,000 - 200,000 × g	HSP90, HSPA13 ^g	Unknown	Perhaps represents composite/aggregation of secreted proteins and extracellular debris	Zhang et al 2018, Zhang et al 2019, this work
NV fractions	Nanoparticle	Non-EV	? (vaults: ~ 70 nm)	100,000 - 200,000 × g	Fibronectin, Histones, MVP, HSPA13 ^g	Unknown	Perhaps represents composite/aggregation of secreted proteins and extracellular debris including vaults	This work

EV, extracellular vesicle; sEV, small extracellular vesicle; MVE, multivesicular endosome; SIM, Structured Illumination Microscopy; ARMM, arrestin-domain-containing protein 1 (ARRDC1)-mediated microvesicle; PS, phosphatidylserine; NV, non-vesicular.

^aThe terms Small EV (<200 nm) and Large EV (>200 nm) for lipid bilayer-delimited extracellular vesicles (EVs) are somewhat arbitrary but widely adopted. In this work, samples were passed through a 220 nm pore filter before isolation of small EV samples.

^bIn this work, we observe Annexin A1-positive vesicles down to about 150 nm in size using SIM, which is at the resolution limit of reliable identification of a vesicle budding from the plasma membrane. Better resolution may allow <150 nm classical microvesicles to be observed.

^cSignificant sedimentation of exosomes/sEVs occurs already at 33,000 × g (Jeppesen et al 2014a). Differential ultracentrifugation will produce enrichment of specific EVs but is incapable of generating samples of purely one type of EV.

^dPresumed, based on the size-range of vesicles bearing Annexin V.

^ePresumed, based on the size-range of vesicles bearing LC3B-PE.

^fIn this work, we demonstrate that at least some ARRDC1-positive sEVs also express CD9 and CD81. In particular, CD9-positive sEVs from the GLI36 cell line strongly expressed ARRDC1. The expression of TSG101 was very low in CD63- and CD81-positive sEVs but robust in CD9-positive sEVs.

^gHSP90 was associated with exomeres (Zhang et al 2018) and is absent from classical exosomes (this work). HSPA13 was associated with exomeres (Zhang et al 2018, Zhang et al 2019), and NV fractions (this work). Other heat shock proteins may be associated with exosomes or sEVs, including HSC70 (Kowal et al 2016, this work).

^hIn this work, we demonstrate that some CD63-positive organelles also stain for LC3B, indicative of amphisomes - the fusion products of MVEs and autophagosomes. However, CD63-positive exosomes do not express LC3B-PE.

ⁱThere is evidence for CD63-dependent and -independent ILVs (van Niel et al 2011), and for CD63-dependent and Hrs-dependent subpopulations of ILVs in the same MVEs (Edgar et al 2014). sEVs that are distinct from classical exosomes must be significant carriers of extracellular proteins previously associated with exosomes (this work), but these sEVs could be non-classical, MVE-derived, exosome.

^jAt least under conditions where apoptosis is not deliberately induced. In this work, we demonstrate that classical exosomes do express Annexin V, but Annexin V is present in gradient purified sEV fractions.

Table S1. Nomenclature of Extracellular Vesicles and Nanoparticles, Related to Figures 1-7.

Nomenclature of terms used to describe extracellular vesicles and nanoparticles in the context of this study. Note that the term 'extracellular vesicles' refers exclusively to lipid bilayer-membrane enclosed vesicles and thus distinct from non-membrane enclosed nanoparticles such as exomeres and the NV fractions, and from lipoprotein complexes such as HDL.

