



RNAi as an insect pest control strategy: challenges and biosafety considerations

Dr. ir. Olivier Christiaens

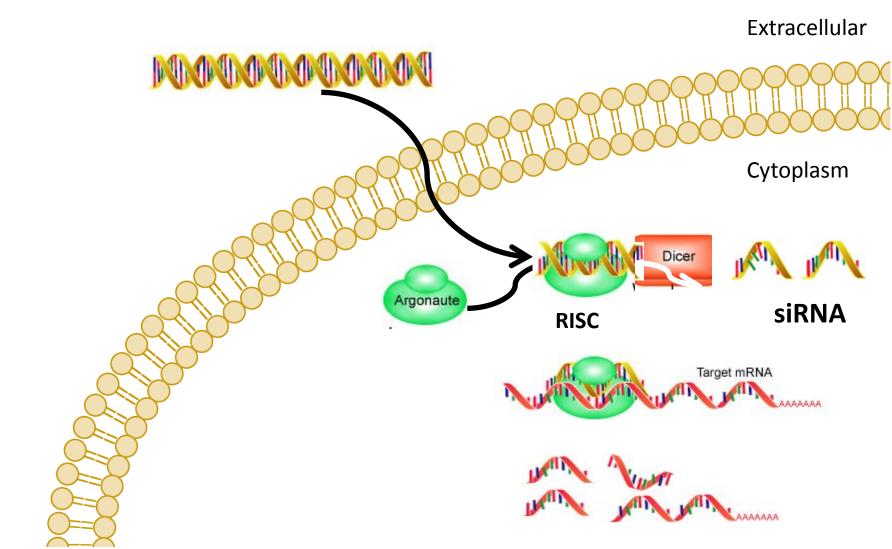
Introduction to RNAi

Basic principle



Introduction to RNAi

Molecular mechanism of the siRNA pathway

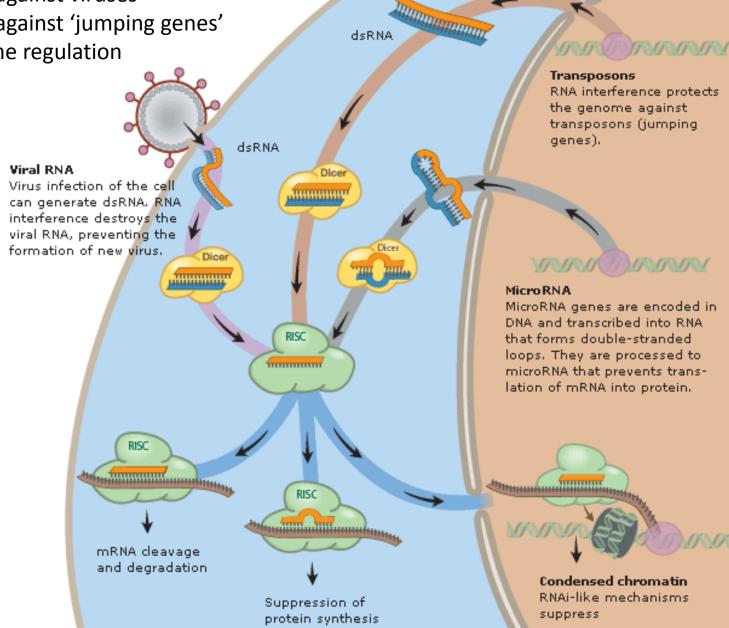


Cellular functions:

CYTOPLASM



- Protection against viruses
- Protection against 'jumping genes'
- Internal gene regulation

