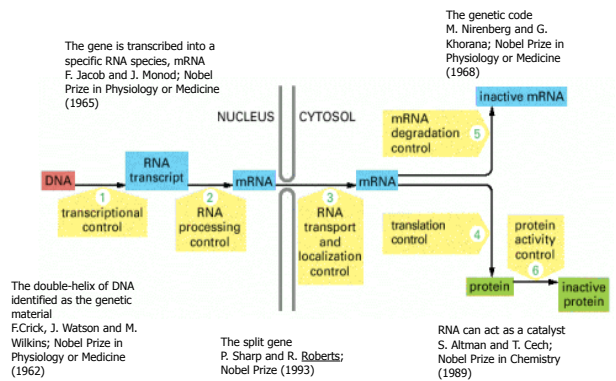


# Small regulatory RNAs - gene silencing



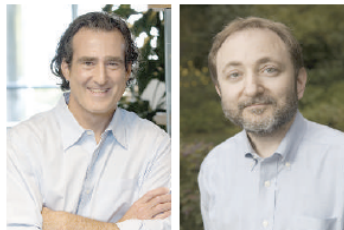
Serena Zacchigna, MD PhD  
Group Leader, Cardiovascular Biology  
ICGEB, Trieste  
zacchign@icgeb.org

# RNA in the CONTROL of GENE EXPRESSION - a Nobel story



## Youthful duo snags a swift Nobel for RNA control of genes

Nobel prize 2006 Physiology and Medicine to Craig Mello and Andrew Fire for their report on RNAi.

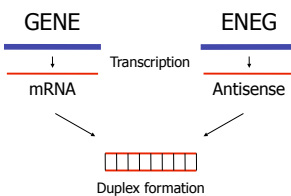


Silence is golden: Craig Mello (left) and Andrew Fire.

## Short history of post-transcriptional gene silencing

- **1962, Singer, Jones, Nirenberg**  
Translation of mRNA can be blocked by complementary (**antisense**) RNA
- **1990, Jorgensen**  
Introduction of transgenes homologous to endogenous genes often results in plants with both genes suppressed (**co-suppression**)
- **1995, Guo and Kempthues**  
Injection of either antisense or sense RNAs in the germline of *C. elegans* is equally effective at silencing homologous target genes
- **1998, Mello and Fire**  
Combination of sense and antisense RNA (= **dsRNA**) is 10 times more effective than ssRNA

## Antisense RNA



When the antisense RNA binds to the complementary mRNA, it forms a double-stranded RNA (dsRNA) complex that is similar to double-stranded DNA. The dsRNA complex does not allow normal translation to occur.

The exact mechanism by which translation is blocked is unknown. Several theories include:

- that the dsRNA prevents ribosomes from binding to the sense RNA and translating (Kimball, Nov 2002)
- that the dsRNA cannot be transported from within the nucleus to the cytosol, which is where translation occurs (Tritton, 1998)
- that dsRNA is susceptible to endoribonucleases that would otherwise not affect single stranded RNA, but degrade the dsRNA (Kimball, Nov 2002)



Right: Flower of a tobacco plant carrying a transgene whose transcript is antisense to one of the mRNAs needed for normal flower pigmentation. Left: Flower of another transgenic plant that failed to have its normal pigmentation altered.  
(van der Krol, et. al., from Nature 333:866, 1988.)

## Longer lasting tomatoes by RNA antisense technology

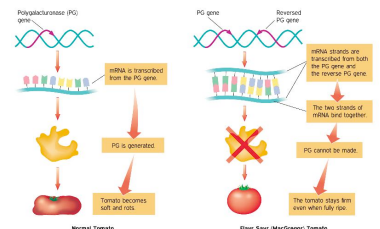


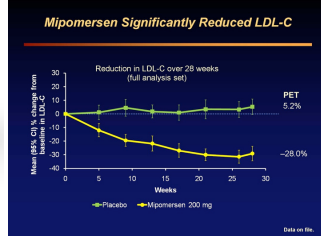
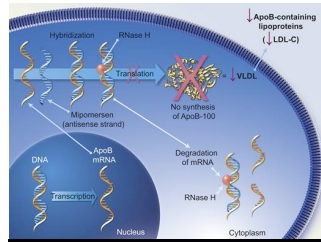
Image shows three sets of tomatoes. The ordinary control tomatoes (extreme left) soften and shrivel up, while texture of gene-silenced tomatoes remains intact for up to 45 days.  
Photo credit: Anis Datta, Subhra Chakraborty, National Institute of Plant Genome Research, New Delhi



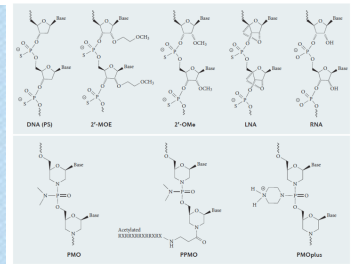
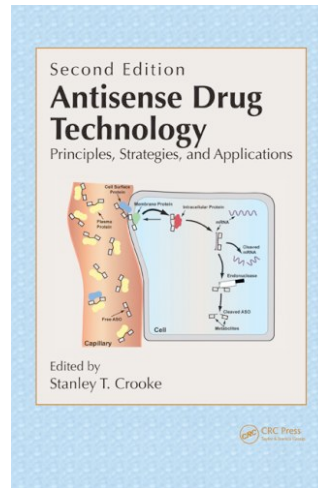
The **Flavr Savr** tomato is a genetically altered tomato developed by Calgene. It contains an antisense RNA which inhibits the expression of a gene that normally causes fruit to soften, therefore, the fruit stays firm longer. This allows producers a greater period of time for transportation and the opportunity for mechanical harvesting with little bruising.

# Antisense approach for lipid management

KYNAMRO® is an oligonucleotide inhibitor of apolipoprotein B-100 synthesis indicated as an adjunct to lipid-lowering medications and diet to reduce low density lipoprotein-cholesterol (LDL-C), apolipoprotein B (apo B), total cholesterol (TC), and non-high density lipoprotein-cholesterol (non-HDL-C) in patients with homozygous familial hypercholesterolemia (HoFH).



Because of the risk of hepatotoxicity, KYNAMRO is available only through a restricted program under a Risk Evaluation and Mitigation Strategy (REMS) called the KYNAMRO REMS.



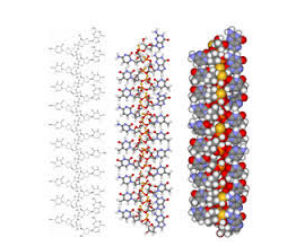
## RNA therapeutics: beyond RNA interference and antisense oligonucleotides

**Ryszard Kole<sup>1</sup>, Adrian R. Krainer<sup>2</sup> and Sidney Altman<sup>3</sup>**  
 Abstract | Here, we discuss three RNA-based therapeutic technologies exploiting various oligonucleotide that bind to RNA by base pairing in a sequence-specific manner yet have different mechanisms of action and effects. RNA interference and antisense oligonucleotides downregulate gene expression by inducing enzyme-dependent degradation of targeted mRNA. Steric blocking oligonucleotides block the access of cellular machinery to pre-mRNA and mRNA without degrading the RNA. Through this mechanism, steric blocking oligonucleotides can redirect alternative splicing, repair defective RNA, restore protein production or downregulate gene expression. Moreover, they can be rationally chemically modified to acquire more drug-like properties. The ability of RNA blocking oligonucleotides to restore gene function makes them best suited for the treatment of genetic disorders. Positive results from clinical trials for the treatment of Duchenne muscular dystrophy show that this technology is close to achieving its clinical potential.