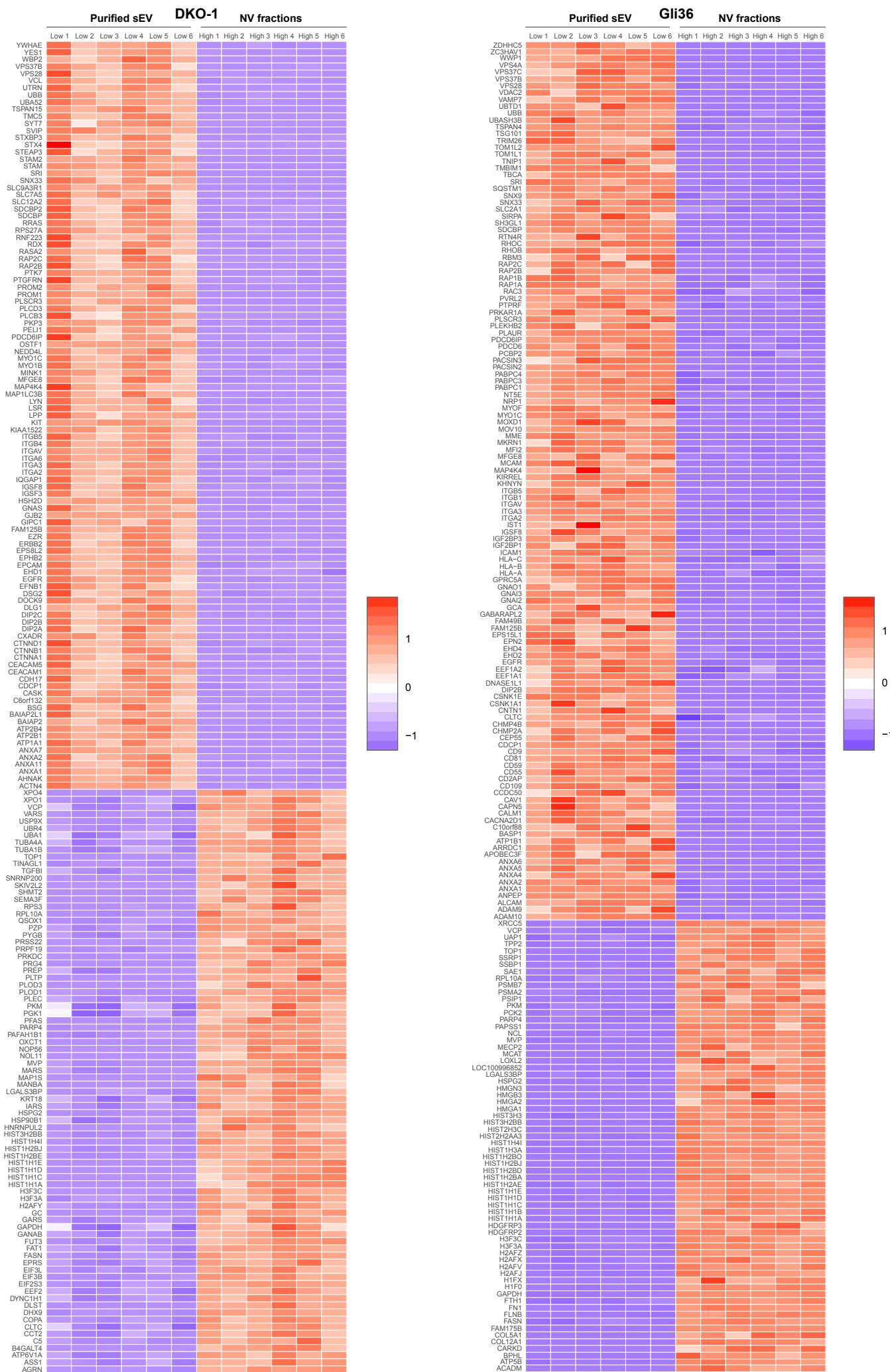


Table S2. Top 200 proteins most significantly differentially expressed between purified sEV and non-vesicular (NV) fractions, Related to Figure 2. Small EVs (low density) and non-vesicular (high density) fractions were separated by high-resolution density gradient centrifugation followed by proteomic analysis. Heat map scale indicates intensity, defined as $(A/\text{spectral counts} - \text{mean spectral counts})/\text{standard deviation}$.