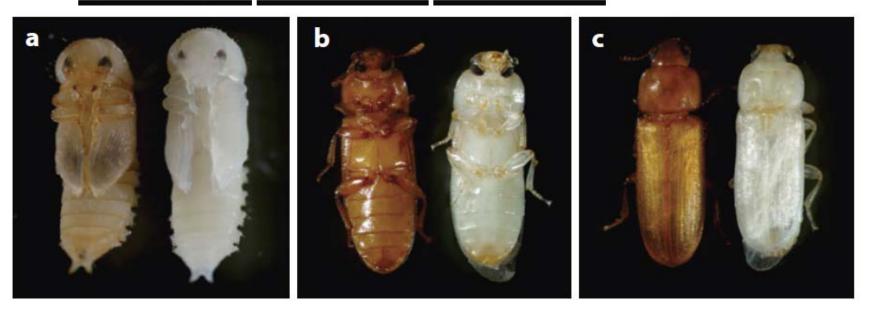


Introduction to RNAi

Applications

Functional genomics

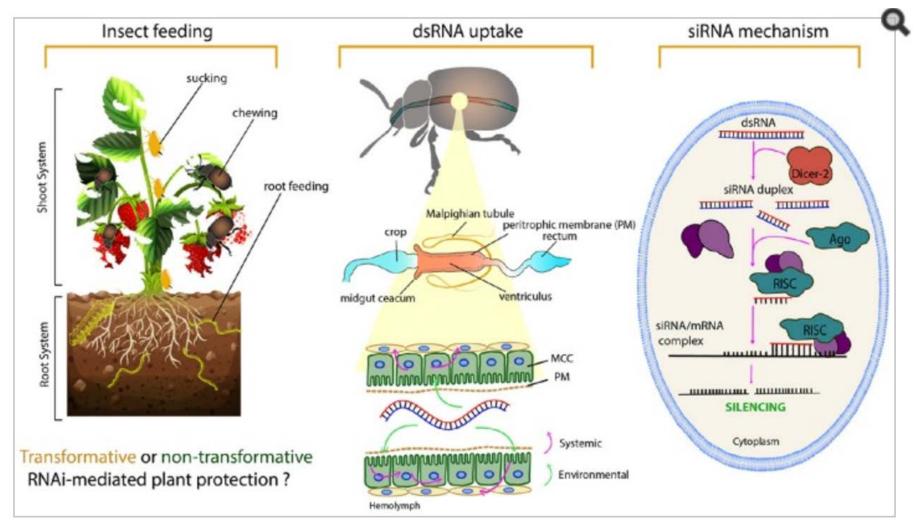
Knockdown of *laccase-2* in T castaneum Arakane *et al.*, 2005, PNAS 102



RNAi of the gene *Laccase 2*, which expresses a phenoloxidase in larvae of *Tribolium castaneum*, prevents tanning after the pupal (a) and imaginal (b, c) molts (2). In each photograph the control is shown at the left and the RNAi knockdown at the right. Photos courtesy of Yas Arakane.

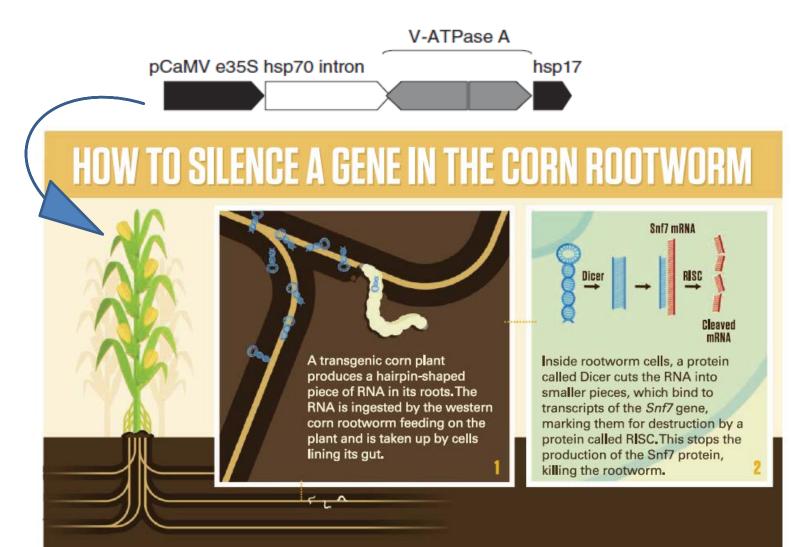
RNAi as a crop protection technology

Silencing essential genes to cause toxicity in insect pests



RNAi as a crop protection technology

First proof of concept: Baum et al., 2007



Possible Applications

2. Crop protection

Baum et al (2007), Nature Biotechnology 25



Possible Applications

2. Crop protection

Monsanto SmartStax Pro (MON87411 X DAS-59122-7)