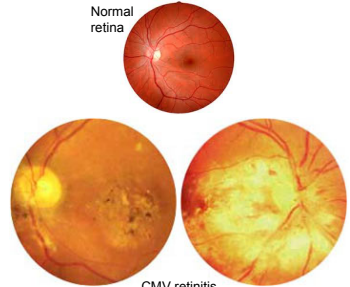
# Fomivirsen

Fomivirsen sodium is a **phosphorothioate oligonucleotide**, 21 nt in length: 5'-GCG TTT GCT CTT CTT CTT GCG-3'

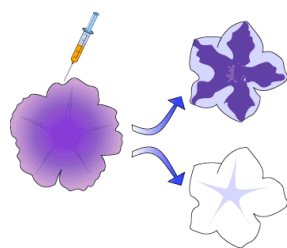
**Vitravene (fomivirsen)** is indicated for CMV infections, and in particular CMV retinitis in patients with acquired immunodeficiency syndrome, who are intolerant of or have a contraindication to other treatment(s) for CMV retinitis or who were insufficiently responsive to previous treatment(s) for CMV retinitis.



Target sequence: IE2 gene of the CMV genome



# Co-suppression



Researchers were trying to deepen the purple colour of the flowers by injecting the gene responsible into the petunias. Surprisingly, instead of a darker flower, the petunias were either variegated or completely white!

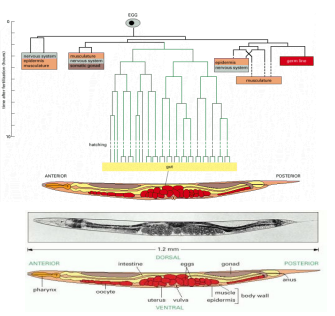
This phenomenon was termed **co-suppression**, since both the expression of the existing gene (the initial purple colour), and the introduced gene (to deepen the purple) were suppressed. Co-suppression has since been found in many other plant species and also in fungi. It is now known that double stranded RNA is responsible for this effect.



# Potent and specific genetic interference by double-stranded RNA in Caenorhabditis elegans

**Andrew Fire<sup>1</sup>, SiQun Xu<sup>1</sup>, Mary K. Montgomery<sup>1</sup>, Steven A. Kostas<sup>1\*</sup>, Samuel E. Driver<sup>1</sup> & Craig C. Mello<sup>2</sup>**  
<sup>1</sup>Carnegie Institution of Washington, Department of Embryology, 115 West University Parkway, Baltimore, Maryland 21210, USA  
<sup>2</sup>Biology Graduate Program, Johns Hopkins University, 3400 North Charles Street, Baltimore, Maryland 21218, USA  
<sup>3</sup>Program in Molecular Medicine, Department of Cell Biology, University of Massachusetts Cancer Center, Two Biotech Suite 213, 373 Plantation Street, Worcester, Massachusetts 01605, USA

Experimental introduction of RNA into cells can be used in certain biological systems to interfere with the function of an endogenous gene<sup>1,2</sup>. Such effects have been proposed to result from a simple antisense mechanism that depends on hybridization between the injected RNA and endogenous messenger RNA transcripts. RNA interference has been used in the nematode *Caenorhabditis elegans* to manipulate gene expression<sup>3,4</sup>. Here we investigate the requirements for structure and delivery of the interfering RNA. To our surprise, we found that double-stranded RNA was substantially more effective at producing interference than was either strand individually. After injection into adult animals, purified single strands had at most a modest effect, whereas double-stranded mixtures caused potent and specific interference. The effects of this interference were evident in both the injected animals and their progeny. Only a few molecules of injected double-stranded RNA were required per affected cell, arguing against stoichiometric interference with endogenous



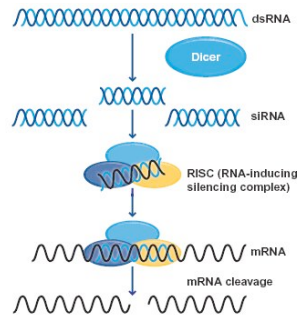
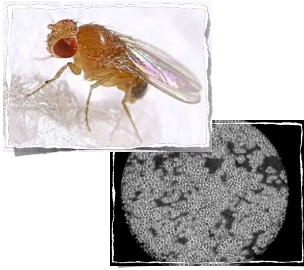
*C. elegans* is a precious tool in developmental biology:  
 - it is tiny and grow rapidly  
 - females are composed of 956 cells  
 - males are composed of 1031 cells  
 - the fate of every cell is characterised

# Conclusions of Fire&Mello's study:

- 1) Silencing is triggered efficiently by **dsRNA**, but weakly or not at all by sense or antisense ssRNAs.
- 2) Silencing is **specific** for an mRNA homologous to the dsRNA; other mRNAs are unaffected.
- 3) The dsRNA has to correspond to **mature mRNA**; neither intron nor promoter sequences trigger a response. This indicated a post-transcriptional, presumably cytoplasmic mechanism.
- 4) The targeted mRNA disappears, suggesting it is **degraded**.
- 5) Only a few dsRNA molecules per cell accomplish full silencing. This indicates that the dsRNA is amplified and/or acts **catalytically** rather than stoichiometrically.
- 6) The effect spreads between tissues and to the progeny, suggesting a **transmission** between cells.

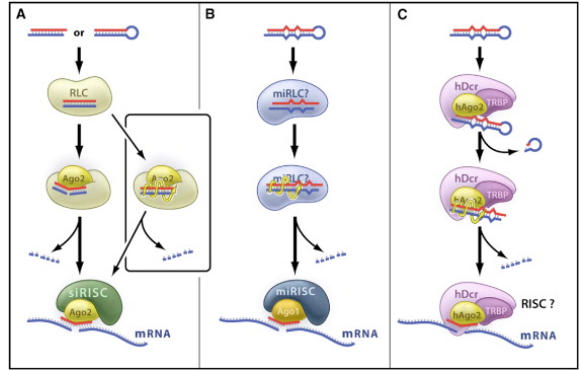
The **unc-22** gene encodes a myofibril protein. Decrease in its activity is known to produce severe twitching movements.  
 Injection of double-stranded *unc-22* RNA, but not single-stranded RNA, into the gonad of *C. elegans* induces the twitching phenotype in the progeny.

# RISC was discovered in Drosophila cells



The molecular machinery responsible for RNAi involves a large complex, called RISC (RNA-induced silencing complex), which is targeted to the mRNA via the antisense RNA. The mRNA is cleaved and subsequently degraded.