DKO-1			Gli36		
Cell	Large EV	Small EV	Non-vesicular	Small EV	Non-vesicular
GAPDH	AHNAK	PDCD6IP	PKM	C3TC	GAPDH
EEF2	GAPDH	PDCD6IP	CLTC	PDCD6IP	HIST1H4I
HSP90AA1	PKM	AHNAK	PKM	PDCD6IP	Miy
HSP90AB1	ACTG1	SOCBP	PKM	GAPDH	PGK1
KRT18	PKM	SOCBP	DYNCH1	NTSE	CLTC
KRT5	PKM	SOCBP	GAPDH	ANPEP	FTH1
EEF1A1	ANXA2	NTSE	MVP	SOCBP	PKM
FLNA	HSPG2	NTSE	FAT1	SOCBP	HSP90AB1
PKM	FLNA	ITGB4	FAFN	HSP48	PKM
AHNAK	FLNB	ITGB4	PGK1	EEF1A1	HIST1H3A
ACTG1	IQGAP1	DIP2B	HSPG2	UBB	FAFN
ENO1	DYNCH1	PKM	HSP90AA1	HSP9A1	VCP
PLEC	ENO1	GAPDH	EEF2	ENO1	FLNB
PLEC	CLTC	PKM	FLNA	PGK1	DYNCH1
TUBB	EZR	CLTC	MVH	HSP90AB1	FN1
PKM	AGRN	ITGA6	PRKDC	PABPC1	HIST2H3C
PKM	MYH9	PKM	HSP90AB1	EEF1A2	TUBB
HSPD1	MYO1C	ITGA6	UBRA	HLA-A	FLNA
TUBB4B	PGK1	ACTG1	AGRN	HSP9A5	PK2
FASN	ALDOA	PTGRN	VCP	HSP90AA1	HSP90AA1
HSP98	SPTAN1	CTNND1	PLEC	PKM	ATP5B
TUBB4A	HSP90AA1	EZR	PLEC	DYNCH1	PAPR4
TUBB2B	ACTN4	MYO1C	UBA1	ACTG1	H3F3A
CLTC	ITGB4	HSP8	TUBB4B	MYO1C	TUBB4B
MYH9	SPTBN1	IQGAP1	ENO1	EH02	HIST2H2AA3
TUBA1B	ITGB4	DYNCH1	TUBB8	PABPC4	ACTG1
SPTAN1	ACTA1	MYOF	HIST1H4I	FTH1	HSP90B1
FLNB	CTNND1	CD81	TUBA1B	MOV10	EEF2
SPTBN1	SPTBN1	EN01	EPRS	MME	HIST1H2BD
DYNCH1	HSP90AB1	SLC12A2	TGFBI	PKM	UBA1
TUBA1C	MYO1B	ANXA2	KIAA1199	HLA-B	HSP8
SPTBN1	MYO1B	PLXNB2	PRP4	CLTC1	H2AFZ
NCL	ACTG2	PGK1	TUBA1C	ANXA6	H2AFJ
PRKDC	PLEC	ITGB1	ACTG1	ANXA2	PYGL
TUBB2A	PLEC	UBB	TUBB4A	TLN1	TUBB2A
TUBB4A	MYOF	ATP1A1	CLTC1	TUBB	SERpine1
EEF1A2	NTSE	RPS27A	TUBB2A	SQSTM1	TUBB4A
HIST1H4I	HSP8	UBA52	TUBB2B	MYOF	HIST1H2B0
ALDOA	MYO1B	FLNB	PDCD6IP	TUBA1B	VIM
HSP90B1	ITGA6	ITGA6	L GALS3B	MF12	HSP5
VCP	ITGA6	CTNNB1	PDCD6IP	FLNA	NAMPt
XRC5C	PLXNB2	MYO1D	GANAB	EGFR	H2AFY
HNRNPU	YWHAZ	HSP90AA1	TUBA4A	SH3GL1	HIST1H2AE
LRPPRC	NTSE	ITGA3	HSP8	CACNA2D1	TOP1
HSP95	POTE	HSP9A1	GANAB	SLC2A2	H2AFY
HNRNPU	FASN	MFGE8	CCT2	ITGB1	TUBA1B
