

RNA-Based Plant Protection

Host-Induced Gene Silencing to Engineer Resistance to FHB

Karl-Heinz Kogel
Phytopathologie, Gießen

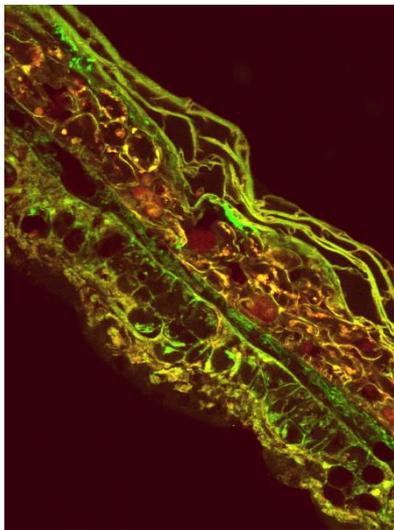
RNA-Based Plant Protection

Is there an agronomic potential for application ?

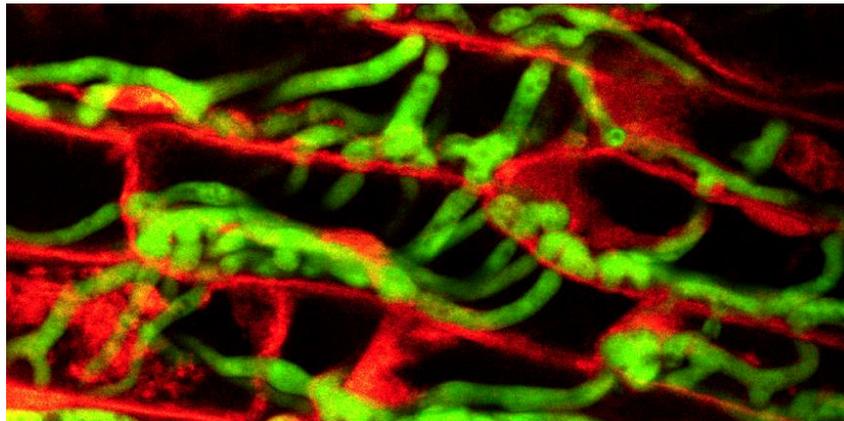
The Plant Pathogen *Fusarium graminearum*



Macroconidia



Toxin syndrome



Necrotrophic growth



Head blight disease

Fusarium species threaten harvest and food safety



Fusarium Head Blight

Toxicity

Zearalenon LD₅₀

7 mg kg⁻¹ (body weight, mouse oral)

Copper treatments LD₅₀

500 - 2000 mg kg⁻¹ (mouse oral)

660 mg kg⁻¹ (birds)

0.052 mg l⁻¹ (fish)

Modern pesticide LD₅₀

>5000 mg kg⁻¹ (mouse oral)

RNA-based crop protection exploits RNA interference

Different mechanisms of RNA interference (gene silencing)

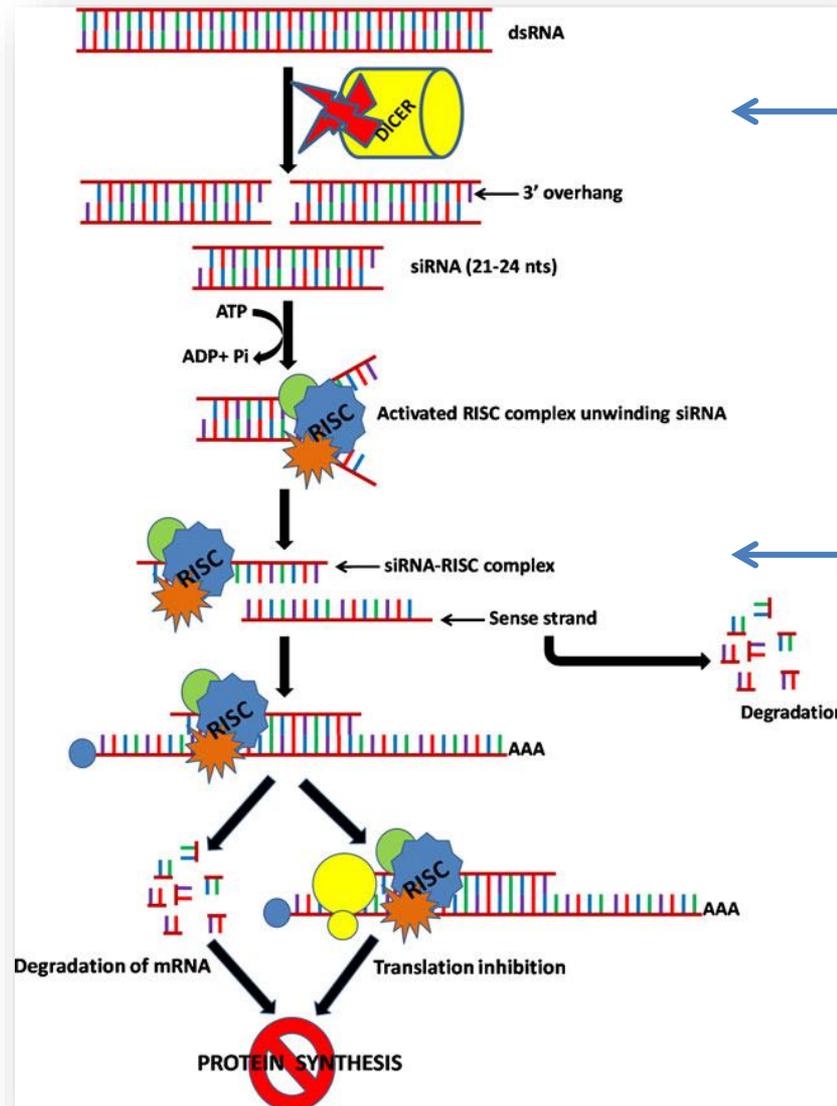
transcriptional gene silencing
(TGS)

post-transcriptional gene silencing
(PTGS)



Post-transcriptional gene silencing

Andrew Z. Fire and Craig C. Mello (Noble Prize 2006)



