Figure S1

<table>
<thead>
<tr>
<th></th>
<th>miR-126*</th>
<th>miR-32</th>
<th>miR-101</th>
<th>miR-15b</th>
<th>miR-550</th>
</tr>
</thead>
<tbody>
<tr>
<td>Microarray</td>
<td>Fold-change</td>
<td>-3.02</td>
<td>-2.80</td>
<td>-2.64</td>
<td>1.26</td>
</tr>
<tr>
<td></td>
<td>B-statistic</td>
<td>2.68</td>
<td>0.63</td>
<td>0.56</td>
<td>-3.17</td>
</tr>
<tr>
<td>qRT-PCR</td>
<td>Fold-change</td>
<td>-3.35</td>
<td>-2.96</td>
<td>-2.35</td>
<td>1.58</td>
</tr>
<tr>
<td></td>
<td>p-value</td>
<td>2.00E-03</td>
<td>9.00E-03</td>
<td>1.63E-02</td>
<td>1.58E-02</td>
</tr>
</tbody>
</table>