UBA1	LDHA	ADAM10	HSP90B1	ITGB1	MYH9
CCT2	EEF1A1	CYFIP1	IARS	TUBA1C	ADAM10
ANXA2	EEF2	IGSF8	HIST1H2B	HSP90B1	HIST3H3
ACTN4	ITGB1	CASK	QSOX1	PP1A	ENO1
EPBS	ATP1A1	EPICAM	FLNA	ANXA5	H2AFV
ACTA1	PDCD6IP	C09	LDHA	HSP9A1	FLNB
GANAB	PDCD6IP	CTNNA1	XP01	EEF2	HIST1H2Bj
XRC5C	CTNNA1	ALDOA	HIST1H2Bj	IGSF8	TUBA1C
GANAB	TUBB	SLC2A1	TUBB2A	DIP2B	L GALS3B
LDHA	EPICAM	MYO1B	WDR1	KRT1	GARS
HNRNPA2B1	RDX	FAT1	KRT18	ITGA6	H2AFX
TUBB8	HSP8	MYO1B	PYGB	ICAM1	H3F3C
KRT19	TUBB4B	PDCD6IP	ALDOA	SNX33	TUBB8
UCHL1	VCP	HSP90AB1	GFP1	TUBA4A	XRC5C
HNRNPA1	SLC2A1	JUP	PYGL	HSP9A1	HIST1H1D
LDHA	HIST1H4I	EPSB1	CAND1	PTPRF	HIST1H1C
LMNA	TUBB	ITGB1	COPA	EH01	EPBS
P4HB	CTTNB1	ITGA3	HIST3H2B	TUBB2A	HIST1H1E
CCT4	KRT8	TLN1	KPNB1	PABPC3	TLN1
TUBB6	UBA1	ACTA1	DIPYSL2	FLNB	TUBB4A
RPL3	UTRN	FLNA	AS1	UBA1	TUBB8
NPM1	TLN1	LDHA	HSP8	TUBB4A	XRC5C
RPL4	ACTG2	SLC3A2	TLN1	TPH1	H1F0
HNRNPK	FAM129B	ATP2B1	KRT8	CD3D1	ACTC1
ACTG2	GPI	RDX	PGM	VPS37B	FLNC
LMNA	ACTBL2	NCKAP1	IP-07	HSP82	GANAB
TUBA3E	KRT18	GNA03	RNPEP	FASN	OAT
HIST1H2B	JUP	UTRN	PSMD2	ACTC1	FSCN1
PDIA3	ACTN1	YWHAZ	QCT1	KRT10	ATIC
HSP95	EPSB1	FASN	ACLY	TSG101	HIST1H2AB
YWHAZ	EEF1A1	NAMPt	FSCN1	ACTG2	
EZR	CFL1	KP93	MYH14	EH04	CS
EIF3A	GNA03	CDCP1	SNRP200	VIM	ALDH1A3
TPH1	ITGA6	MYH9	HSP9A1	HLA-C	EEF1A1
TUBB3	GNA02	RAP1B	TUBA8	MSN	MIPEP
POTE	TUBA1B	EGFR	GRY2	C02AP	NDRG1
SNINP200	TUBB2B	POTE	LDHA	HIST1H4I	HY0U1
TLN1	GFP1	NEOD4L	EIF3L	MYH9	BANF1
PGK1	TUBB4A	CD55	VARS	CD81	HSP8
DSPI	POTEJ	ACTN	CCT4	HLA-C	HIST1H1B
SYNCRIP	VCL	MINK1	TUBA3E	SNX9	PDCD6IP
NPM1	ANXA1	ACTG1	PGM1	ADAM10	AARS
CFL1	SFN	TPH1	PRSS23	CFL1	UAP1
CCT8A	LDHA	VCP	GARS	ANXA11	FTL
HSP94	HSP9A1	AP2B1	ERO1L	VCP	PDCD6IP
IP07	TUBB2A	PTK7	CSE1L	VPS28	HIST3H2B
HMG1	POTEF	RAP1A	HUWE1	ALDOA	PXDN
VCL	ANXA5	FAM129B	EEF1A1	CD55	TUBA3E
CALR	MYO6	SDCBP2	SDCBP	PYGL	SSBP1