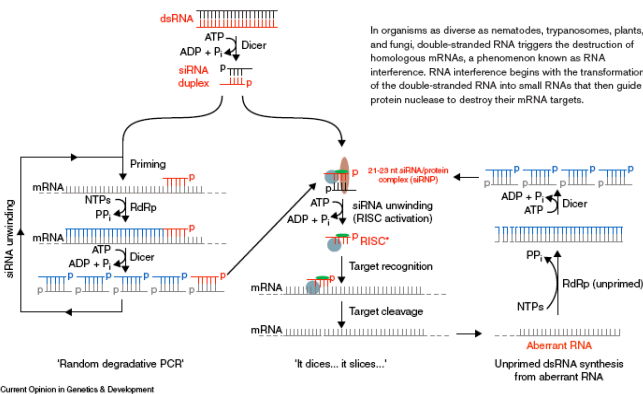
# RNA loading and activation in RISC



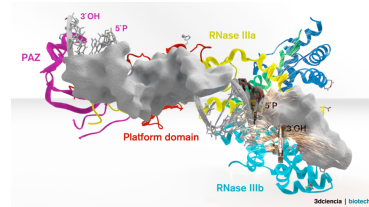
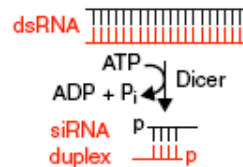
RISC contains at least one member of the **argonaute** protein family, which have endonuclease activity and cut the mRNA.

# RNAi: nature abhors a double-strand

György Hutvagner and Phillip D Zamore\*

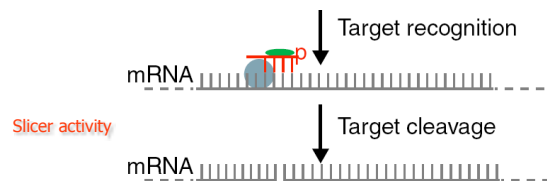


# It dices...



RNAi is initiated by the ATP-dependent, processive cleavage of long dsRNA into 21-25 nt ds-fragments, termed **small interfering RNAs (siRNAs)**. This cleavage is mediated by the enzyme Dicer (a member of the RNase III family of dsRNA-specific endonucleases).

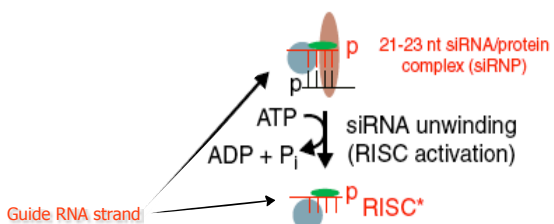
# It slices...



Finally, in a step that requires little or no ATP, RISC\* recognizes and cleaves the target RNA, complementary to the guide strand of the siRNA.

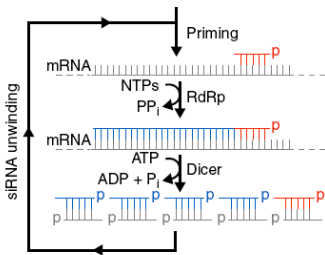
The siRNA duplexes are incorporated into a protein complex that is not yet competent to mediate RNAi.

ATP-dependent **unwinding of the siRNA duplex** remodels the complex to generate an **active RNA-induced silencing complex (RISC)** - the asterisk denotes active conformation



# Random degradative PCR

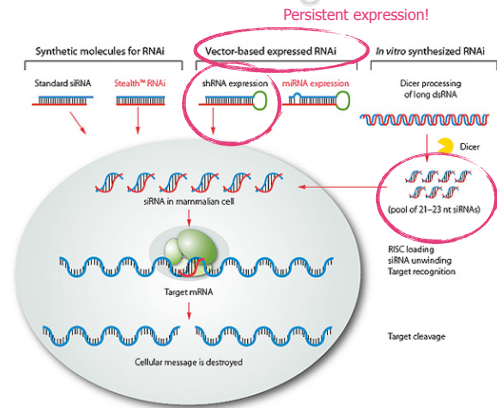
The discovery of RNA-dependent RNA polymerases (RdRPs) in plants, worms and fungi explains the remarkable efficacy of dsRNA in gene silencing - in worms RNAi not only spread throughout the entire animal, but also can be inherited through multiple generations.



In *Drosophila* embryos, 35 molecules of dsRNA can silence a target mRNA present at >1000 copies per cell.

In the "random degradative PCR" model, the RdRp uses the guide siRNA strand as a primer to synthesize new RNA, using the target RNA as a template and thereby converting it into dsRNA, that can be then processed by Dicer. This in turn releases new siRNAs to prime additional rounds of synthesis and target destruction.

# RNAi in mammalian cells - a precious tool for gene silencing

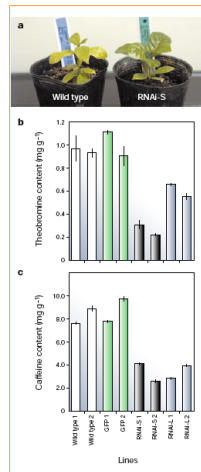


## RNA interference

### Producing decaffeinated coffee plants

Three N-methyltransferases are involved in caffeine biosynthesis - CaXMT1, CaMXMT1 (theobromine synthase) and CaDXMT1 (caffeine synthase).

Coffee plants in which expression of CaMXMT1 is repressed by RNAi have a caffeine content reduced by up to 70%.



# siRNA/shRNA therapeutics in clinical trials

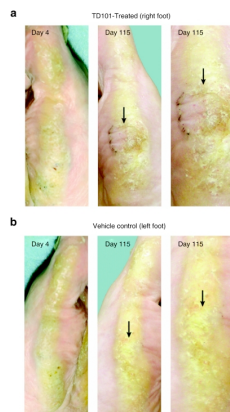
Company	Drug	Delivery route	Target	Vehicle	Disease	Phase
Baxter	SPC049 (LNA)	IPC	miR-122	Naked LNA	HCV	IIIa
Opko Health	Bevrantrah	IVT	VEGF	Naked siRNA	AMD/DMR	III
Adogen/Novus	AGN-548	IVT	VEGF-R1	Naked siRNA	AMD	II
Quark/Phar	QP-666	IVT	HTT/mn	Naked siRNA	AMD/DMR	II
Quark/Pharma	QP-1007	IVT	Caspase 2	Naked siRNA	NAION	I
TransDerm/IPC	TD101	Intradermal injection	KRT6A/NO71K	Naked siRNA	Pachyonychia Congenita	II
Schering	SYL0012	siRNA drops	ADRB2	Naked siRNA	Intercellular Pressure	II
Schering	SYL0005	Cyphalamin drops	TRPV1	Naked siRNA	Dry eye syndrome	I
Zelcor	ExoBifTM	Inhalation	β2k kinase	unknown	Asthma	II
Alnylam/Chibret	ALN-RSV01	Subcutaneous/IV intratumoral	RSV	Naked siRNA	RSV	IIIb
Martin Biotech	CEQ008	Oral	Beta-actin	siRNA in E. coli	FAP/colorectal cancer	I
Silimed Ltd	siGrpD	IV	ERASGrpD	LODIER	PDAC	I
Tekmira	TKM-ApoB	IV	Apo B	SNALP	Hypocholosterolemia	I
Tekmira	TKM-PLK1	IV	PLK1	SNALP	Solid tumours	I
Alnylam/Tekmira	ALN-VSP01	IV	KSP and VEGF	SNALP	Solid tumours	I
Alnylam	ALN-FT001	IV	FTF	SNALP	FTF medication	I
University of Heidelberg	Res-AM	IV	Res-AM	Antisense liposome	CMV	I
Danaher	Ata007	IV	PCN3	siRNA-liposomes	Advanced solid tumor	I
Quark/Pharma	QNP	IV	PS3	Naked siRNA	AKI and PCP	II
Caladrius Pharma	CALAA-01	IV	RRM2	Cyclohexan iminopyridine, TP, and PEG	Solid tumours	I
Genadine Inc.	PANG	Ex vivo IV	Parv and CMV	SNP	Extracorporeal Shockwaves	II
Solid tumor cancer	siRNA	Ex vivo intratumoral injection	LMP1, LMP2, MEK1	Transfection	Metastatic melanoma	I
City of Hope/Bectec	Res-AM	IV	Res-AM	Antisense liposome	CMV	I

## First-in-human mutation-targeted siRNA phase Ib trial of an inherited skin disorder

TransDerm has designed the first mutation-specific siRNA to be used for human therapy.