- Cry3Bb1
- Cry34Ab1/Cry34Ab2
- Glyphosate resistance
- dsRNA targeting the snf7 gene
- Approval for cultivation in USA, Canada and Brazil



Regulatory Toxicology and Pharmacology Volume 81, November 2016, Pages 77-88



Ecological risk assessment for DvSnf7 RNA: A plantincorporated protectant with targeted activity against western corn rootworm

Pamela M. Bachman A ⊠, Kristin M. Huizinga, Peter D. Jensen, Geoffrey Mueller, Jianguo Tan, Joshua P. Uffman, Steven L. Levine

<u>Transgenic Res</u>. 2013; 22: 1207–1222. Published online 2013 Jun 8. doi: <u>10.1007/s11248-013-9716-5</u> PMCID: PMC3835954

Characterization of the spectrum of insecticidal activity of a doublestranded RNA with targeted activity against Western Corn Rootworm (*Diabrotica virgifera virgifera* LeConte)

Pamela M. Bachman,^{®1} Renata Bolognesi,² William J. Moar,¹ Geoffrey M. Mueller,¹ Mark S. Paradise,¹ Parthasarathy Ramaseshadri,² Jianguo Tan,¹ Joshua P. Uffman,¹ JoAnne Warren,¹ B. Elizabeth Wiggins,² and Steven L. Levine¹

Author information
Article notes
Copyright and License information

Pest Manag Sci. 2017 Sep;73(9):1883-1899. doi: 10.1002/ps.4554. Epub 2017 Mar 17.

Evaluation of SmartStax and SmartStax PRO maize against western corn rootworm and northern corn rootworm: efficacy and resistance management.

Head GP¹, Carroll MW¹, Evans SP¹, Rule DM², Willse AR¹, Clark TL¹, Storer NP², Flannagan RD¹, Samuel LW¹, Meinke LJ³.

RNAi as a crop protection technology

Challenges

Delivery

• Genetically modified plants: regulation, public opinion





- Virus-mediated delivery
- Sprayable biopesticide: Cost? Stability of dsRNA in the field?
- Non-GM in planta applications (stem injection, root drenching)

RNAi as a crop protection technology

Challenges

Efficiency of (oral) RNAi in arthropods:

•Very variable

Coleoptera > Diptera, Hemiptera, Orthoptera > Lepidoptera

• Multiple factors

dsRNA stability in the digestive system

Cellular uptake efficiency

Viral infections

RNAi effector gene repertoire

dsRNA design

RNAi-mediated crop protection: potential

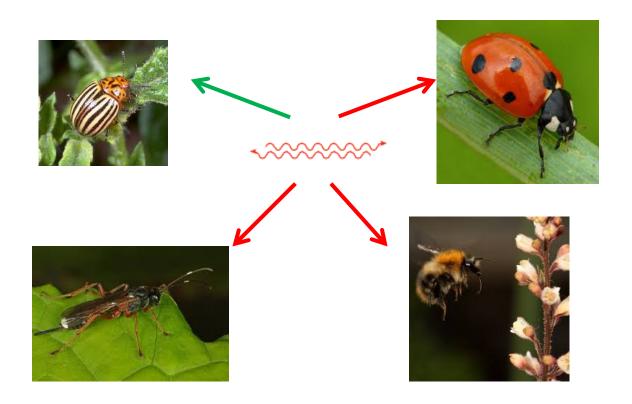
- Molecule ubiquitously present in nature
- Highly species specific in theory
- Persistency in the environment

Potential environmental risks associated with RNAi technology

- Gene silencing in non-target organisms
- Environmental fate of dsRNA
- RNAi machinery saturation?
- Immune stimulation?

Gene silencing in non-target organisms

- Knockdown of any gene in a non-target organism
- Due to sequence homology
- dsRNA design is important

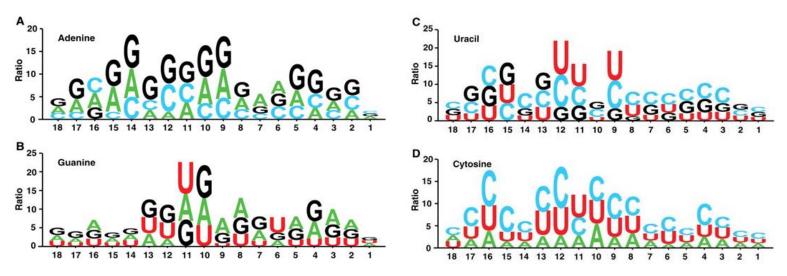


Gene silencing in non-target organisms

How specific is RNAi?