Key enzymes:

DICER

RNase III –type endonuclease
4 DICER-like proteins in Arabidopsis

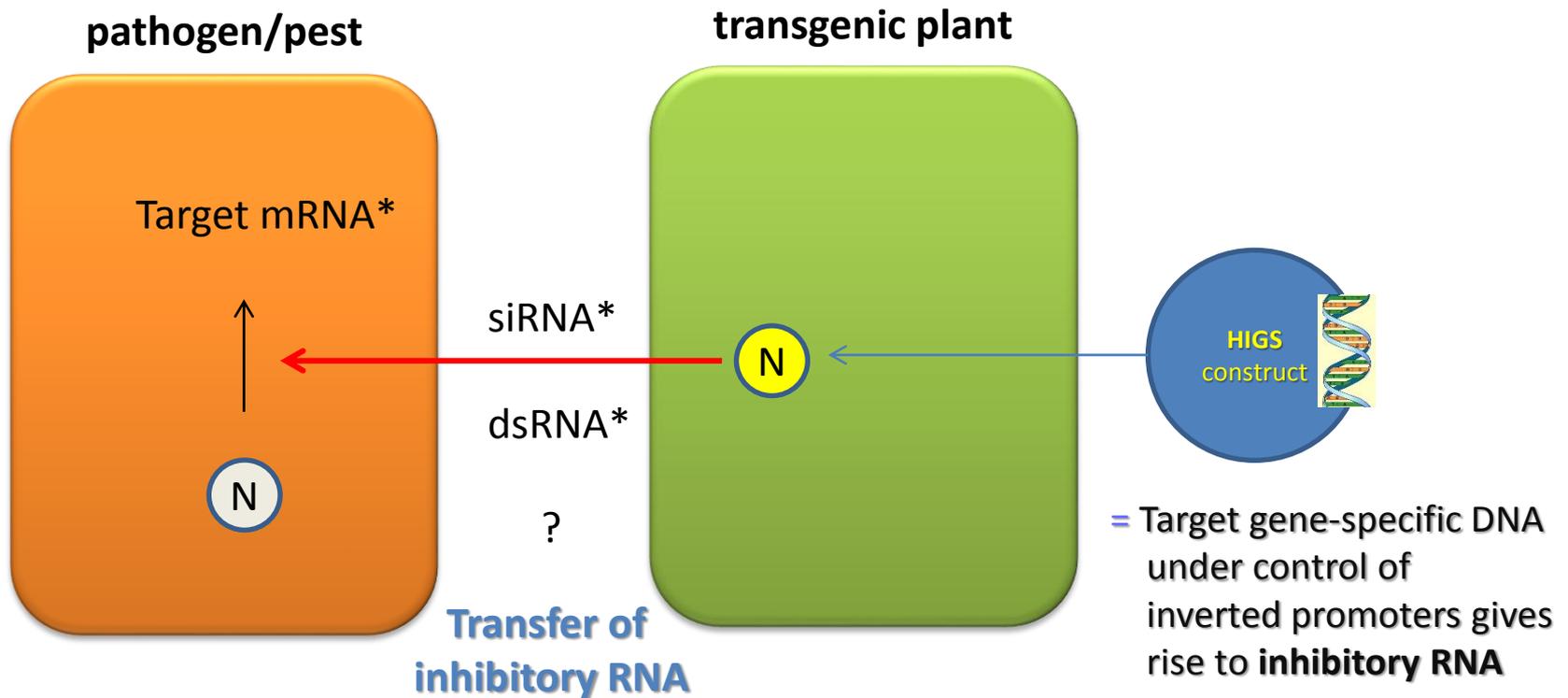
Argonaute

RNase III –type endonuclease
10 Argonautes in Arabidopsis

Image from Jagtap et al. 2011

Host-Induced Gene Silencing

oversimplified model



*Lethal target gene

siRNA: DICER-released small interfering RNA; ds: double stranded RNA; N: nucleus

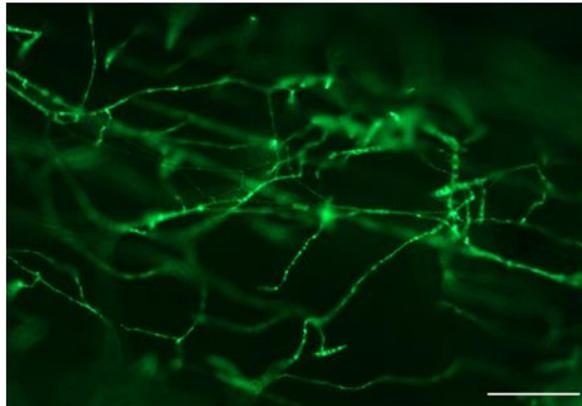
Proof-of-Concept

dsRNA-GFP plants inoculated with δ GFP-tagged *Fusarium graminearum*

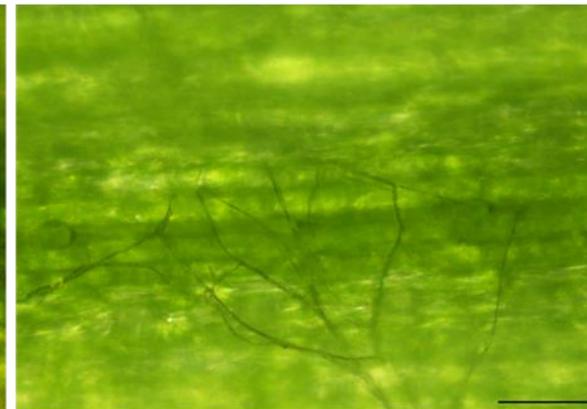
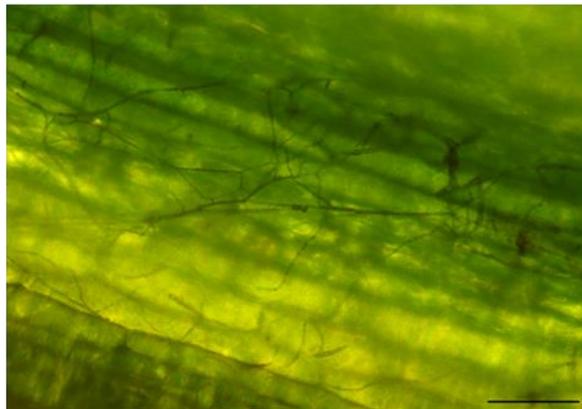
Control*

