

miR-34a is not a tumor suppressor

Hervé Seitz

IGH (UMR 9002 CNRS and university of Montpellier), Montpellier, France

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Slideshow accessible for download at:

<http://www.igh.cnrs.fr/equip/Seitz/CRCT21.pdf>

Defining "gene
regulation"

microRNAs

miR-34 in cancer

miR-34a is not a
general repressor
of proliferation

miR-34a is rarely
mutated in cancers

Acknowledgements

Supplementary
data

Defining “gene regulation”

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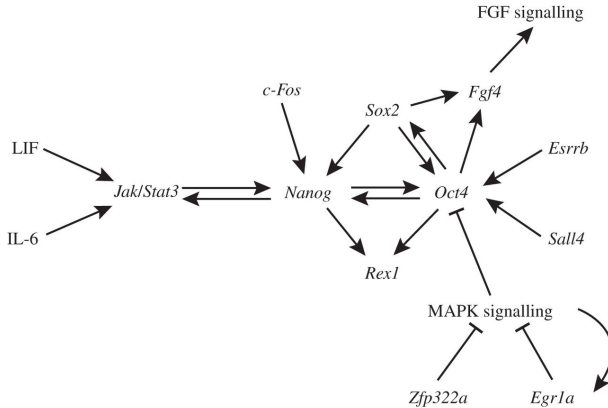
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(Parfitt and Shen, 2014)

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(illustration by D. Goodsell, Scripps Research Institute and Rutgers University)

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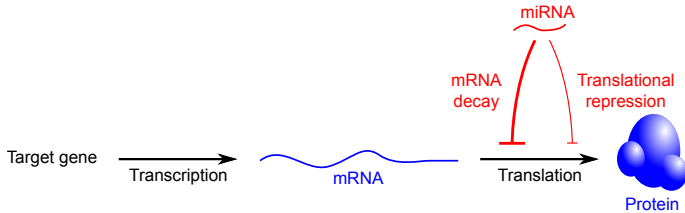
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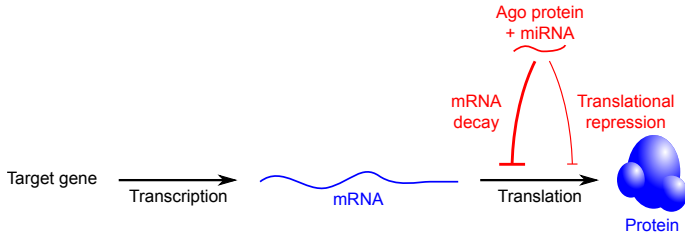
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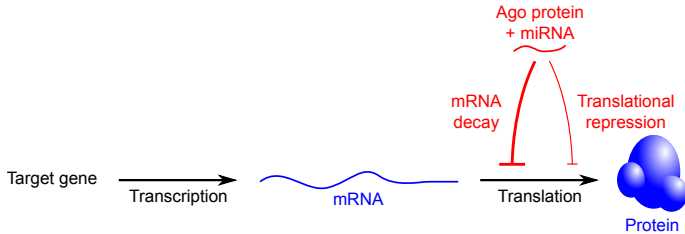
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miRNA sequences are known (for the most part); can we identify their targets ?

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- ▶ Physically bind the miRNA effector complex.
 - ▶ CLIP protocol
- ▶ mRNA and protein level respond to miRNA.
- ▶ Sequence complementarity to the miRNA.

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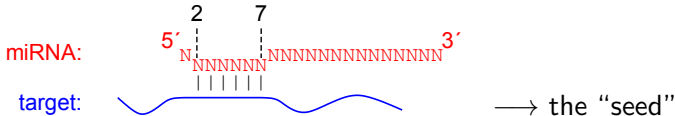
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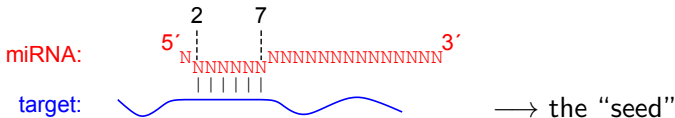
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microRNAs

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Comparative genomics (if a seed match is conserved, it must be functional).

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Functionality of miRNA/target interactions

- ▶ Experimental methods (CLIP and avatars): hundreds of targets per miRNA.
- ▶ Computational methods (seed matches, conserved in evolution): hundreds of targets per miRNA.

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miRNA-mediated repression is very modest (usually < 2 -fold): lower than well tolerated fluctuations in gene expression (e.g., haplosufficiency).

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Both experimental and computational target prediction methods capture mostly false positives (Pinzón *et al.*, 2017).