- No real consensus (rules for miRNA are clear, not for siRNA)
- Some mismatches could be allowed depending on:

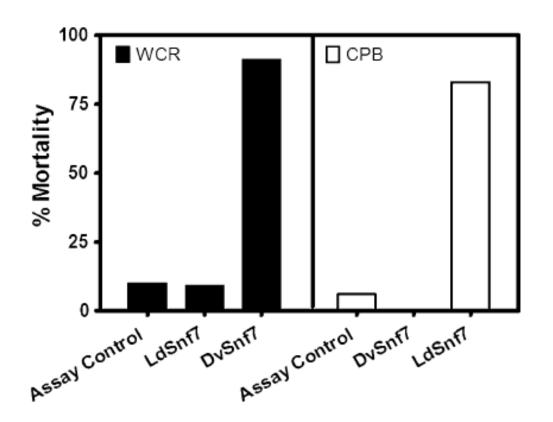
Location in the siRNA: seed region vs non-seed region Type of mismatch



Huang et al., 2009

Gene silencing in non-target organisms

How specific is RNAi?



Bachman et al., 2013 – Transgenic Res

Gene silencing in non-target organisms

How specific is RNAi?

Order	Family	Subfamily	Species	Bioassay duration (days)	Endpoints	LC ^a ₅₀ or no observed effect concentration ^b (ng/mL or g diet)
Coleoptera	Chrysomelidae	Galerucinae	D. virgifera virgifera	12	Survival	1.2 ^a
	Chrysomelidae	Galerucinae	D. undecimpunctata howardi	12	Survival	4.4 ^a
	Chrysomelidae	Chrysomelinae	L. decemlineata	12	Survival, Growth	5,000 ^b
	Tenebrionidae	Tenebrioninae	T. castaneum	30	Survival, Growth	5,000 ^b
	Coccinellidae	Coccinellinae	C. maculata	24	Survival, Growth, Development	3,000 ^b
	Coccinellidae	Epilachninae	E. varivestis	28	Survival, Growth, Development	3,000 ^b
	Carabidae	Harpalinae	P. chalcites	35	Survival, Growth, Development	5,000 ^b
Hemiptera	Anthocoridae	Anthocorinae	O. insidiosus	9	Survival, Growth, Development	5,000 ^b
Hymenoptera	Eulophidae	Entedoninae	P. foveolatus	21	Survival	3,000 ^b
	Pteromalidae	Pteromalinae	N. vitripennis	20	Survival	5,000 ^b
Lepidoptera	Noctuidae	Noctuinae	S. frugiperda	8	Survival, Growth	500 ^b
		Heliothinae	H. zea	12	Survival, Growth	5,000 ^b
	Crambidae	Pyraustinae	O. nubilalis	12	Survival, Growth	5,000 ^b
	Bombycidae	Bombycinae	B. mori	14	Survival, Growth	5,000 ^b

Bachman et al., 2013 – Transgenic Res

Gene silencing in non-target organisms

How specific is RNAi?

 Table 2
 Percent identity, numbers of single nucleotide polymorphisms (SNPs) and number of 21 nt matches of Snf7 orthologs from ten Coleoptera in the families Chrysomelidae and Tenebrionidae

Species	Subfamily, Tribe	Percent identity to DvSnf7 dsRNA	No. SNPs	No. 21 nt matches (or longest contiguous sequence)	
D. virgifera virgifera	Galerucinae, Luperini	100	0	221	
D. undecimpunctata howardi	Galerucinae, Luperini	98.8	3	186	
A. vittatum	Galerucinae, Luperini	95.0	12	69	
C. trifurcata	Galerucinae, Luperini	90.8	22	18	
G. calamriensis	Galerucinae, Galerucini	90.8	22	3	
A. lacertosa	Galerucinae, Alticini	81.7	44	0, (17 nt)	
C. quadrigemina	Chrysomelinae, Chrysomelini	82.1	43	0, (19 nt)	
M. ochroloma	Chrysomelinae, Chrysomelini	79.6	49	0, (19 nt)	
L. decemlineata	Chrysomelinae, Chrysomelini	78.3	52	0, (14 nt)	
T. castaneum	Tenebrioninae, Tribolini	72.1	67	0, (11nt)	