Table S5. Top 100 most abundant proteins identified in cells, large EV and high-resolution density gradient purified sEV and non-vesicular fractions, Related to Figure 2.

Proteomic analysis of DKO-1 cells, large EVs, small EVs and non-vesicular fractions (left), and Gli36 small EVs and non-vesicular fractions (right). Small EVs and non-vesicular fractions were separated by high-resolution density gradient fractionation. Proteins are ranked from top to bottom based on the number of spectral counts.

Table S6. Differential expression of extracellular miRNA between sEV and non-vesicular (NV) fractions, Related to Figure 2.

DKO-1			Gli36		
Enriched in sEV			Enriched in NV		
miRNA	Fold enrichment	p-value (FDR)	miRNA	Fold enrichment	p-value (FDR)
miR-1249	Infinite	0.005121695	miR-1291	14.504	1.91292E-07
miR-23a-5p	Infinite	0.002511446	miR-4479	10.377	0.011221734
miR-6087	18.512	0.00013301	miR-1307-5p	10.150	8.34514E-20
miR-4516	18.190	0.020737292	miR-22-3p	10.095	1.11159E-38
miR-1260b	12.783	0.001518903	miR-3074-5p	9.794	1.50141E-36
let-7d-5p	12.229	8.24324E-14	miR-24-3p	9.233	6.37581E-35
let-7e-5p	10.010	1.87266E-12	miR-193a-5p	8.566	1.00825E-09
miR-5100	8.929	0.00132335	miR-9-3p	8.494	7.45737E-10
let-7b-5p	8.496	2.01003E-13	miR-629-5p	6.588	7.97809E-25
miR-3178	8.395	0.013110886	miR-129-5p	6.384	3.71236E-05
miR-92a-1-5p	7.330	5.38855E-06	miR-10b-3p	6.174	3.12024E-07
miR-25-5p	7.090	3.34438E-05	let-7i-3p	5.932	0.002671421
miR-30b-3p	6.686	0.002309714	miR-3191-5p	5.372	0.005121695
miR-4508	5.818	0.003926944	miR-101-3p	5.162	6.61305E-18
miR-3182	5.522	9.77007E-05	miR-3190-3p	5.130	0.002671421
miR-492	5.251	0.016848952	miR-335-5p	4.859	6.73765E-16
let-7c	4.999	3.99027E-08	miR-1246	4.649	0.008675648
miR-98-5p	4.883	9.07274E-09	miR-10a-3p	4.447	8.09872E-16
miR-197-3p	4.862	1.91292E-05	miR-378c	4.288	2.48378E-09
let-7a-5p	4.816	7.87557E-07	miR-3605-3p	3.790	0.000219997
miR-1301	4.788	8.35249E-05	miR-629-3p	3.389	0.000826114
miR-139-3p	4.621	0.004441666	miR-320e	3.330	0.003394286
miR-1306-5p	4.584	0.014890867	miR-130b-3p	3.328	0.001518903
miR-1323	4.194	0.031742701	miR-320d	3.315	0.001029466
miR-374b-5p	3.874	0.000281787	miR-95	3.180	7.35126E-09
miR-374c-3p	3.874	0.000281787	miR-378d	3.161	0.000606018
miR-641	3.654	0.001038684	miR-107	3.059	1.45028E-06
miR-185-5p	3.490	6.79962E-06	miR-92b-3p	2.878	0.001816149
miR-516b-5p	3.342	0.005355404	miR-193b-5p	2.758	0.004859571
miR-576-5p	3.300	0.005355404	miR-215	2.574	1.14086E-05
miR-29b-1-5p	3.290	0.012357964	miR-140-3p	2.568	1.77963E-06
miR-424-3p	3.154	0.001816149	miR-181a-5p	2.438	4.95424E-05
miR-331-3p	3.023	0.016116464	miR-328	2.437	6.64054E-05
miR-1247-3p	2.928	0.000171117	miR-92b-3p	2.356	0.011375891
miR-196a-5p	2.862	0.000944707	miR-181b-5p	2.315	7.90088E-05
