

Table S2

**List of down-regulated probes/genes identified by cDNA  
microarray comparing CLL with high and low miR-155  
expressions**

<b>GeneBank ID</b>	<b>Gene name</b>	<b>Description</b>	<b>Fold Change</b>	<b>q-value(%)</b>
AL521634	CoQ10A	coenzyme Q10 homolog A	0.71	0
<b>AL530441</b>	<b>CSNK1G2</b>	<b>casein kinase 1 gamma 2</b>	<b>0.79</b>	<b>0</b>
<b>NM_001319</b>	<b>CSNK1G2</b>	<b>casein kinase 1 gamma 2</b>	<b>0.77</b>	<b>0</b>
AI989996	KDM2B	lysine demethylase 2B	0.68	0
N32834	GLIPR1	GLI pathogenesis-related 1	0.45	3.16
<b>AA130132</b>	<b>KLF3</b>	<b>Kruppel-like facotr 3 (basic)</b>	<b>0.11</b>	<b>3.16</b>
BC006270	HECTD4	HECT domain E3 ubiquitin protein ligase 4	0.72	3.16
<b>NM_014897</b>	<b>ZNF652</b>	<b>Homo sapiens zinc finger protein 652</b>	<b>0.61</b>	<b>3.16</b>
AF289565	KLF3-AS1	KLF3 antisense RNA 1	0.74	3.16
<b>BF438116</b>	<b>KLF3</b>	<b>Kruppel-like facotr 3 (basic)</b>	<b>0.17</b>	<b>5.23</b>

Genes with predicted miR-155 binding sites are in bold.

FDR=false discovery rate, <0.05