

<b>miRNAs</b>	<b>No. of target genes</b>	<b>No. of target genes identified by at least 5 out of 8 prediction tools*</b>
hsa-miR-24-3p	641	139
hsa-miR-130a-3p	838	338
hsa-miR-505-3p	273	3
hsa-miR-142-3p	336	118
hsa-miR-455-5p	182	21
<b>Total</b>	<b>2270</b>	<b>619</b>

\* Prediction tools include VALIDATION, DIANA, MIRANDA, MIRBRIDGE, PICTAR, PITA, RNA22, and TARGETSCAN

<http://mirsystem.cgm.ntu.edu.tw/index.php>