miR-26a-5p	2.590	0.000326817	miR-29a-3p	2.280	0.00015546
miR-1180	2.522	0.00132335	miR-3184-5p	2.280	1.4716E-05
miR-125a-3p	2.507	0.018563296	miR-152	2.269	3.71236E-05
miR-196b-5p	2.353	0.00211413	miR-1287	2.261	0.0443721
miR-652-3p	2.206	0.031366541	miR-192-5p	2.201	0.000432721
miR-132-3p	2.091	0.006074342	miR-3615	2.182	0.000321644
miR-320a	1.975	0.007662384	miR-423-3p	2.149	6.64054E-05
let-7i-5p	1.970	0.039864818	miR-203b-3p	2.139	0.00115975
miR-27a-5p	1.819	0.039609269	miR-375	2.106	8.68019E-05
Enriched in NV			miR-500a-3p	2.024	0.020431581
Enriched in sEV			miR-320c	1.934	0.01540494
Enriched in NV			miR-3928	1.930	0.033329499
Enriched in sEV			miR-378a-3p	1.928	0.002243453
Enriched in NV			miR-1304-3p	1.909	0.033329499
Enriched in sEV			miR-651	1.890	0.02257969
Enriched in NV			miR-103a-3p	1.878	0.001152357
Enriched in sEV			miR-103b	1.878	0.001152357
Enriched in NV			miR-484	1.862	0.007635951
Enriched in sEV			miR-20a-5p	1.829	0.004196148
Enriched in NV			miR-584-5p	1.811	0.011512276
Enriched in sEV			miR-92a-3p	1.811	0.005121695
Enriched in NV			miR-210	1.771	0.019523623
Enriched in sEV			miR-146a-5p	1.762	0.015843339
Enriched in NV			miR-221-3p	1.720	0.0443721
Enriched in sEV			miR-28-3p	1.715	0.011190766
miRNA	Fold enrichment	p-value (FDR)	miRNA	Fold enrichment	p-value (FDR)
miR-1236-5p	Infinite	2.38577E-12	miR-4661-5p	Infinite	0.027843937
miR-1225-5p	Infinite	1.25306E-05	miR-215	59.698	1.87176E-06
miR-3676-5p	Infinite	0.001888493	miR-452-5p	53.656	0.000115088
miR-3620-3p	Infinite	0.003500437	miR-335-5p	35.639	0.000886052
miR-3940-3p	Infinite	0.006605787	miR-335-3p	35.545	4.04217E-05
miR-4700-3p	Infinite	0.006756923	miR-148a-3p	35.514	1.10994E-36
miR-574-5p	Infinite	0.007004365	miR-200a-5p	24.988	6.4363E-05
miR-6511b-5p	Infinite	0.023992038	miR-200b-3p	23.785	2.06953E-50
miR-4443	190.216	8.83084E-11	miR-200c-3p	19.975	4.58537E-52
miR-1275	106.311	1.61415E-29	miR-375	19.483	1.59982E-50
miR-1229-5p	70.218	0.002860958	miR-1291	17.893	0.001072026
miR-324-5p	49.582	0.000375731	miR-22-3p	17.802	6.88043E-79
miR-23a-5p	42.661	3.59764E-39	miR-10b-5p	16.450	1.19132E-49
miR-365a-5p	32.555	1.95513E-57	miR-3074-5p	15.061	6.15373E-71
miR-378a-5p	31.077	2.51746E-14	miR-224-5p	15.030	1.42675E-17
miR-197-5p	29.640	3.07266E-29	miR-184	14.606	1.40258E-11
miR-877-3p	24.591	0.01126995	miR-200b-5p	14.231	5.82675E-05
miR-671-5p	22.256	2.72141E-19	miR-193a-5p	13.711	5.6144E-66
miR-34a-5p	20.585	0.00213908	miR-320d	12.881	0.000199904
miR-149-3p	18.990	2.71965E-20	miR-24-3p	11.854	1.84227E-62
miR-365b-5p	17.913	2.93753E-38	miR-3909	11.754	0.000684607
miR-30b-3p	17.480	1.42828E-22	miR-1304-3p	10.863	0.000248823
miR-769-3p	16.750	0.013097097	miR-200a-3p	10.715	3.19293E-40