dsGFP-RNA
plants

excitation
395 nm



visible light



*Golden Promise inoculated with Fg-GFP

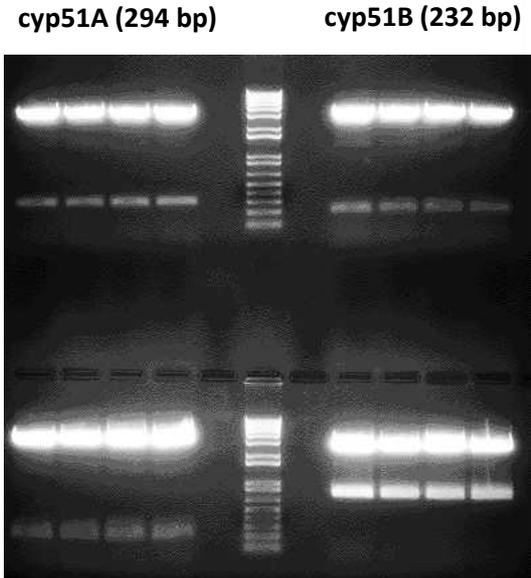
δ GFP = Green Fluorescence Protein

Host-Induced Gene Silencing

**Selection of the right target gene
is a critical success factor
of HIGS applications**

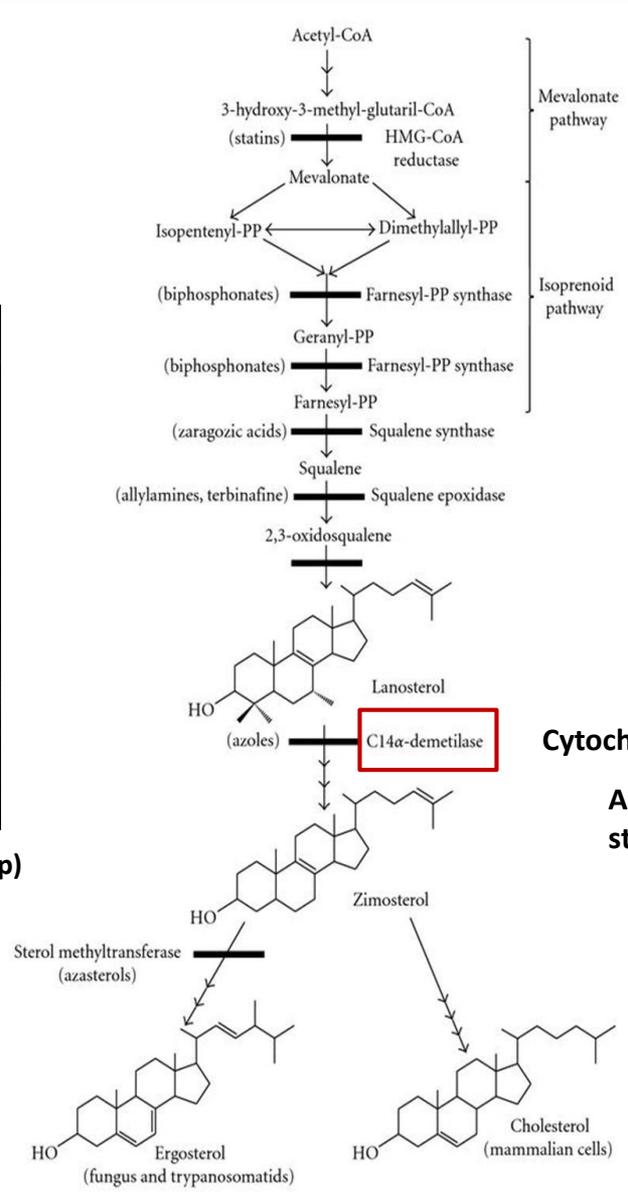
The target is a critical success factor for HIGS applications

Fusarium graminearum
has three *CYP51* genes



cyp51A (294 bp) **cyp51B (232 bp)**
cyp51C (250 bp) **cyp51ABC (791 bp)**

Gene fragments were amplified from genomic DNA with specific primers and cloned into pGEM-T easy vector



Ergosterol biosynthesis

Cytochrome P450 sterol 14 α -demethylase (CYP51)

**Azole fungicides =
sterol demethylation Inhibitors (DMI)**

The inhibitory dsRNA *CYP3RNA*

Clone sequences of *CYP51A* (294nt)

```
CGGTCCATTGACAATCCCCGCTCTTTGGTAGCGATGTCGTATACGATTGTCCCAACTCGAAGCTCATGGAACAAAAGAAGTTTGTCAAGTTTGGCCTTACGCAAAA  
AGCACTCGAGTCACACGTCCAGTTAATCGAGCGAGAGGTTCTTGACTACGTCGAAACTGATCCATCCTTTTCTGGCAGAAGTAGCACCATCGATGTCCCAAGGC  
AATGGCTGAGATAACAATCTTTACTGCCTCACGTTCTTTGCAGGGTGAGGAAGTTCGGAGAAAACACTACTGCCGAGTTTGCTGC
```

Clone sequences of *CYP51B* (220nt)

```
CAGCAAGTTTGACGAGTCCCTGGCCGCTCTCTACCACGACCTCGATATGGGCTTCACCCCCATCAACTTCATGCTTCACTGGGCCCCCTCTCCCCTGGAACCGTA  
AGCGCGACCACGCCAGCGCACTGTTGCCAAGATCTACATGGACACTATCAAGGAGCGCCGCGCCAAGGGCAACAACGAATCCGAGCATGACATGATGAAGCA  
CCTTATGAACTCT
```