The TD101 siRNA is directed at the mRNA encompassing the dominant mutation (N171K) in the keratin 6a gene (KRT6A). This mutation causes pachyonychia congenita, a rare skin disorder characterized by painful calluses on weight-bearing areas and hypertrophic nails.

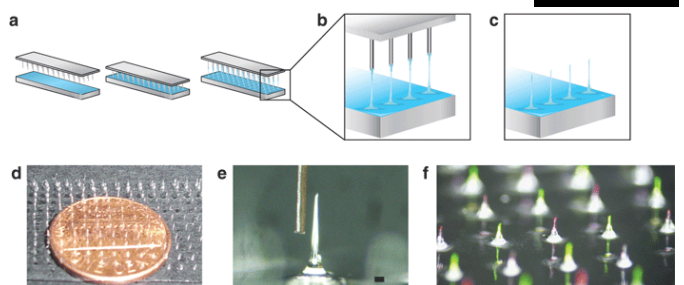
The siRNA therapy is administered by local injection. Since the Phase Ib therapy (NCT00716014) was well tolerated and reduced the callus, TransDerm is developing less painful alternatives for delivering the drug, such as an ointment with lipid-based carriers (GeneCreme) and a dissolvable microneedle array (Protrusion Array Device).



## TRANSDERM, INC

### Protrusion Array Device (PAD)

It consists of a loadable ordered grid of needle-like microprotrusions formed from injection-safe soluble polymers.

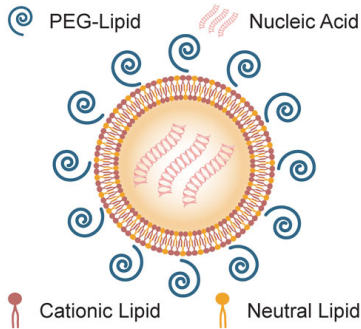


(a) Pin template and glass slide covered with a thin film of 20% polyvinyl alcohol (PVA) solution (left panel). The pin template is placed in contact with the PVA solution (middle panel). Microneedles are produced by withdrawing the pins as the film is drying, forming fiber-like structures (right panel). (b) Enlarged view of fibers. (c) Protrusions are subsequently trimmed to the desired length and tip shape. (d) PAD supported by a glass substrate, with a penny to show scale. (e) Micrograph of one microneedle after trimming to 1 mm length, showing beveled structure that facilitates skin penetration, with a human hair to show scale (bar = 80 μm). (f) PAD needles loaded alternately with fluorescein (green) and R-phycoerythrin (red).



# siRNA lipid nanoparticles (LNPs)

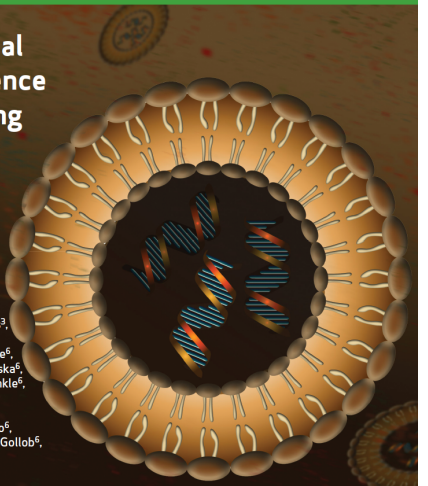
While unmodified siRNAs have been injected locally into the eye and other organs in early trials, those released directly into the bloodstream are degraded by enzymes and are unable to cross cell membranes. One strategy for smuggling siRNAs through the blood and into diseased cells is to embed them in lipid nanoparticles (LNPs).



In vivo, siRNA LNPs generally end up in the liver. The liver is highly vascularized and its endothelium is peppered with pores about 100 nanometers in diameter, wide enough for 70- to 80-nanometer LNPs to slip through en route to hepatocytes. Moreover, once the LNPs are released into the bloodstream, they are rapidly coated with apolipoprotein E (ApoE), which binds to receptors on hepatocytes and eases cell entry of the nanoparticles.

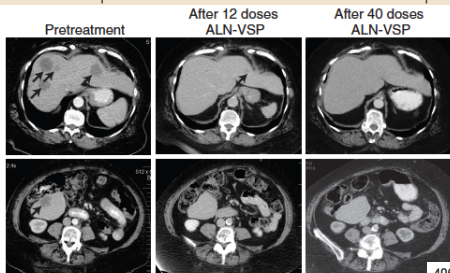
## First-in-Humans Trial of an RNA Interference Therapeutic Targeting VEGF and KSP in Cancer Patients with Liver Involvement

Josep Taberner<sup>1</sup>, Geoffrey I. Shapiro<sup>4</sup>, Patricia M. LoRusso<sup>1</sup>, Andres Cervantes<sup>2</sup>, Gary K. Schwartz<sup>3</sup>, Glen J. Weiss<sup>5</sup>, Luis Paz-Ares<sup>2</sup>, Daniel C. Cho<sup>2</sup>, Jeffrey R. Infante<sup>10</sup>, Maria Alsina<sup>1</sup>, Minal M. Gounder<sup>8</sup>, Rick Falzone<sup>6</sup>, Jamie Harrop<sup>5</sup>, Amy C. Seila White<sup>6</sup>, Iva Toudjarska<sup>6</sup>, David Bumcrot<sup>6</sup>, Rachel E. Meyers<sup>6</sup>, Gregory Hinkle<sup>6</sup>, Nenad Svrzikapa<sup>6</sup>, Renta M. Hutabarat<sup>6</sup>, Valerie A. Clausen<sup>6</sup>, Jeffrey Cehelsky<sup>6</sup>, Saraswathy V. Nochur<sup>6</sup>, Christina Gamba-Vitalo<sup>6</sup>, Akshay K. Vaishnav<sup>6</sup>, Dinah W.Y. Sah<sup>6</sup>, Jared A. Gollob<sup>6</sup>, and Howard A. Burris III<sup>10</sup>



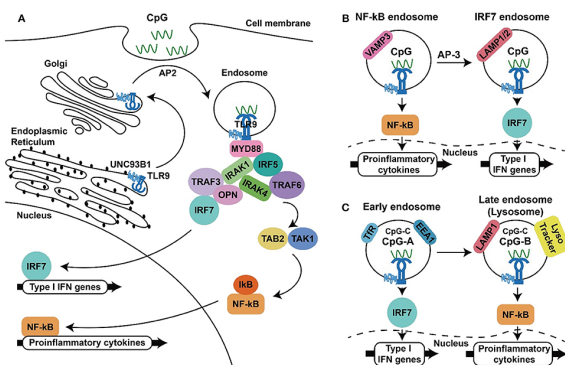
### ABSTRACT

RNA interference (RNAi) is a potent and specific mechanism for regulating gene expression. Harnessing RNAi to silence genes involved in disease holds promise for the development of a new class of therapeutics. Delivery is key to realizing the potential of RNAi, and lipid nanoparticles (LNP) have proved effective in delivery of siRNAs to the liver and to tumors in animals. To examine the activity and safety of LNP-formulated siRNAs in humans, we initiated a trial of ALN-VSP, an LNP formulation of siRNAs targeting VEGF and kinesin spindle protein (KSP), in patients with cancer. Here, we show **detection of drug in tumor biopsies, siRNA-mediated mRNA cleavage in the liver, pharmacodynamics suggestive of target downregulation, and antitumor activity, including complete regression of liver metastases in endometrial cancer.** In addition, we show that biweekly intravenous administration of ALN-VSP was safe and well tolerated. These data provide proof-of-concept for RNAi therapeutics in humans and form the basis for further development in cancer.