miR-30c-1-3p	16.153	0.010617826	miR-1307-5p	9.487	7.44722E-19
miR-1296	15.663	0.022946487	miR-629-5p	9.358	9.49043E-17
miR-5100	14.119	2.55966E-14	miR-378i	9.172	1.3915E-12
miR-92b-5p	13.471	1.85341E-23	miR-625-5p	9.001	0.004073497
miR-664b-5p	13.311	0.038311784	miR-429	8.703	1.36806E-06
miR-1226-5p	13.283	0.019381796	miR-141-3p	7.625	1.55299E-07
miR-2277-5p	12.573	2.48366E-10	miR-3605-3p	7.587	1.52876E-11
miR-331-3p	11.439	0.000272393	miR-92b-3p	7.541	5.23634E-30
miR-29b-1-5p	11.037	3.22655E-12	miR-378c	7.331	1.81494E-12
miR-1908	10.123	9.64036E-18	miR-192-5p	7.007	1.86856E-09
miR-1236-3p	9.068	5.19374E-13	miR-625-3p	6.177	0.000203018
miR-3182	8.536	0.00059701	miR-320c	6.119	5.63863E-06
miR-503-3p	8.476	9.53712E-05	miR-27b-5p	5.868	0.000101367
miR-652-3p	8.244	6.82654E-06	miR-378a-3p	5.857	5.85365E-13
miR-532-3p	7.980	0.00336107	miR-140-3p	5.526	2.88794E-26
miR-193b-5p	7.439	4.76071E-20	miR-339-3p	5.513	1.67232E-23
let-7b-5p	7.327	1.21871E-26	miR-3158-5p	5.352	0.008418044
miR-140-5p	7.275	2.44209E-22	miR-152	5.276	4.44765E-14
let-7e-5p	6.944	1.56845E-20	miR-10a-5p	5.275	1.14245E-29
miR-195-5p	6.768	3.37288E-05	miR-10a-3p	5.239	2.03899E-10
miR-935	6.483	0.0025355	miR-320b	5.091	4.3029E-09
miR-342-5p	6.407	0.037863457	miR-363-3p	4.756	6.57425E-06
miR-93-3p	6.305	0.005811746	let-7d-3p	4.679	1.82846E-10
miR-193b-3p	6.203	2.85747E-09	miR-378d	4.111	0.001877832
miR-25-5p	5.731	5.63155E-06	let-7f-5p	3.881	1.06414E-19
miR-1260b	5.501	0.03745708	miR-128	3.770	2.91475E-13
miR-151a-5p	5.246	5.25486E-06	miR-9-3p	3.691	8.12733E-09
miR-744-5p	5.149	1.05909E-14	miR-3615	3.625	2.93304E-06
miR-138-5p	5.099	0.000692993	miR-708-3p	3.503	0.023523389
miR-339-5p	4.965	0.000272393	miR-125a-5p	3.470	2.44266E-17
miR-374b-5p	4.764	9.76307E-10	miR-181b-5p	3.431	9.09106E-09
miR-374c-3p	4.764	9.76307E-10	miR-130b-3p	3.405	0.000225281
miR-874	4.553	0.001167393	miR-3158-3p	3.232	0.039211927
miR-444-3p	4.436	0.000802099	miR-1468	3.221	0.009693567
miR-23b-5p	4.390	0.004068123	miR-130b-5p	3.090	4.8246E-07
miR-16-5p	4.241	1.63666E-11	miR-30e-3p	3.070	5.18956E-11
miR-99b-3p	4.053	1.65409E-07	miR-30a-3p	3.041	3.98371E-12
miR-4521	3.938	0.043802516	miR-22-5p	2.887	1.75393E-06
miR-92a-1-5p	3.916	2.55567E-07	miR-27b-3p	2.843	9.09835E-08
miR-125a-3p	3.884	0.002746171	miR-584-5p	2.609	0.009895457
miR-126-5p	3.870	0.009895457	miR-107	2.508	0.00011606
miR-769-5p	3.809	1.99282E-07	miR-501-3p	2.500	0.022367484
miR-365b-3p	3.724	0.048480747	miR-345-5p	2.479	0.003479991
miR-365a-3p	3.724	0.048480747	miR-130a-3p	2.427	3.86817E-05
miR-374a-5p	3.455	4.26559E-08	miR-95	2.412	0.000120622
let-7d-5p	3.245	1.31339E-07	miR-103a-3p	2.410	4.37759E-09
miR-425-3p	3.223	0.006307783	miR-103b	2.410	4.37759E-09
miR-455-3p	3.104	0.001975098	miR-2682-5p	2.40	