Clone sequences of *CYP51C* (238nt)

```
ATTGGAAGCACCGTACAATATGGCATCGACCCGTACGCTTTTTTCTTCGACTGCAGAGATAAATACGGCGACTGCTTTACCTTTATTCTCCTTGGCCAAATCAACGA  
CTGTCTTTCTTGGTCCCAAGGGCAATGACTTTATCCTCAACGGCAAACACGCCGATCTCAACGCCGAGGACGTTTATGGGAAACTTACCACGCCCGTGTGGTG  
AGGAGGTTGTTTATGACTGCTCCAATG
```



****CYP3RNA***

dsRNA: *CYP3RNA*

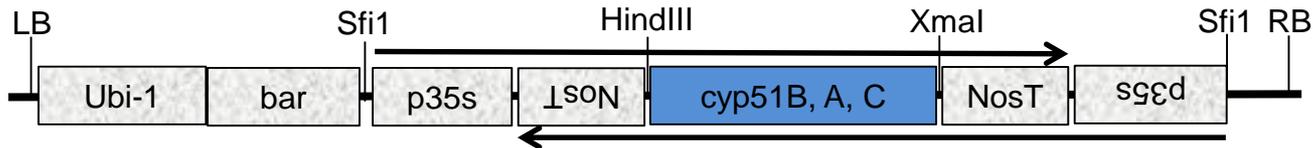