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Murine miR-34 family members:

miR-34b-5p AGGCAGUGU--CAUUGAGCUGAUUG-
miR-34c-5p AGGCAGUGU--AGUUGAGCUGAUUGC
miR-34a-5p UGGCAGUGU--C-UUAGCUGGUUGU
miR-449a UGGCAGUGUAUUGUUGAGCUGG---
miR-449b-5p AGGCAGUGUAUUGUAGCUGGC---
miR-449c-5p AGGCAGUGUAUUGCUAGC-GGCGUGU

Seed

miR-34a-5p UGGCAGUG--UC-UUAGCUGGUUGU
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miR-34c-5p AGGCAGUG--UAGUUAGCUGAUUGC
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2007: miR-34a, miR-34b and miR-34c are transcriptionally controlled by p53. ► 7 simultaneous publications

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miR-34 miRNAs may participate in the anti-proliferative
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miR-34a: the most highly expressed member in human cell
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→ *miR-34a*, first described tumor-suppressing miRNA.

miR-34 in cancer

Problems:

- ▶ In mouse, a triple KO (miR-34a, b and c) and a sextuple KO (miR-34a, b and c and miR-449a, b and c) do not have increased cancer frequency ([Concepcion et al., 2012](#); [Song et al., 2014](#)).

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miR-34b (0.06 ppm)
miR-449b (2 ppm)
miR-449c (3 ppm)
miR-34c (5 ppm)
miR-449a (36 ppm)
miR-34a (246 ppm)

HCT-116:



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- ▶ miR-34 miRNAs are poorly expressed in most cells.

▶ Additional cell lines, tissues and body fluids

Poorly-expressed miRNAs fail to repress their targets, while being easily detectable ([Mullokandov et al., 2012](#)).

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- ▶ Supra-physiological miRNA over-expression is prone to false positives (reviewed in [Mockly and Seitz, 2019](#)).
- ▶ miR-34 miRNAs are poorly expressed in most cells.
- ▶ MRX34 clinical trial failed to improve cancer patients' condition.

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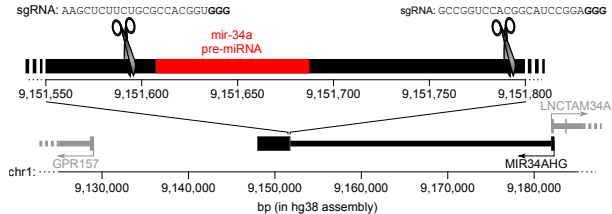
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In human HCT-116 and HAP1 cells:



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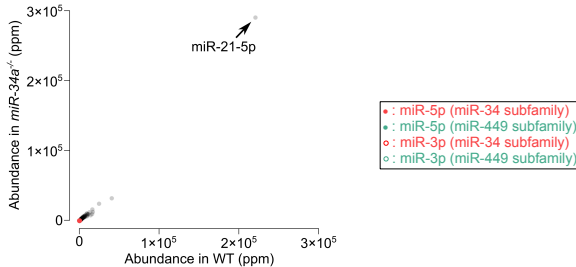
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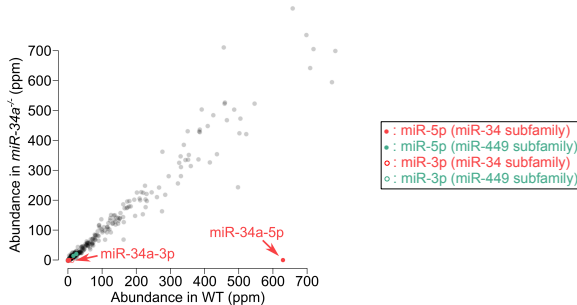
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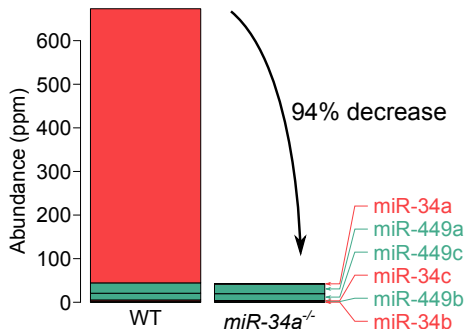
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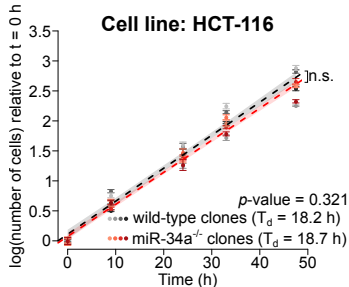
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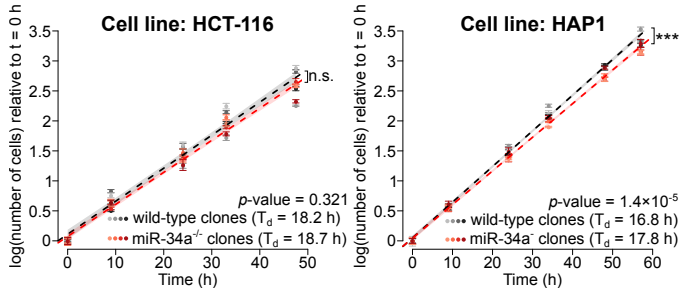
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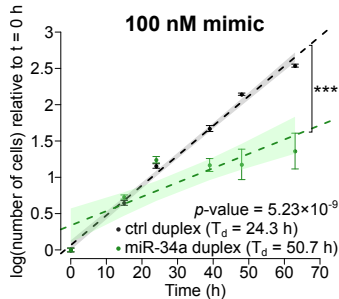
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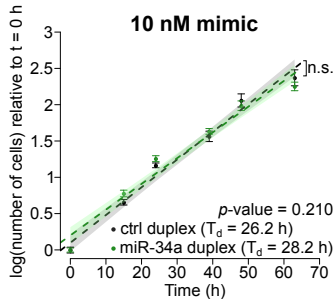
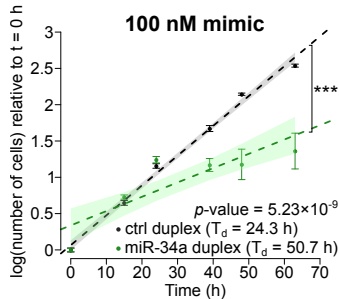
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Time post-transfection (h):	1	24	48	1	24	48	1	24	48	1	24	48
miR-34a mimic (100 nM):	-	-	-	+	+	+	-	-	-	+	+	+
Quantified miRNA:	miR-34a						miR-21					