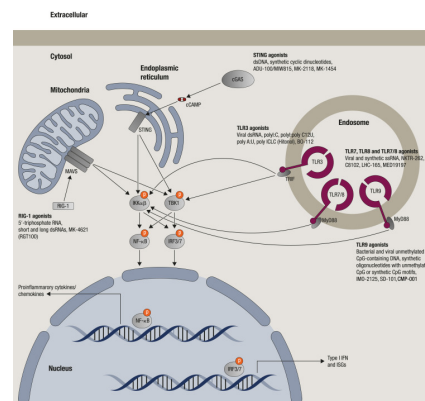


Drug name	Clinical advance	Mechanism of action	Therapeutic group
Ampligen (Jiang et al., 2008)	Launched—2017	TLR3 Agonists (Jiang et al., 2008; Izboren et al., 2016)	Bladder cancer, breast cancer, CRC, ovarian cancer, pancreatic cancer, prostate cancer, renal cancer
Genesense (Jin et al., 2019)	Pre-registered	Apoptosis inducers, BCL2	Breast cancer, CRC, gastric cancer, liver cancer, lung cancer, pancreatic cancer, prostate cancer, renal cancer
Leifolimid	Phase III	TLR9 agonists	CRC, small cell lung cancer
IMO-2125 (Wang et al., 2016)	Phase III	Cytokine, TLR9 agonists	CRC, head and neck cancer, NSCLC
LY-900033 (Mitsuru-Caleiro et al., 2004)	Phase III	PK3A	Breast cancer, NSCLC, ovarian cancer
Inxeltat sodium	Phase III/III	Telomerase reverse transcriptase inhibitors	Breast cancer, liver cancer, neurologic cancer, NSCLC, ovarian cancer
Oncomy-NG3	Phase III/III	MYC	Bladder cancer, breast cancer, lung cancer, pancreatic cancer, prostate cancer, renal cancer
BNT-122	Phase II		Bladder cancer, breast cancer, CRC, NSCLC, pancreatic cancer, renal cancer
NCI-4650	Phase II		Digestive/Gastrointestinal cancer
siG12D/LODER	Phase II	KRAS (Gly12Asp mutant)	Pancreatic cancer
Daravarsen	Phase II	STAT3	Bladder cancer, CRC, liver cancer, NSCLC, pancreatic cancer
ATU-027	Phase II	PKM3	Digestive/Gastrointestinal cancer, pancreatic cancer
EGEN-001	Phase II		Brain cancer, CRC, ovarian cancer, pancreatic cancer
Apatorsen sodium	Phase II	Heat shock protein 27, HSPB1	Bladder cancer, breast cancer, NSCLC, ovarian cancer, pancreatic cancer, prostate cancer
ISIS-EF4ERx	Phase II	EIF4E	NSCLC, prostate cancer
AEG-35156	Phase II	BIRC4	Breast cancer, liver cancer, NSCLC, pancreatic cancer
ACT-GPO-777	Phase II	Anti-nucleolin (NCL)	Lung cancer, pancreatic cancer, renal cancer
ISISLM	Phase II	TLR9 agonists	CRC, renal cancer
ISIS-23722	Phase II	BIRC5 (survivin)	NSCLC, prostate cancer
GTI-2040	Phase II	FRM2	Bladder cancer, breast cancer, CRC, NSCLC, prostate cancer, renal cancer
Agatolmod sodium	Phase II	TLR9 agonists	Breast cancer, NSCLC, prostate cancer, renal cancer
ISIS-2503	Phase II	HRAS	Breast cancer, CRC, NSCLC, pancreatic cancer
CGP-69848A	Phase II	RAF1	Breast cancer, ovarian cancer
Poly I: CLC	Phase II	TLR3 agonists	CRC, liver cancer, neurologic cancer, ovarian cancer, pancreatic cancer, prostate cancer
ARB-1598	Phase I	TLR9 agonists	CRC, head and neck cancer, NSCLC
Emuipcap pegol	Phase I/II	Anti-CCL2 (C-C motif chemokine 2; MCP-1)	Pancreatic cancer, solid tumors
Archein	Discontinued	AKT1	Liver cancer, ovarian cancer, pancreatic cancer, renal cancer
AS TRPM2 ODN	Discontinued	CLU	Breast cancer, NSCLC, prostate cancer

## Adjuvant Effect of Toll-Like Receptor 9 Activation



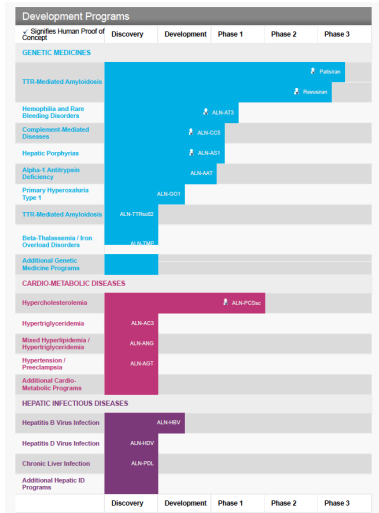
## Intratatumoral immunotherapy: activation of nucleic acid sensing pattern recognition receptors





Alnylam now concentrates on liver-based diseases, with more than 15 RNAi therapies in clinical development for 3 strategic areas:

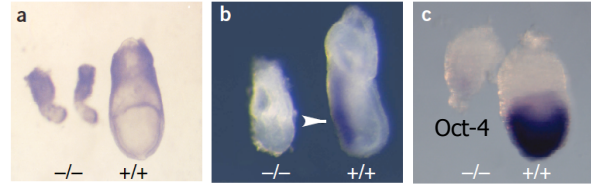
1. Genetic Medicine: treatment of rare diseases
2. Cardio-Metabolic: liver-expressed disease targets for unmet needs in dyslipidemia, hypertension, non-alcoholic steatohepatitis (NASH), type 2 diabetes
3. Hepatic Infectious Diseases: HBV, HDV