buffer



In planta experiment

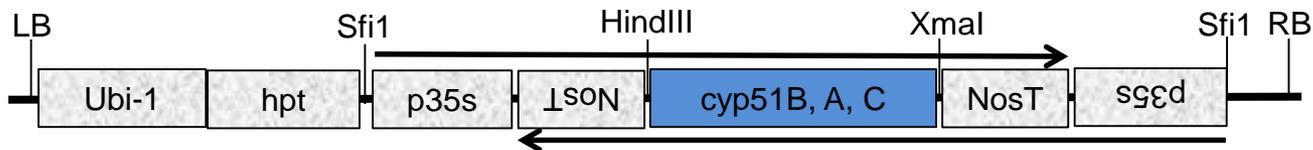
Vector for plant transformation



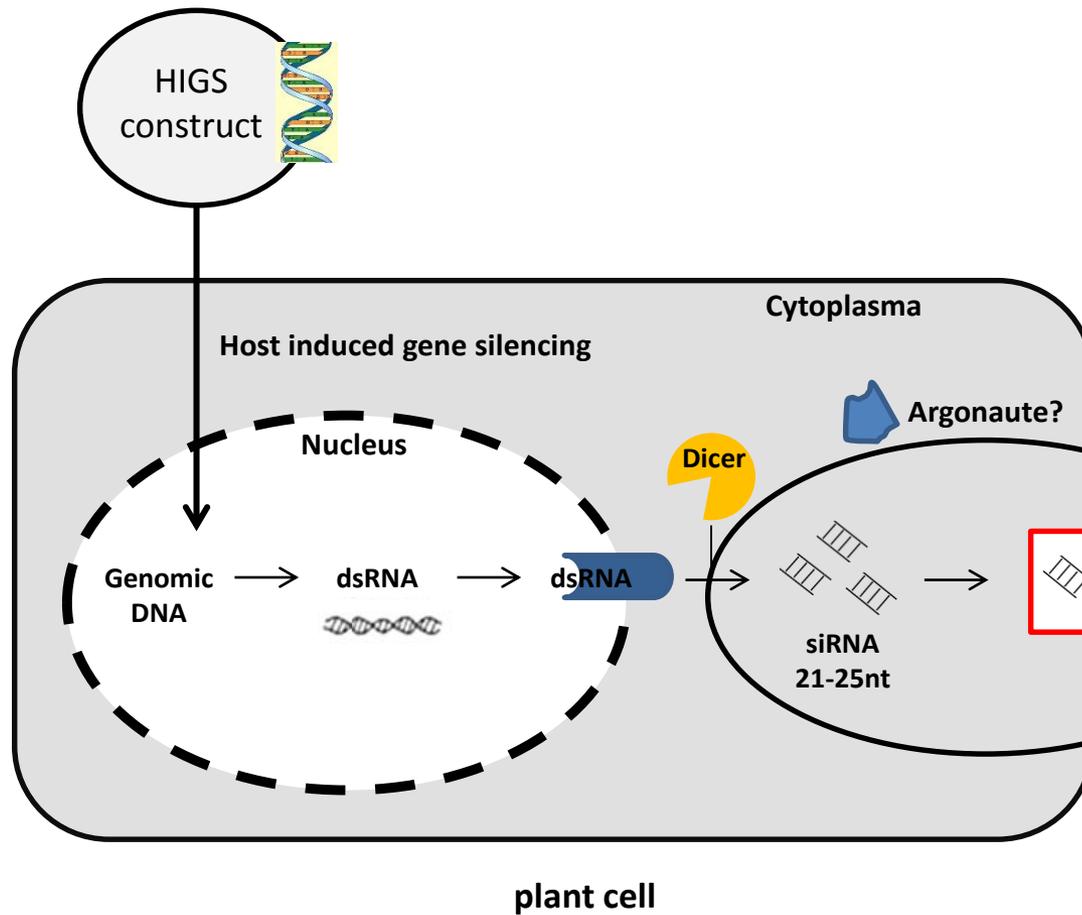
Arabidopsis



Barley

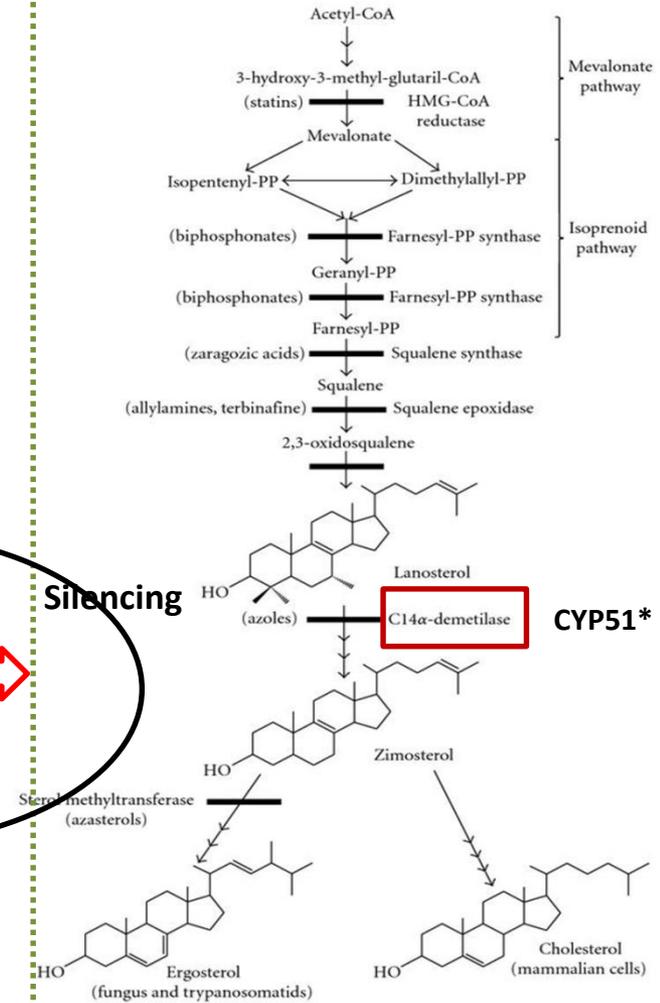


CYP3RNA processing



fungus cell

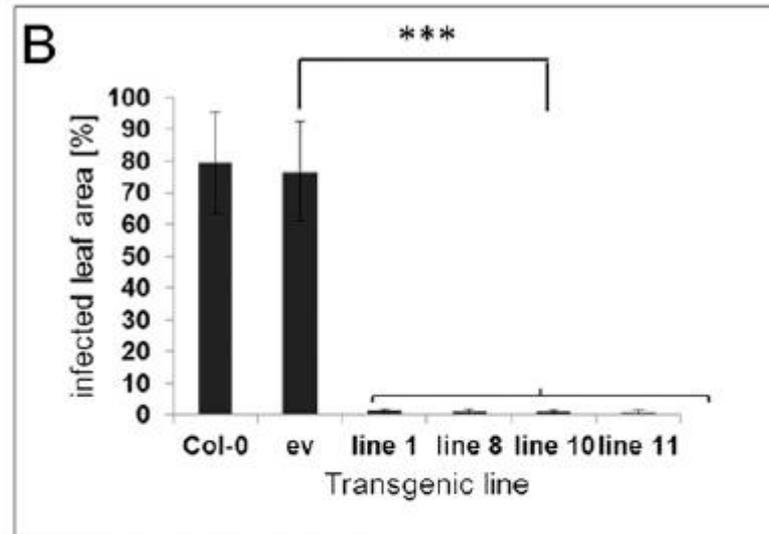
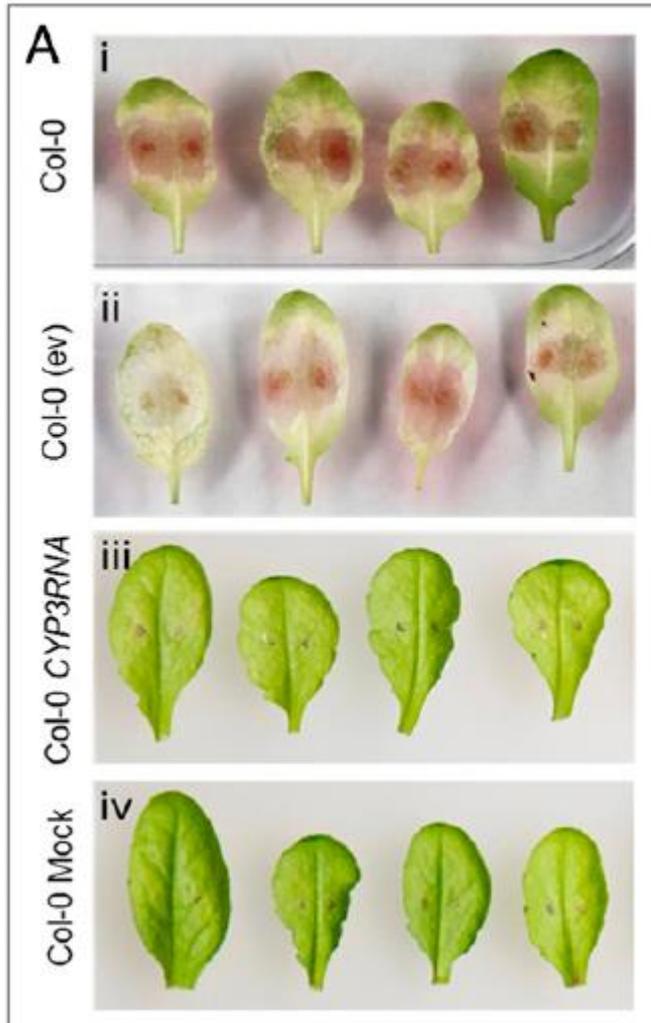
Ergosterol biosynthesis