The inclusion of a representative of the Tenebrionidae is to provide context for the degree of sequence divergence outside the Chrysomelidae

Bachman et al., 2013 – Transgenic Res

Gene silencing in non-target organisms

How specific is RNAi?

Species from which dsRNA was synthesized	dsRNA concentration fed to WCR (ng/mL)	Control survival %	dsRNA treatment survival %	Statistical significance	Activity	21 nt matches
A. vittatum	1,000	90	8	<i>p</i> < 0.05	Active	Yes
C. trifurcata	500	76	15	p < 0.05	Active	Yes
G. calamariensis	5,000	89	13	p < 0.05	Active	Yes
M. ochroloma	500	76	72	p > 0.05	Not Active	No
C. quadrigemina	5,000	94	88	p > 0.05	Not Active	No

Table 3 Summary of results from heterospecific Snf7 dsRNAs fed to WCR in 12-day continuous exposure bioassays

The concentrations of heterospecific dsRNAs were prepared at 100–1,000 times the conspecific dsRNA 12-day LC_{50} for WCR (as reported in Bolognesi et al. 2012)

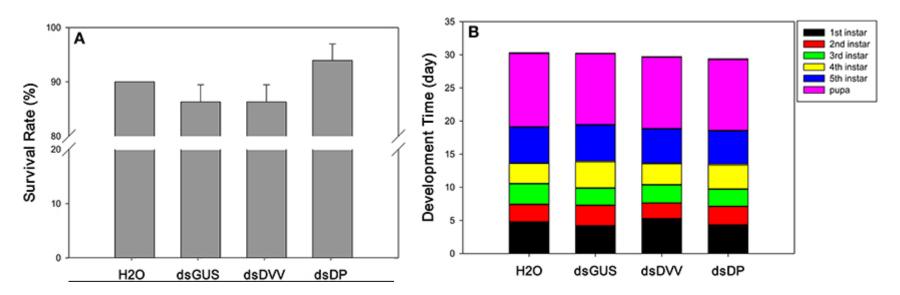
ORIGINAL RESEARCH ARTICLE

Front. Plant Sci., 22 February 2017 | https://doi.org/10.3389/fpls.2017.00242

Dietary Risk Assessment of *v-ATPase A* dsRNAs on Monarch Butterfly Larvae

Luipeng Pan^{1,2}, Xiaowei Yang², Keith Bidne³, Richard L. Hellmich³, Blair D. Siegfried⁴ and W Xuguo Zhou^{2*}

- Worst case scenario: Very conserved gene
 - Very conserved region
 - 1600 x Diabrotica virgifera LC50



ORIGINAL RESEARCH ARTICLE

Front. Plant Sci., 22 February 2017 | https://doi.org/10.3389/fpls.2017.00242

Dietary Risk Assessment of *v-ATPase A* dsRNAs on Monarch Butterfly Larvae

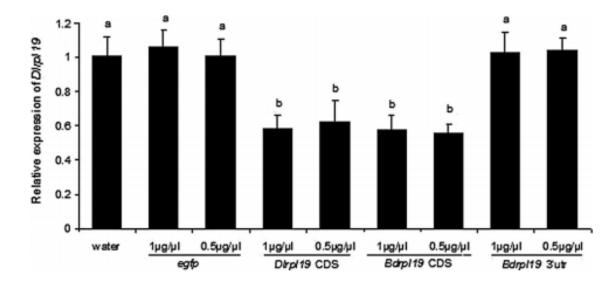
L Huipeng Pan^{1,2}, L Xiaowei Yang², Keith Bidne³, Richard L. Hellmich³, Blair D. Siegfried⁴ and Xuguo Zhou^{2*}

- Worst case scenario: Very conserved gene
 - Very conserved region
 - 1600 x Diabrotica virgifera LC50
- RNAi efficiency in Lepidoptera is a factor here

Gene silencing in non-target organisms

How specific is RNAi?