## The endogenous role of RNAi

### Dicer is essential for mouse development

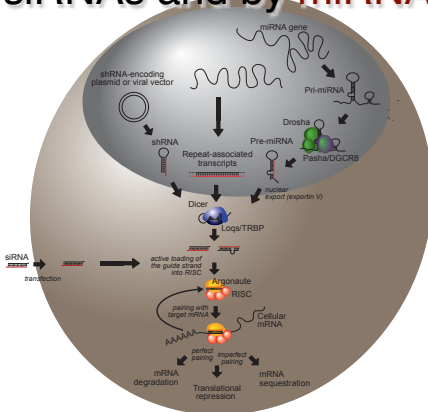
Emily Bernstein<sup>1,2</sup>, Sang Yong Kim<sup>1</sup>, Michelle A Carmell<sup>1,2</sup>, Elizabeth P Murchison<sup>1</sup>, Heather Alcorn<sup>3</sup>, Mammie Z Li<sup>4</sup>, Alea A Mills<sup>1</sup>, Stephen J Elledge<sup>4</sup>, Kathryn V Anderson<sup>3</sup> & Gregory J Hannon<sup>1</sup>



E7.5 embryos - lack of stem cell development

NATURE GENETICS VOLUME 35 | NUMBER 3 | NOVEMBER 2003

## RNAi in mammalian cells works by siRNAs and by miRNAs

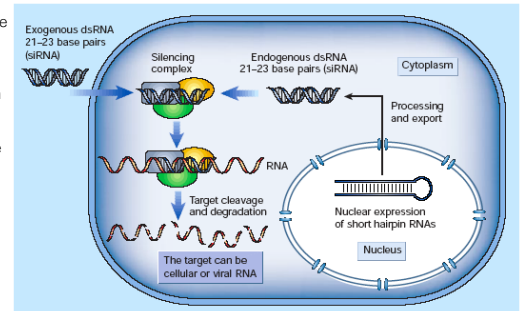


miRNAs are produced by the successive action of two RNaseIII ribonucleases. After transcription, primary miRNAs are cleaved in the nucleus by Drosha. Pre-miRNAs bind exportin V and is exported to the cytoplasm, where Dicer binds the base of the pre-miRNA stem. Dicer cleavage liberates a duplex comprising the miRNA and miR\*.

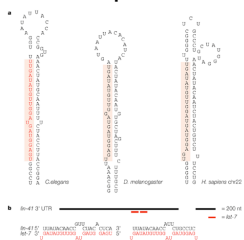
The miRNA is then unwound and selectively incorporated into RISC to search for targets by its seed sequence.

miRNAs, siRNAs and shRNAs differ in their biogenesis, not in their function

Short dsRNAs can be introduced into cells from the outside, or are produced within the cell nucleus from longer precursors forming hairpin structures, which are cleaved to generate shorter RNAs (21-23 bp), that are then exported to the cytoplasm

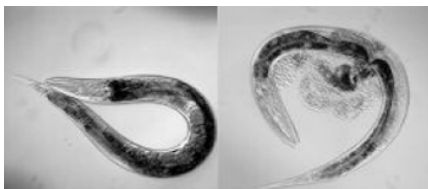


## Developmental control by miRNAs

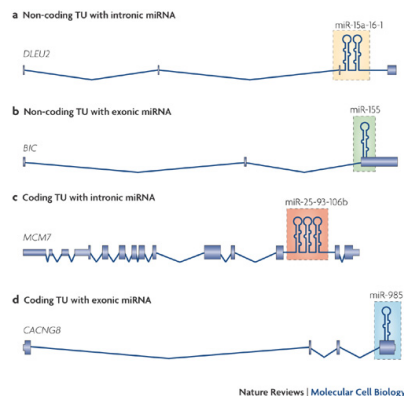


Worms with a mutated form of the microRNA let-7 (right) have severe growth problems, rupturing as they develop.

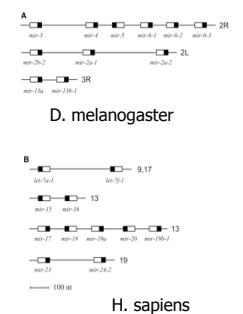
Predicted let-7 precursor RNAs of *Caenorhabditis elegans*, *Drosophila melanogaster* and *Homo sapiens*. The region that corresponds to the mature let-7 RNA is shaded pink



## Genes coding for microRNAs

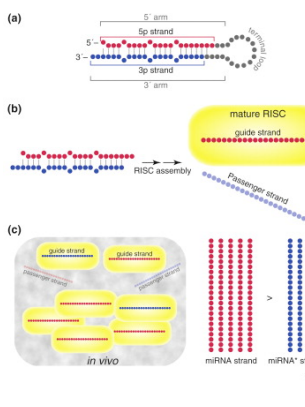


### Gene clusters for miRNAs:



Nature Reviews | Molecular Cell Biology

# Nomenclature for miRNA strands

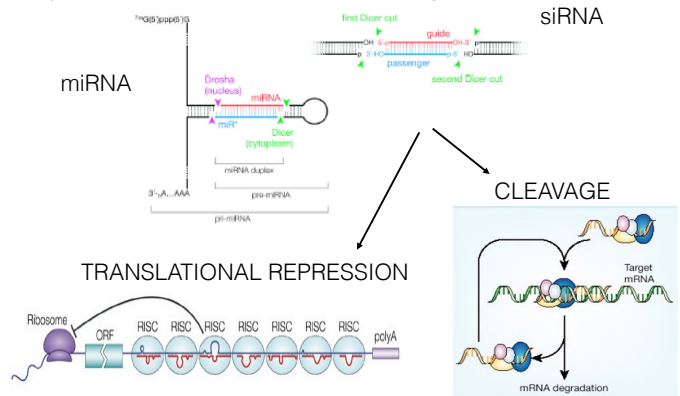


**5p and 3p strands:** 5p and 3p designate the strands derived from the 5' arm and 3' arm of a pre-miRNA, respectively.

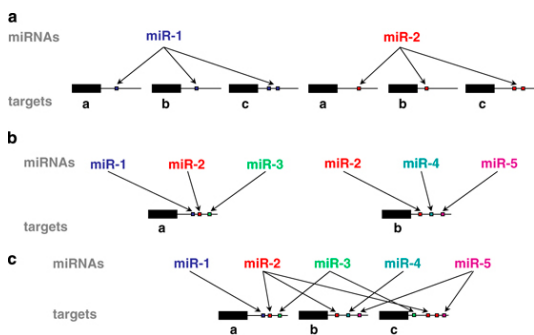
**Guide and passenger strands:** the guide strand is retained in the mature RISC whereas the passenger strand is discarded upon unwinding. Which strand is selected as the guide is independent of the original orientation within the pre-miRNA (i.e. 5p or 3p) or long dsRNA precursors, but does depend on the thermodynamic asymmetry, the 5' nucleotide identity and the structure of the small RNA duplex.

**miRNA and miRNA\* strands:** the miRNA strand is the more abundant (and thereby more frequently cloned) strand overall in vivo whereas the miRNA\* strand is the less abundant strand. Note that a passenger strand is quickly degraded as soon as it is discarded from pre-RISC whereas a guide strand is protected from nucleases in the mature RISC. Consequently, the strand that is more likely to serve as the guide strand tends to accumulate and therefore become the 'miRNA strand'.

