

Supplementary Table 2. Top ten miRNAs in undifferentiated EC cells and their associations with malignancy.

The relative expression of top up- and downregulated miRNAs in undifferentiated 2102Ep cells compared to Ntera2 cells. Comparative analysis was carried out using the 2^{-ddC_t} method (Livak and Schmittgen, 2001). Expression levels for miRNAs in undifferentiated Ntera2 cells were equated to 1 and levels in undifferentiated 2102Ep cells are presented proportionally to this.

miRNA	Relative Expression	Malignancy	miRNA	Relative Expression	Malignancy
Upregulated			Downregulated		
miR-9	1055.7	Lung [66] Breast [49]	miR-296	-53.2	Oral [SR8]
miR-32	1324.0	Lung [66] Prostate [56]	miR-211	-18.1	Lung [66]
miR-9*	1419.0	Breast [45] Breast [SR1]	miR-139	-7.4	Tongue [65] Neural [SR9]
miR-127	1754.9	Leukaemia [SR2] Tongue [66]	miR-181b	-4.89	Colorectal [SR10] Ovary [33]
miR-134	1999.4		miR-199a	-4.1	Lung [66]
miR-520a*	2006.4		miR-363	-3.5	Lymphoma [80]
miR-365	2130.7	Breast [SR1] Leukaemia [51]	miR-302C*	-2.9	Embryonal
miR-373	2378.6	Prostate [SR3] Metastasis [SR4]	miR-302b	-2.7	Carcinoma [SR11] Lymphoma [52]
miR-148a	2399.0	Leukaemia [SR5]	miR-19a	-2.6	Lung [53] Pancreas [SR12]
miR-371	2835.0	Prostate [65]	miR-190	-2.5	Liver [SR13] Leukaemia [55]
miR-182	3000.4	Melanoma [SR6]	miR-17-5p	-2.4	Colorectal [57]
miR-519c	3571.2		miR-25	-2.3	Gastric [60]

miR-335	4500.4	Ovary [58] Myeloma [74]	miR-100	-2.3	Ovary [37] Prostate [SR14]
miR-518c	6371.7	Tumorigenesis [63]	let-7e	-2.3	Ovary [61] Tumorigenesis [63]
miR-17-3p	6675.6	Lymphoma [52] Colorectal [62]	miR-214	-2.2	Ovary [38]
miR-372	8647.5	Germ Cell [SR7] Tongue [91]	let-7i	-2.2	Ovary [39]
miR-154*	15922.9	Leukaemia [SR2] Tongue [66]	miR-212	-2.1	
			miR-29a	-2.1	Metastasis [SR15] Leukaemia [SR15]

Supplementary Table 3 References

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