Fig. 4 Expression levels of Dlrpl19 after feeding D. longicaudata Dlrpl19 CDS dsRNA, Bdrpl19 CDS dsRNA or Bdrpl19 3'UTR dsRNA. Feeding D. longicaudata Bdrpl19 CDS dsRNA or Dlrpl19 CDS dsRNA significantly decreased Dlrp19 gene expression when compared to adults fed Bdrpl19 3'UTR dsRNA, egfp dsRNA or RNAfree water after 24 h



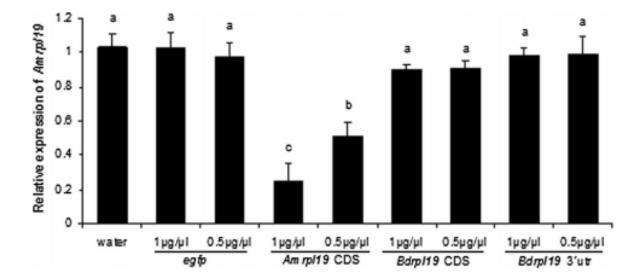
The effects of RNA interference targeting *Bactrocera dorsalis* ds-*Bdrpl19* on the gene expression of *rpl19* in non-target insects

Aie Chen · Weiwei Zheng · Wenping Zheng · Hongyu Zhang

Gene silencing in non-target organisms

How specific is RNAi?

Fig. 5 Expression levels of Amrpl19 after feeding A. mellifera Amrpl19 CDS dsRNA, Bdrpl19 CDS dsRNA or Bdrpl19 3'UTR dsRNA. Amrpl19 CDS dsRNA significantly decreased the Amrpl19 level when compared with Bdrpl19 CDS dsRNA, Bdrpl19 3'UTR dsRNA, egfp dsRNA and RNA-free water after 24 h



The effects of RNA interference targeting *Bactrocera dorsalis* ds-*Bdrpl19* on the gene expression of *rpl19* in non-target insects

Aie Chen · Weiwei Zheng · Wenping Zheng · Hongyu Zhang

Bioinformatics useful in risk assessment?

• Knowledge on the ecosystems and species present

What insects are present in a certain crop? Will these insects be exposed to the dsRNA?

What kind of amount of dsRNA can we expect these

species to take in?



Explore this journal >

Open Access

External scientific report

Establishing a database of bio-ecological information on non-target arthropod species to support the environmental risk assessment of genetically modified crops in the EU

Michael Meissle, Fernando Álvarez-Alfageme, Louise A. Malone, Jörg Romeis

First published: 13 September 2012 Full publication history



View issue TOC Volume 9, Issue 9 September 2012 334E

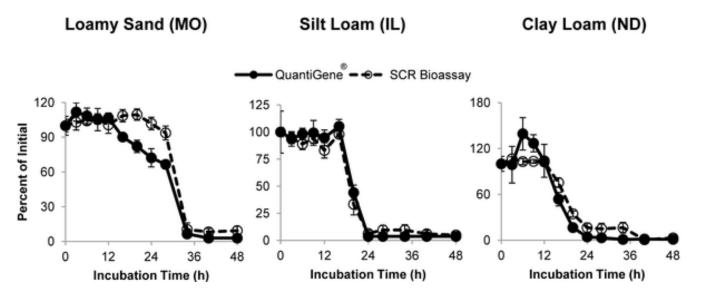
cts

Potential risks associated with RNAi-based (crop protection) technology

Environmental fate of dsRNA

Dubelman et al., 2014 - PLOS One

- Investigated the persistency of dsRNA in the soil
- 3 soil types



DsRNA persistency in soil + lyophilized corn tissue

Potential risks associated with RNAi-based (crop protection) technology

Saturation of RNAi machinery

- Number of RISC complexes is limited in a cell
- Uptake of large amounts can saturate RNAi machinery
- Sequence-independent
- Potential effects on efficiency of internal gene regulation?
- Potential effects on ability to fight off viral infections?

