

Table S1: Dataset characteristics of whole embryo lysate sample set data (raw Ct values) and conditioned media samples (base 2 logarithmic transformations) according to the BestKeeper analytical tool. *Abbr.* CP – crossing point (cycle threshold)

<i>CP data of housekeeping Genes (Whole Embryo Lysates):</i>							
	U6	let-7a	miR-16	miR-26a	miR-103	miR-106	miR-191
n	7	7	7	7	7	7	7
geo Mean [CP]	28.58	34.37	30.50	34.92	32.36	33.01	33.45
ar Mean [CP]	28.76	34.39	30.56	34.96	32.49	33.10	33.53
min [CP]	25.39	32.20	27.41	32.22	28.69	30.05	30.30
max [CP]	35.01	35.92	33.45	36.61	36.58	37.26	36.47
std dev [± CP]	2.89	1.08	1.64	1.43	2.58	2.07	2.17
CV [% CP]	10.04	3.14	5.37	4.10	7.94	6.26	6.46
min [x-fold]	-9.12	-4.50	-8.54	-6.49	-12.74	-7.78	-8.85
max [x-fold]	86.32	2.93	7.72	3.21	18.64	18.99	8.11
std dev [± x-fold]	7.40	2.11	3.12	2.70	5.97	4.21	4.49

<i>CP data of housekeeping Genes (Conditioned Media):</i>				
	U6	miR-16	miR-103	miR-106
n	6	6	6	6
geo Mean [CP]	5.50	5.31	3.92	2.91
ar Mean [CP]	5.57	5.44	4.24	3.16
min [CP]	4.05	4.06	2.62	1.96
max [CP]	6.41	7.50	7.17	5.07
std dev [± CP]	0.75	1.05	1.60	1.22
CV [% CP]	13.55	19.22	37.67	38.53
min [x-fold]	-2.73	-2.38	-2.47	-1.93
max [x-fold]	1.88	4.57	9.56	4.47
std dev [± x-fold]	1.69	2.06	3.02	2.33