Table S7. Differential expression of miRNA between sEVs and parental cells, Related to Figure 2.

DKO-1			Gli36		
Enriched in sEV			Enriched in Cell		
miRNA	Fold enrichment	p-value (FDR)	miRNA	Fold enrichment	p-value (FDR)
miR-3655	Infinite	4.70978E-13	miR-1275	Infinite	9.81446E-09
miR-105-5p	Infinite	2.04769E-08	miR-3662	Infinite	1.24759E-08
miR-4680-3p	Infinite	1.31609E-06	miR-573	Infinite	0.00065315
miR-4492	Infinite	8.60079E-06	miR-652-5p	Infinite	0.00318554
miR-708-3p	Infinite	0.000157405	miR-4791	Infinite	0.00352673
miR-767-5p	Infinite	0.000161031	miR-3170	Infinite	0.005831891
miR-223-5p	Infinite	0.019209336	miR-4517	Infinite	0.009013818
miR-4488	1804.609	1.49977E-10	miR-1257	Infinite	0.013635142
miR-3178	422.181	5.09375E-35	miR-4521	27.560	3.08294E-09
miR-552	283.755	1.25171E-95	miR-1305	26.512	0.002824
miR-6087	262.881	8.51081E-05	miR-3145-3p	25.235	0.001106078
miR-4516	255.845	8.66747E-21	miR-556-3p	24.370	2.02458E-05
miR-492	206.371	3.782E-53	miR-3116	22.543	0.007677828
miR-3656	145.064	1.21256E-11	miR-19b-1-5p	22.436	0.009511301
miR-1244	111.079	1.26063E-08	miR-2277-5p	21.970	4.12297E-28
miR-4508	91.519	3.27675E-54	miR-5701	18.417	0.000203264
miR-4680-5p	74.011	7.12838E-06	miR-3944-3p	16.542	0.042765388
miR-4709-3p	72.361	1.85865E-05	miR-3619-5p	14.771	1.21377E-07
miR-622	44.538	1.61814E-10	miR-188-5p	12.827	2.78424E-06
miR-205-5p	41.887	0.000278353	miR-4802-3p	11.573	7.58056E-12
miR-4700-5p	39.351	4.05856E-08	miR-4687-3p	11.379	0.011941098
miR-320e	35.159	3.37923E-81	miR-181b-3p	11.272	1.5175E-24
miR-320d	33.232	0	miR-671-5p	11.257	1.01949E-41
miR-1246	32.538	2.0532E-153	miR-18a-5p	10.597	1.62546E-43
miR-139-3p	31.587	4.67079E-48	miR-4470	9.766	0.012175725
miR-139-5p	31.327	0	miR-1296	9.129	4.75862E-19
miR-4497	28.123	3.00828E-17	miR-582-5p	8.571	8.67023E-06
miR-125a-5p	27.806	0	miR-7-5p	8.195	1.96983E-62
miR-95	24.348	2.67207E-67	miR-3529-3p	8.184	7.71032E-62
miR-4647	24.301	0.000355041	miR-1914-5p	8.135	0.005130632
miR-543	23.631	0.02386302	miR-16-1-3p	7.533	2.57177E-06
miR-3929	23.380	0.00020019	miR-548u	6.735	0.013705458
miR-4709-5p	21.275	0.043838734	miR-449c-5p	6.662	9.81144E-06
miR-3617-5p	20.627	9.37596E-08	miR-185-3p	6.283	3.55976E-15
miR-135b-5p	15.809	3.2324E-164	miR-491-5p	6.150	2.69102E-09
miR-125a-3p	15.024	1.1762E-84	miR-15a-5p	6.146	1.76282E-13
miR-127-3p	14.979	0.042644537	miR-500a-5p	6.080	0.011770198
miR-509-3p	13.908	0.03328933	miR-33b-3p	5.952	0.000199165
miR-153	13.583	0.000149454	miR-4659a-3p	5.763	0.003599157
miR-4724-5p	12.996	0.000101085	miR-4446-3p	5.688	0.02358975
miR-4510	11.846	5.60716E-05	miR-5001-5p	5.633	4.96868E-07
miR-3605-3p	11.724	3.13392E-37	miR-301a-3p	5.423	0.000822508
miR-320c	10.690	1.65482E-05	miR-17-5p	5.074	2.88082E-32
miR-1468	10.514	2.63989E-21	miR-3200-3p	4.926	1.12411E-10
miR-3614-5p	9.232	0.003394943	miR-362-5p	4.774	1.34627E-12
miR-664a-5p	9.177	9.17537E-46	miR-326	4.736	0.00095297
miR-99b-5p	8.674	0	miR-3187-3p	4.647	1.4951E-05
miR-181a-5p	8.266	2.2839E-200	miR-107	4.489	4.52082E-27
let-7e-5p	8.159	9.0465E-187	miR-15b-5p	4.470	6.92977E-33
miR-516a-5p	8.126	3.37996E-05	miR-542-3p	4.359	8.53609E-16
miR-141-3p	7.820	0	miR-877-3p	4.323	9.89344E-08
miR-3182	7.612	3.10853E-24	miR-548n	4.322	0.040189372
miR-3605-5p	7.433	0.000232411	miR-500a-3p	4.290	1.53307E-21
miR-224-5p	7.397	2.9691E-174	miR-373-3p	4.206	1.21078E-08
miR-338-5p	7.351	3.97394E-16	miR-4474-3p	4.181	0.006004349
miR-1277-3p	7.244	0.013084336	miR-106b-5p	4.045	2.15194E-32
miR-10b-5p	7.239	1.9747E-151	miR-4664-3p	4.034	7.78091E-11
miR-375	7.233	0	miR-301a-5p	4.017	4.13582E-18
let-7c	6.781	7.2929E-194			
miR-934	6.757	3.37263E-07			
miR-99a-5p	6.613	2.77785E-17			
miR-146b-5p	6.520	1.3862E-131			
let-7b-5p	6.493	1.1294E-191			
miR-523-3p	6.422	0.033308577			
let-7a-5p	6.363	2.0037E-161			
miR-27a-3p	6.363	5.6018E-189			
miR-184	6.146	0.001677301			
miR-23a-3p	5.955	7.832E-168			
miR-93-5p	5.921	0.029833036			
miR-616-3p	5.701	0.019209336			
miR-320b	5.689	1.49E-139			
miR-451a	5.629	1.92422E-06			
miR-203a	5.608	8.9862E-162			
miR-203b-5p	5.596	2.4618E-161			
miR-142-5p	5.534	6.4154E-15			
miR-4708-5p	5.501	0.020320854			
miR-129-5p	5.276	0.00193896			
miR-642a-3p	5.197	2.57271E-08			
miR-642b-5p	5.197	2.57271E-08			
miR-664a-3p	4.966	5.81721E-13			
miR-483-5p	4.924	0.003890983			
miR-514a-3p	4.651	6.64653E-05			
miR-3074-3p	4.570	7.72471E-22			
miR-6716-3p	4.565	0.00016912			
miR-100-5p	4.487	9.55594E-73			
miR-328	4.432	2.09627E-16			
miR-3065-3p	4.401	8.51598E-12			
miR-222-3p	4.365	1.86175E-76			
miR-143-3p	4.305	1.02415E-08			
miR-210	4.056	3.47579E-46			

Table S7. Differential expression of miRNA between sEVs and parental cells, Related to Figure 2.

Significant differential expression of miRNAs between density gradient purified sEVs and their parental cell type. Included are miRNAs with at least a four fold-change enrichment and an FDR-adjusted p-value < 0.05.