* Cytochrome P450 Sterol 14 α -Demethylase

CYP3RNA expression inhibits infection

Arabidopsis

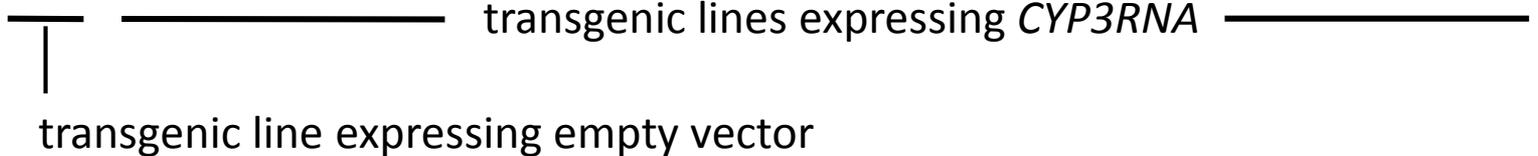


CYP3RNA expression inhibits infection

Barley



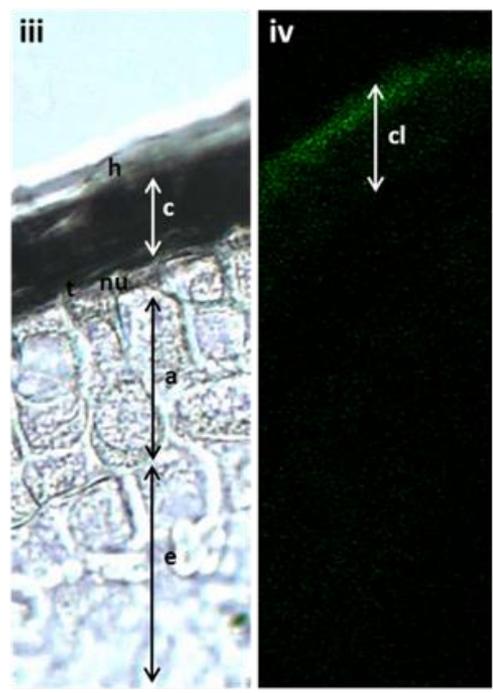
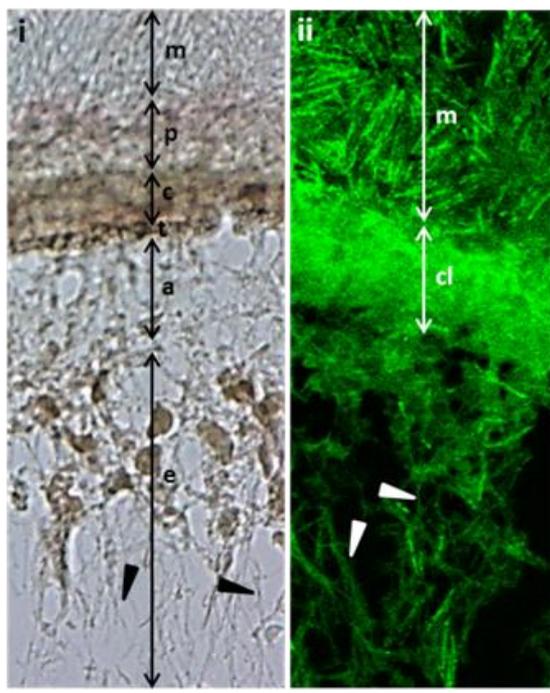
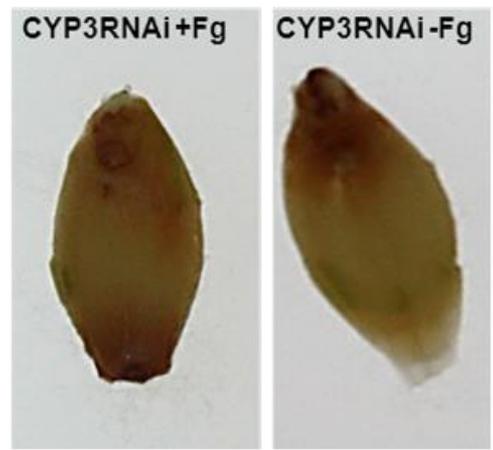
Wild
type



wt = cv. Golden Promise

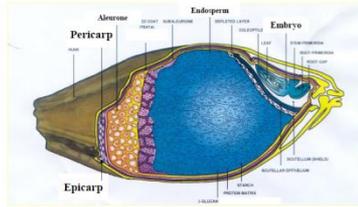
Strong inhibition of Fusarium Head Blight

Barley



Control

CYP3RNA



Strong silencing of fungal *CYP51* expression *in planta*

Barley

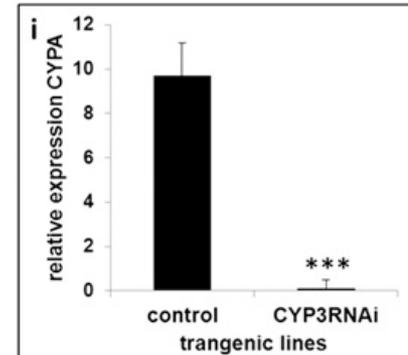
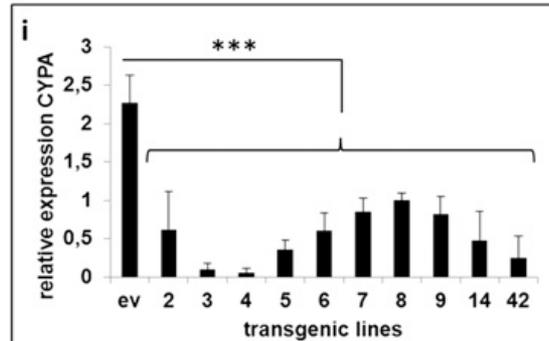
F. usarium infected



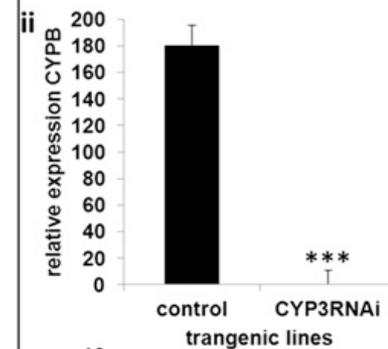
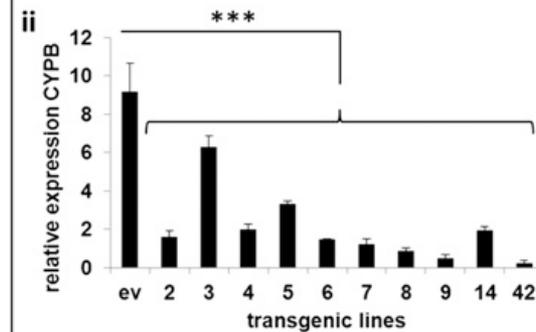
Leaves