Like siRNAs, miRNAs can cleave their mRNA targets when the two are extensively complementary, but repress mRNA translation when they are not

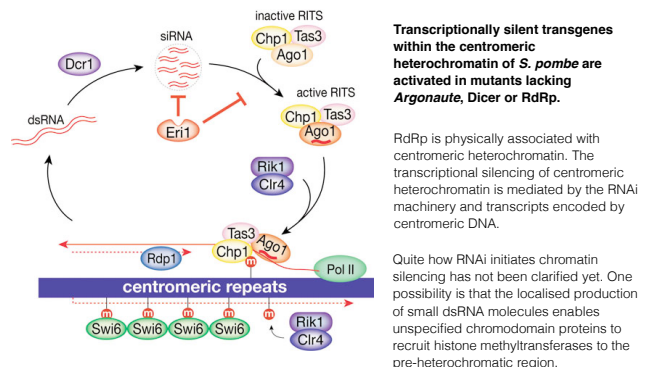


# miRNAs target multiple genes and genes are targeted by multiple miRNAs



(a) miRNAs have multiple targets. (b) Many genes have seed matches for multiple miRNAs in their 3'UTRs. (c) A complex network of mutual interactions between miRNAs and mRNAs.

# RITS Connects RNAi and Heterochromatin Formation Machinery



**Transcriptionally silent transgenes within the centromeric heterochromatin of *S. pombe* are activated in mutants lacking *Argonaute*, *Dicer* or *RdRp*.**

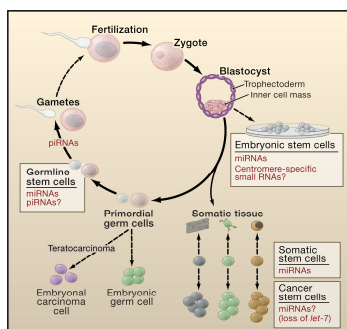
RdRp is physically associated with centromeric heterochromatin. The transcriptional silencing of centromeric heterochromatin is mediated by the RNAi machinery and transcripts encoded by centromeric DNA.

Quite how RNAi initiates chromatin silencing has not been clarified yet. One possibility is that the localised production of small dsRNA molecules enables unspecified chromodomain proteins to recruit histone methyltransferases to the pre-heterochromatic region.

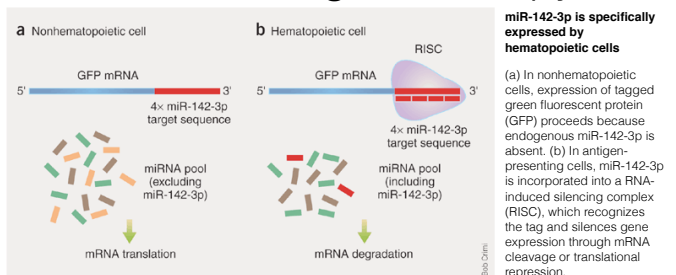
# Small RNAs: Keeping Stem Cells in Line

Bradford M. Stadler<sup>1</sup> and Hannelore Ruchholz-Baker<sup>1\*</sup>  
<sup>1</sup>Department of Biochemistry and Institute for Stem Cell and Regenerative Medicine, University of Washington, 1705 NE Pacific Street, Health Science Building, Room J-587, Seattle, WA 98195, USA  
 \*Correspondence: hannelor@u.washington.edu  
 DOI 10.1038/nrn1209a.02.009

Stem cells and RNA silencing have emerged as areas of intense interest for both basic and clinical research. Recently these fields have converged with reports implicating small regulatory RNAs in the maintenance and pluripotency of stem cells.



# RISC control for gene therapy



**miR-142-3p is specifically expressed by hematopoietic cells**

(a) In non-hematopoietic cells, expression of tagged green fluorescent protein (GFP) proceeds because endogenous miR-142-3p is absent. (b) In antigen-presenting cells, miR-142-3p is incorporated into a RNA-induced silencing complex (RISC), which recognizes the tag and silences gene expression through mRNA cleavage or translational repression.

Endogenous microRNA regulation suppresses transgene expression in hematopoietic lineages and enables stable gene transfer

Brian D Brown<sup>1</sup>, Mary Anna Venneri<sup>1</sup>, Anna Zingale<sup>1</sup>, Lucia Sergi Sergi<sup>1</sup> & Luigi Naldini<sup>1,2</sup>



## Eat Less, Live Longer? miRNAs Link Calorie Restriction To Longevity



Caloric restriction (CR) is the most effective environmental method to increase lifespan (and to prevent late-onset diseases!)

Dietary restriction extends lifespan in *S. cerevisiae*, *C. elegans*, *D. melanogaster*, rodents and primates.

**CR = 60-70% of what an animal would eat at libitum**

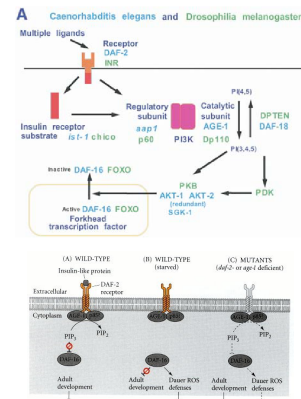
In rodents CR results in as much as a 50% increase in rodent longevity

Physiological effects of CR: acute phase followed by an adaptive period of several weeks to reach a stable, altered physiological state characterized by lower body temperature, lower blood glucose and insulin levels and reduced fat and weight.

The CR animals are more resistant to external stressors, including heat and oxidative stress; organs are typically smaller (except for the brain)

CR animals are resistant to disease, including **cancer** and **infections**

## Mutants in the IIS pathway with extended lifespan (~50-80%) in *C. elegans* and *D. melanogaster*



### *C. elegans*

age-1: catalytic subunit of PI3 kinase  
daf-2: Insulin/IGF1 receptor  
daf-16: fork-head (FOXO) transcription factor

Mutations in the GH axis, which in turn impair the IIS activity

### *Drosophila*

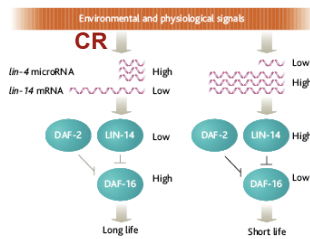
#### IIS receptor

Insulin receptor substrate (chico)

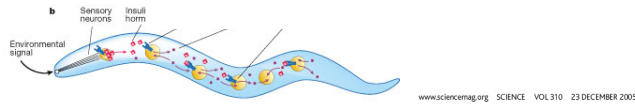
Ablation of neurosecretory cells producing insulin-like ligands

Overexpression of the forkhead transcription factor (dFOXO) in the fat body  
all increase lifespan up to 85%

## Eat Less, Live Longer? miRNAs Link Calorie Restriction To Longevity

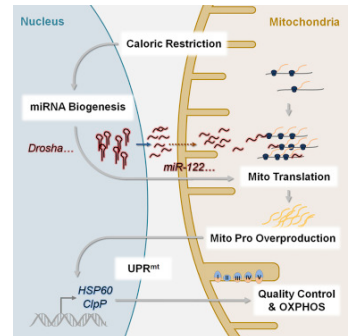


Regulation of adult life span by the *lin-4* microRNA. (Left) When *lin-4* microRNA activity is high, expression of *lin-4* mRNA and protein are low. Hence, the DAF-2 transcription factor is active and promotes long life. (Right) When *lin-4* activity is low, *lin-4* activity is high, and DAF-16 is inhibited, resulting in short life. *lin-4* and *lin-14* gene products may work downstream of or in parallel to DAF-2 (the insulin-like receptor) to modulate DAF-16. Proteins are depicted as oval shapes.