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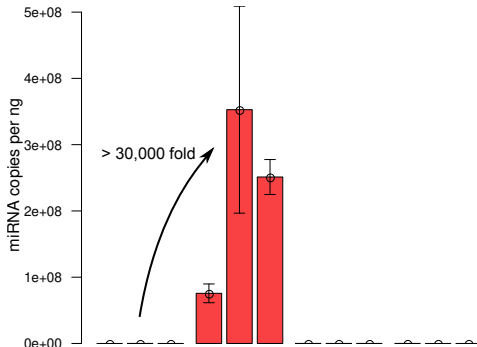
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miR-34a and miR-21 RT-ddPCR



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miR-34a mimic (100 nM):	-	-	-	+	+	+	-	-	-	+	+	+
Quantified miRNA:	miR-34a						miR-21					

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The observed anti-proliferative effect was artifactual (due to unrealistic over-expression).

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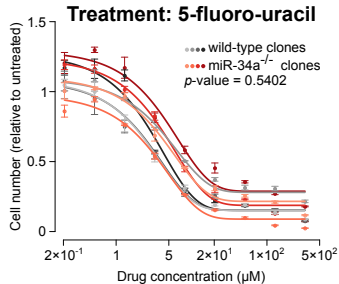
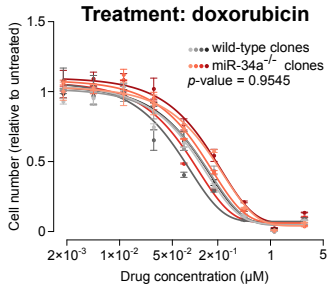
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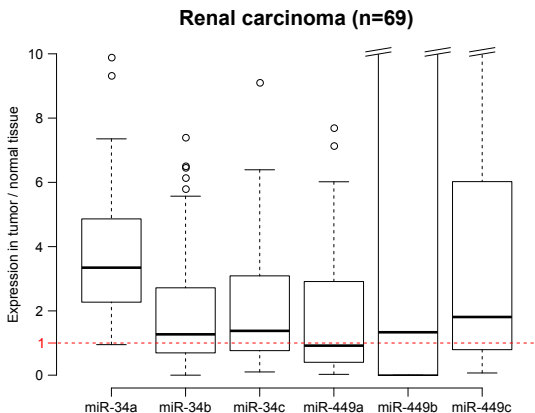
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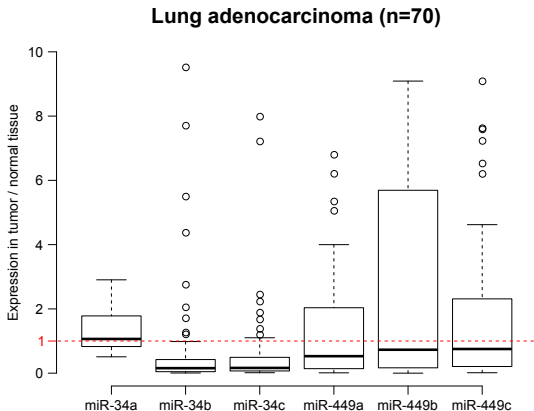
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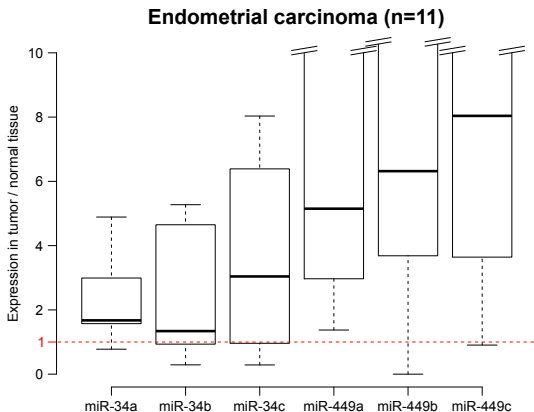
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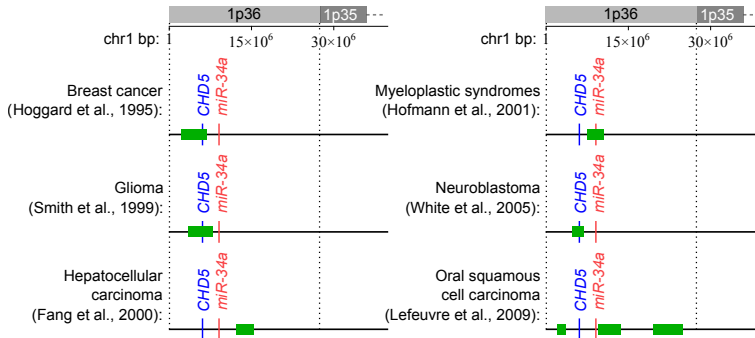
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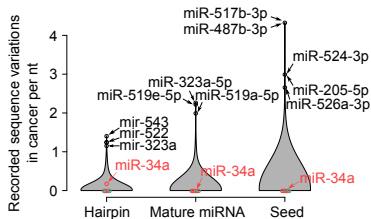
