

**Table S3. Candidate tumor-suppressive miRNAs binding to *MCM3*.**

MicroRNA	miRBase accession No.	Log <sub>2</sub> fold change GSE230229	Normalized read count GSE230229		FDR GSE230229	<i>p</i> value GSE230229
			LUAD tissues	Normal lung tissues		
<i>hsa-miR-1208</i>	MIMAT0005873	-2.95	0.00	2.95	0.002	<0.001
<i>hsa-miR-516b-5p</i>	MIMAT0002859	-2.50	1.12	3.62	0.100	0.022
<i>hsa-miR-373-3p</i>	MIMAT0000726	-2.09	0.00	2.09	0.152	0.038
<i>hsa-miR-4516</i>	MIMAT0019053	-1.38	3.56	4.94	0.164	0.042
<i>hsa-miR-3613-3p</i>	MIMAT0017991	-1.25	5.14	6.40	0.016	0.003

LUAD: lung adenocarcinoma

FDR: false discovery rate