## Caloric Restriction induces miRNAs to improve mitochondrial proteostasis

- CR increases miRNA biogenesis and the global expression of miRNAs in mitochondria
- miRNAs are critical for CR-induced activation of mitochondrial translation
- CR-induced miRNAs cause overproduction of mtDNA-encoded proteins and induce UPR<sup>mt</sup>



## Treatment of HCV Infection by Targeting MicroRNA

Harry L.A. Janssen, M.D., Ph.D., Hendrik W. Reesink, M.D., Ph.D., Eric J. Lawitz, M.D., Stefan Zeuzem, M.D., Maribel Rodriguez-Torres, M.D., Keyur Patel, M.D., Adriaan J. van der Meer, M.D., Amy K. Patock, Ph.D., Alice Chen, B.A., Yi Zhou, Ph.D., Robert Persson, Ph.D., Barney D. King, M.D., Sakari Kauppinen, Ph.D., Arthur A. Levin, Ph.D., and Michael R. Hodges, M.D.

N Engl J Med 2013; 368:1685-1694 May 2, 2013 DOI: 10.1056/NEJMoa1209026



### Miraviren Mode of Action

- Miraviren is a LNA modified phosphorothioate anti-sense oligonucleotide targeting and blocking miR-122
- First drug to exploit a microRNA target for therapeutic use
- As a host targeting agent miraviren poses a high barrier to resistance
- Miraviren should work in all HCV genotypes because miR-122 binding sites are conserved



### HCV replication depends on miR-122 expression.

The use of miraviren in patients with chronic HCV genotype 1 infection showed prolonged dose-dependent reductions in HCV RNA levels without evidence of viral resistance. (Funded by Santaris Pharma; ClinicalTrials.gov number, NCT01200420.)

## MIRNA THERAPEUTICS

pioneering microRNA Replacement Therapy

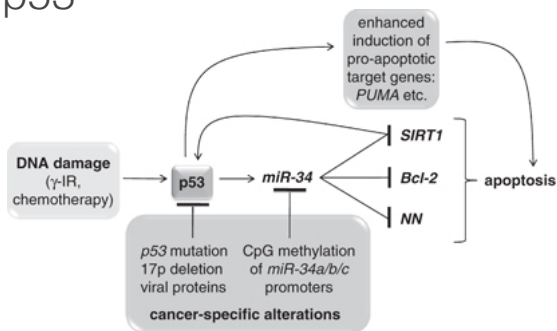
### MRX34 is a first-in-class cancer therapy and the first microRNA mimic to enter clinical trials.

Mirna has secured an exclusive license from Marina Biotech, Inc. to the patent estate covering the SMARTICLES® liposomal delivery technology for several of our lead microRNA product candidates, including miR-34, let-7 and two other undisclosed targets. The SMARTICLES formulation offers key efficacy and safety benefits, including the ability to deliver high numbers of microRNA mimic molecules to cancers cells in the liver, spleen and other highly vascularized tissues, as well as bone marrow and malignant lymphocytes.



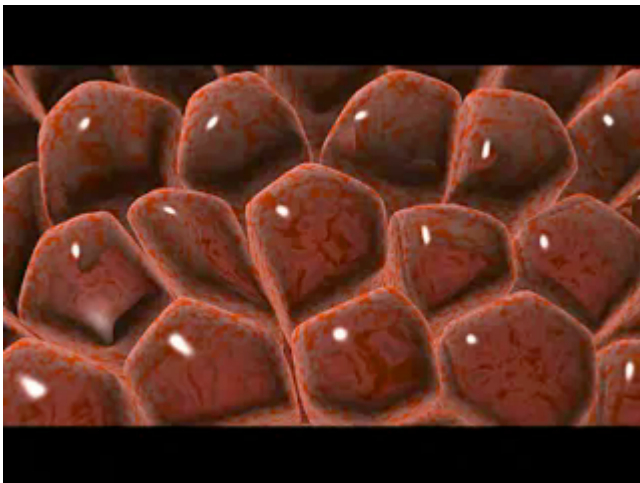
Program	Key Oncogenic Targets	Indication	Discovery	In Vivo Formulation	Preclinical	Phase 1	Phase 2
MRX34 miR-34 mimic	BCL2, E2F3, HDAC1, MET, MEK1, CDK4, TGF-α, WNT1/3, NOTCH-1	Primary liver cancer & solid cancers with liver metastases Hematological malignancies	█	█	█	█	█
miR-Rxlet-7 let-7 mimic	RAS, MYC, HMGA2, TGFβ1, MYCN, Cyclin D2, IL6, IGF1R3		█	█	█	█	
miR-Rx06	UNDISCLOSED		█	█	█	█	
miR-Rx07	UNDISCLOSED		█	█	█	█	
miR-Rx16 miR-16 mimic	BCL2, VEGF-A, Cyclin-D1, HMGA1, FGFR1, CDK6, BMI1		█	█	█	█	

The miR-34 gene family is a mediator of tumor suppression by p53



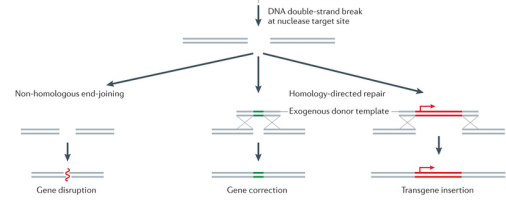
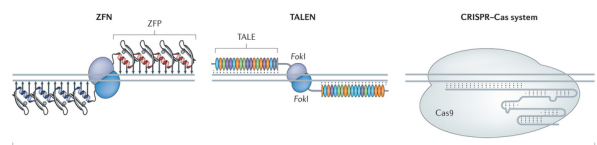
siRNA/miRNA cancer therapy in clinical trials

Drug	Target	Disease	Company	Stage
<b>siRNA Cancer therapeutics in clinical trials</b>				
CALAA-01	M2 subunit of ribonucleotide	Solid tumors	Calando Pharmaceuticals	Ongoing Phase I, Not recruiting
ALN-VSP02	VEGF and KSP	Solid tumors involving the liver	Aplyam Pharmaceuticals	Completed Phase I
Atu027	Protein Kinase 3 (PKN3)	Solid tumors	Silence Therapeutics AG	Completed Phase I
TKM 080301	Polo-like kinase 1	Solid tumors	Tekmira Pharmaceutical	Recruiting Phase I
siG12D LODER	KRAS	Pancreatic ductal adenocarcinoma	Silenced Ltd	Phase II, Not yet open
siRNA-Epha2-DOPC	EPHA2	Solid tumors	M.D. Anderson Cancer Center	Phase I, not yet open
<b>MiRNA Cancer therapeutics in clinical trials</b>				
MRX34	miR-34 mimic	Liver cancer or metastatic cancer with liver involvement	Mima Therapeutic, Inc.	Recruiting Phase I



### Genome editing technology

- zinc finger nucleases (ZFNs)
- transcription activator-like effector nucleases (TALENs)
- clustered regularly interspaced short palindromic repeat (CRISPR)/Cas system



Nature Reviews Genetics 15, 541–555 (2014)



Serena Zacchigna, MD PhD  
 Group Leader, Cardiovascular Biology  
 ICGEB, Trieste  
[zacchign@icgeb.org](mailto:zacchign@icgeb.org)