

**Table 2. Pseudogenes aberrantly expressed in human cancer and validated miRNAs binding to their cognate wt genes.**

Pseudogene	Role in human cancer	wt gene	Validated miRNA families	Conservation of the binding site between wt and pseudo
<i>PTENP1</i>	see text	<i>PTEN</i> : see text	<i>miR-17</i> <i>miR-19</i> <i>miR-21</i> <i>miR-26</i> <i>miR-214</i> <i>miR-216</i> <i>miR-217</i>	yes yes yes yes yes no no
$\psi$ <i>CX43</i>	specifically expressed in breast cell lines (not in normal mammary epithelium) <sup>1</sup>	<i>CONNEXIN 43 (CX43)</i> : one of the monomers that compose gap junctions. CX43 expression is aberrantly lost in cancer.	<i>miR-1</i> <sup>2</sup>	yes
<i>NA88-A</i>	specifically expressed in melanoma cell lines (not in normal melanocytes) <sup>3</sup>	<i>HPX42B</i>	-	
<i>OCT4-pg1</i> <i>OCT4-pg5</i>	specifically expressed in cancer cell lines and tissues (not in normal tissues) <sup>4</sup>	<i>OCT4</i> : transcription factor expressed in embryonic stem cells where it plays a critical role in maintaining the pluripotent and self-renewing state. Oct4 is aberrantly expressed in cancer cells.	<i>miR-145</i> <sup>5</sup>  <i>miR-470</i> <sup>6</sup>	yes  <i>miR-470</i> is mouse-spec
<i>NANOGP8</i>	specifically expressed in cancer cell lines and tissues (not in normal fibroblasts and fetal liver) <sup>7</sup>	<i>NANOG</i> : transcription factor expressed in embryonic stem cells where it plays a critical role in maintaining the pluripotent and self-renewing state. Oct4 is aberrantly expressed in cancer cells	<i>miR-134</i> <sup>8</sup>  <i>miR-296</i> <sup>6</sup>	no*  <i>miR-296</i> binding sites are not conserved between human and mouse
$\psi$ <i>BRAF</i>	specifically expressed in thyroid tumor samples (especially if they don't carry BRAF mutations), and not in normal thyroid <sup>9</sup>	<i>BRAF</i> : Ser/Thr kinase that serves as downstream effector of RAS in the MAPK signaling cascade. Mutations that render BRAF constitutively active are common in cancer.	-	-

The conservation of *miR-17*, *19*, *21*, *26* and *214* binding sites in *PTENP1* has been discussed elsewhere (**Fig. 1**). The asterisk indicates those wt/pseudogene pairs that show an overall low sequence conservation (<60%).

**References 1-9**

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