Potential risks associated with RNAi-based (crop protection) technology

Immune stimulation

- In mammals: long dsRNA fragments can incite an immune response when taken up into the cell
- Will this affect the fitness of other organisms as well?
- In insects: unknown
- More research necessary

Insects 2013, 4(1), 90-103; doi:10.3390/insects4010090

Open Access

Article

Non-Target Effects of Green Fluorescent Protein (GFP)-Derived Double-Stranded RNA (dsRNA-GFP) Used in Honey Bee RNA Interference (RNAi) Assays

Francis M. F. Nunes ^{1,†,*} , Aline C. Aleixo ^{1,†,*} , Angel R. Barchuk ² , Ana D. Bomtorin ¹ , Christina M. Grozinger ³ and Zilá L. P. Simões ⁴

Conclusions

- RNAi seems promising, compared to current pest control methods
- Bioinformatics: useful tool, but not enough
- DsRNA disappears quickly from the environment
- Gaps in our knowledge
- Level of exposure to siRNAs and dsRNAs must be carefully considered
- Each specific event will have to be tested for effects in non-target organisms:
 - Effects on plant
 - Effects on agroecosystem
 - Effects on humans after ingestion of specific siRNA/dsRNA

2014: RNAi workshop in Brussels organized by EFSA

- Experts, regulatory bodies, industry and NGOs were invited
- Three break-out sessions:
 - 1) Molecular characterization
 - 2) Food/feed risk assessment
 - 3) Environmental risk assessment
- Report: http://onlinelibrary.wiley.com/doi/10.2903/sp.efsa.2014.EN-705/epdf

2015: Calls for Tenders

EXTERNAL SCIENTIFIC REPORT

APPROVED: 16 may 2017

doi:10.2903/sp.efsa.2017.EN-1246

Literature review of baseline information to support the risk assessment of RNAi-based GM plants

Jan Paces¹, Miloslav Nic², Tomas Novotny², Petr Svoboda¹

¹ Institute of Molecular Genetics of the Academy of Sciences of the Czech Republic (IMG)





EFSA Tender on Environmental Safety

Contract – OC/EFSA/GMO/2015/02 - Literature review of baseline information on RNAi that could support the food/feed and environmental risk assessment of RNAibased GM plants;



Lot 2 – Literature review of scientific information on RNAi to support the environmental risk assessment of RNAi-based GM plants" (EFSA-Q-2016-00329)

- Goal: Gather knowledge on RNAi in invertebrate species
- Consortium for the ERA lot:
 - UGENT (Dr. Olivier Christiaens and Prof. Guy Smagghe)
 - ENEA (Prof. Salvatore Arpaia and Dr. Isabella Urru)
 - ABI (Prof. Kaloyan Kostov and Dr. Teodora Dzhambazova)
 - JT Environmental Consultants (Dr. Jeremy Sweet)

EFSA Tender on Environmental Safety

- Systematic literature search (+/- 13.500 papers retrieved)
- Review the available literature
 - Cellular uptake mechanisms in invertebrates
 - RNAi efficiency
 - Risks for non-target organisms
 - Exposure routes
 - Available genomic data
- Expected publication of ERA lot: early 2018



Further references

- EPA White Paper on RNAi technology as a pesticide (2013)
- Roberts AF, Devos Y, Zhou X, Lemgo G (2015) Biosafety research for nontarget organism risk assessment of RNAi-based GE plants. *Frontiers in Plant Science*, 6:958,doi:10.3389/fpls.2015.00958
- Casacuberta JM, Devos Y, du Jardin P, Ramon M, Vaucheret H, Nogué F (2015) Biotechnological uses of RNA interference in plants: risk assessment considerations. *Trends in Biotechnology*, 33: 145-147
- Ramon M, Devos Y, Lanzoni A, Liu Y, Gomes A, Gennaro A, Waigmann E (2014) RNAi-based GM plants: food for thought for risk assessors. *Plant Biotechnology Journal*, 12: 1271-1273
- Lundgren JG, Duan, JJ (2013). RNAi-based insecticidal crops: potential effects on nontarget species. Bioscience 63:657-665

Thank you for your attention!

Dr. Ir. Olivier Christiaens Lab for Agrozoology Ghent University – Belgium olchrist.christiaens@ugent.be

