

Table S6. Candidate tumor-suppressive miRNAs binding to *MCM6*.

MicroRNA	miRBase accession No.	Log ₂ fold change GSE230229	Normalized read count GSE230229		FDR GSE230229	<i>p</i> value GSE230229
			LUAD tissues	Normal lung tissues		
<i>hsa-miR-551b-5p</i>	MIMAT0004794	-3.39	1.03	4.42	0.022	0.004
<i>hsa-miR-218-1-3p</i>	MIMAT0004565	-3.07	1.05	4.12	0.061	0.012
<i>hsa-miR-6813-3p</i>	MIMAT0027527	-2.65	0.00	2.65	0.007	0.001
<i>hsa-miR-6768-5p</i>	MIMAT0027436	-2.13	4.12	6.25	0.076	0.016
<i>hsa-miR-373-3p</i>	MIMAT0000726	-2.09	0.00	2.09	0.152	0.038
<i>hsa-miR-144-3p</i>	MIMAT0000436	-2.06	10.85	12.91	0.107	0.024
<i>hsa-miR-30c-2-3p</i>	MIMAT0004550	-1.92	8.27	10.19	0.119	0.027
<i>hsa-miR-1-5p</i>	MIMAT0031892	-1.62	3.76	5.37	0.047	0.009
<i>hsa-miR-145-3p</i>	MIMAT0004601	-1.24	11.26	12.50	0.157	0.039
<i>hsa-miR-548ao-5p</i>	MIMAT0021029	-1.12	4.55	5.67	0.003	<0.001
<i>hsa-miR-484</i>	MIMAT0002174	-1.04	7.73	8.77	0.055	0.011
<i>hsa-miR-2110</i>	MIMAT0010133	-0.69	6.91	7.59	0.141	0.034

LUAD: lung adenocarcinoma

FDR: false discovery rate