



Description	Gene Name	Accession	FC (peptide-to-spectrum matches [PSM] values)		p-Value
			Scrambled	miR-101a-3p	
Proline-, glutamic acid- and leucine-rich protein 1	PELP1	Q8IZL8	1	4	0.0506
AT-rich interactive domain-containing protein 3A	ARID3a	Q99856	1	3	0.0454
WD repeat-containing protein 43	WDR43	Q15061	1	3	0.1870
Nuclear pore membrane glycoprotein 210	NUP210	Q8TEM1	1	2.285	0.0370
Elongation factor 1-alpha 1	EEF1A1	P68104	1	1.674	0.0428

Supplementary Figure S12. The number of proteins detected in global proteomics analyses pulled down in miR-101a-3p promoter transfected cells compared to scrambled transfected cells (a) and their abundance in fold change in a heatmap (b). c) The selected top 5 proteins that are abundantly binding to miR-101a-3p promoter.