

Table 3. Conservation of validated miRNA binding sites in cancer-related target genes.

wt genes	corresponding pseudogene(s)	validated miRNA families	conservation of the binding site between wt and pseudo
<i>CCND3</i>	<i>CCND3P</i>	<i>miR-16</i> ¹	no*
<i>CDK4</i>	<i>CDK4P</i>	<i>miR-34</i> ²	yes
<i>DNMT3A</i>	<i>DNMT3AP1</i>	<i>miR-29</i> ³ <i>miR-143</i> ⁴	no no
<i>E2F3</i>	<i>E2F3P1</i>	<i>miR-17</i> ⁵ <i>miR-34</i> ⁶	yes no
<i>c-MYC</i>	<i>MYCL3</i>	<i>let-7</i> ⁷ <i>miR-145</i> ⁸	no* no*
<i>OCT4</i>	<i>OCT4-pg1,2,3,4,5,6</i>	<i>miR-145</i> ⁹	yes
<i>KRAS</i>	<i>KRAS1P</i>	<i>let-7</i> ¹⁰ <i>miR-143</i> ¹¹	yes yes
<i>PTEN</i>	<i>PTENP1</i>	<i>miR-17</i> ¹² <i>miR-19</i> ^{13,14} <i>miR-21</i> ¹⁵ <i>miR-26</i> ¹⁶ <i>miR-214</i> ¹⁷ <i>miR-216</i> ¹⁸ <i>miR-217</i> ¹⁸	yes yes yes yes yes no no
<i>FOXO3</i>	<i>FOXO3B</i>	<i>miR-182</i> ¹⁹	yes

A list of miRNA families with a well recognized oncogenic or oncosuppressor role was obtained merging the most recent reviews about microRNAs and cancer²⁰⁻²⁴.

The validated targets of these miRNAs that have at least 1 pseudogene (<http://www.genecards.org>) are listed above. The conservation of the binding sites of the validated miRNAs in the pseudogene(s) is also reported. The asterisk indicates those wt/pseudogene pairs that show an overall low sequence conservation (<60%).

The conservation of *miR-17*, *19*, *21*, *26* and *214* binding sites in *PTENP1* has been discussed elsewhere (Fig. 1). Analogously, the conservation of *miR-145* binding sites in *OCT4* pseudogenes has been described in Table 2.

References 1-24

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