Roots

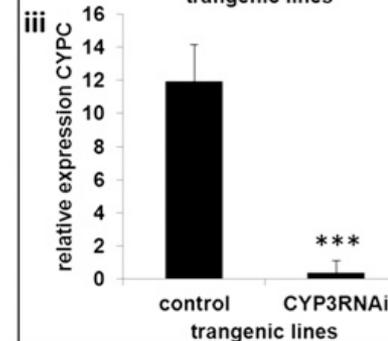
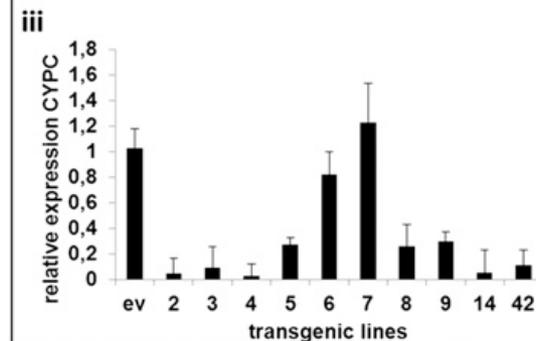
CY51A
expression



CY51B
expression



CY51C
expression



Normalized with fungal β -tubulin

Off-target analysis

Table S1. Prediction of CYP3RNA off-target transcripts

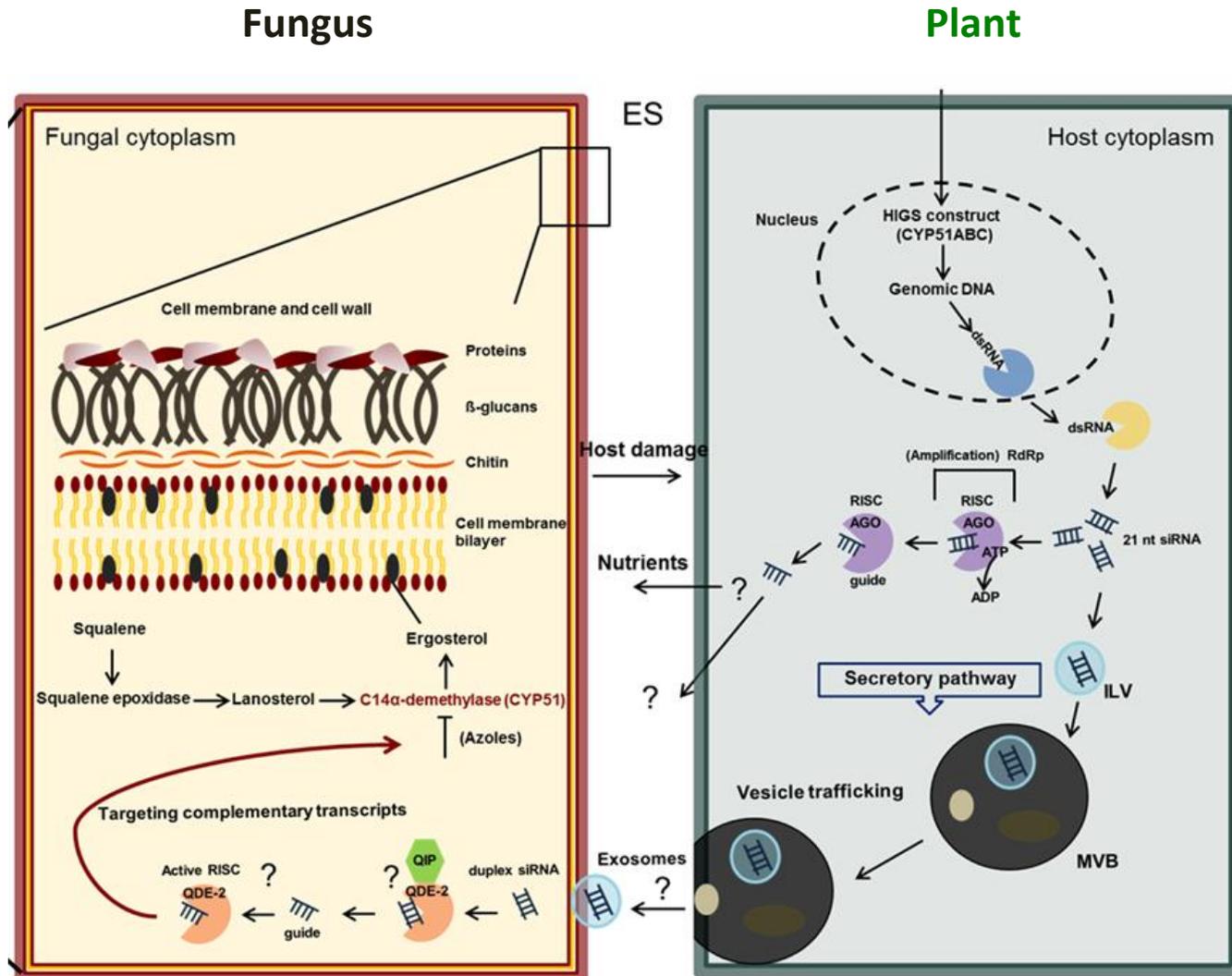
Organism	Query name*		All hits [†]	Efficient hits [‡]
	Gene	Description		
<i>Fusarium graminearum</i>	FGSG_01000 [§]	<i>CYP51B</i>	200	46
	FGSG_04092 [§]	<i>CYP51A</i>	274	126
	FGSG_11024 [§]	<i>CYP51C</i>	218	95
<i>Arabidopsis thaliana</i>	AT2G17330	<i>CYP51A1</i>	0	0
	AT1G11680	<i>CYP51A2</i>	0	0
<i>Hordeum vulgare</i>	Published database (1)		0	0
<i>Hyaloperonospora arabidopsidis</i>	Published database (2)		0	0
<i>Rhizophagus irregularis</i>	Published database (3)		0	0
<i>Piriformospora indica</i>	Published database (4)		0	0
<i>Homo sapiens</i>	Published database (5)		0	0
<i>Fusarium cerealis</i> isolate NRRL13721	JN416614 [¶]	<i>CYP51A</i>	190	83
<i>Fusarium austroamericanum</i> isolate NRRL28718	JN416607 [¶]	<i>CYP51A</i>	117	50
<i>Fusarium vorosii</i> isolate 67C1	JN416608 [¶]	<i>CYP51A</i>	116	50
<i>Fusarium acaciae-mearnsii</i> isolate NRRL26752	JN416603 [¶]	<i>CYP51A</i>	94	43

*Simulations were run using Si-Fi software (v3.1) for predicting off-targets prediction (<http://labtools.ipk-gatersleben.de>).