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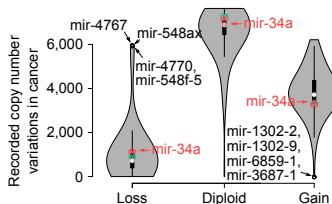
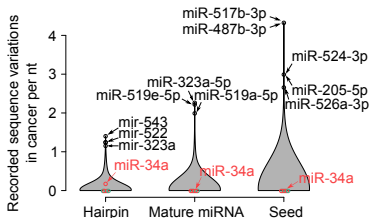
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Sophie
Mockly

Élisabeth
Houbon



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► CLIP protocol

► References: p53 controls miR-34a, b and c

► miR-34 family expression in human cell lines, tissues, body fluids and murine tissues

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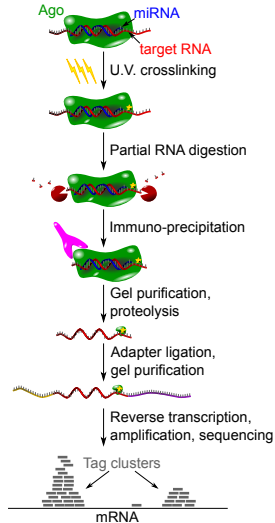
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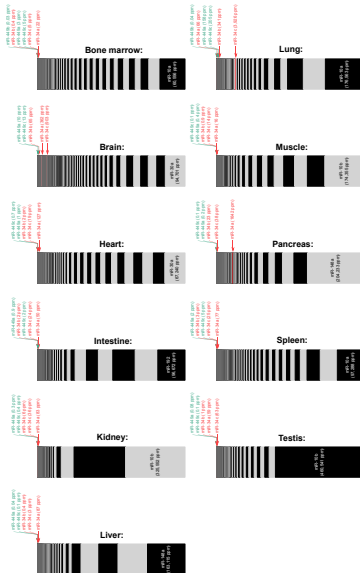
Figure 1 displays a schematic representation of the 12 cell lines used in the study. Each cell line is represented by a horizontal bar divided into segments of different colors (black, grey, white) representing different genomic regions. Red arrows indicate specific genomic locations and associated data. The cell lines are: A549, Hep G2, Caco-2, IMR90, H1299, Kelly, H2170, RKO, HEK293, SW480, HeLa, and U-2 OS.

Supplementary data

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