†Number of 21-mer siRNA sequences with perfect match to the query sequence.

‡Number of 21-mer siRNA sequences with perfect match to the query sequence that fulfill additional criteria for efficient RNAi (See Si-Fi software).

What type of inhibitory RNA is transferred ?

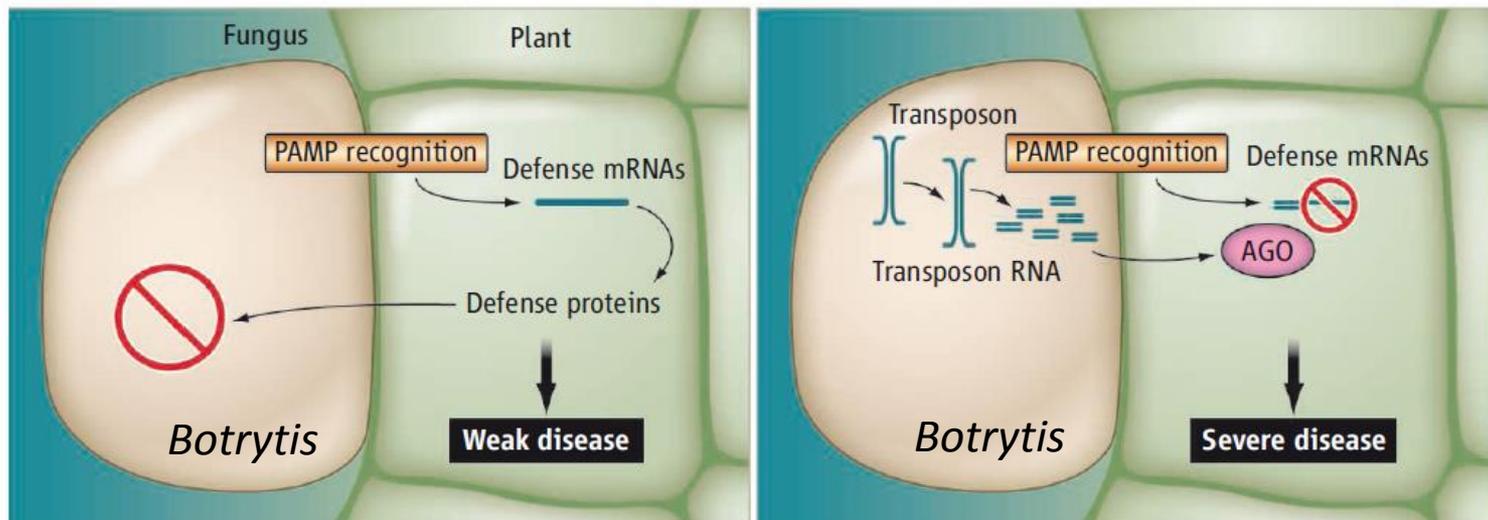


ILV, intraluminal vesicles, MVB, multivesicular bodies

Outlook – HIGS amenable to plant breeding ?

No example has been found so far showing that a crop produces small RNAs to target its pathogen/pest

However: *Botrytis cinerea* targets plant defense genes by small RNAs



From: Baulcombé 2013
Comments on Weiberg et al. 2013

It is too early to speculate whether breeding approaches on these plant targets could be a realistic strategy.

Acknowledgments



Dr. Aline Koch



Dr. Jafar Imani
transgenic barley



M.Sc. Eltayb Abdellatef
RNAi mutants



Dagmar Biedenkopf
Technical assistance

Excellent effects of HIGS against the grain aphid *Sitobion avenae**



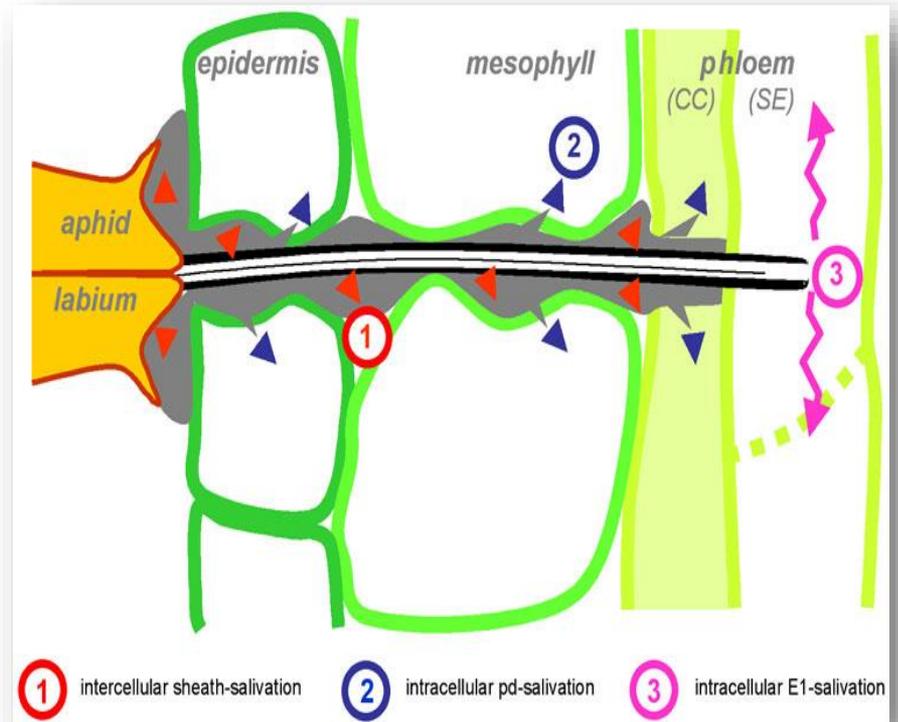
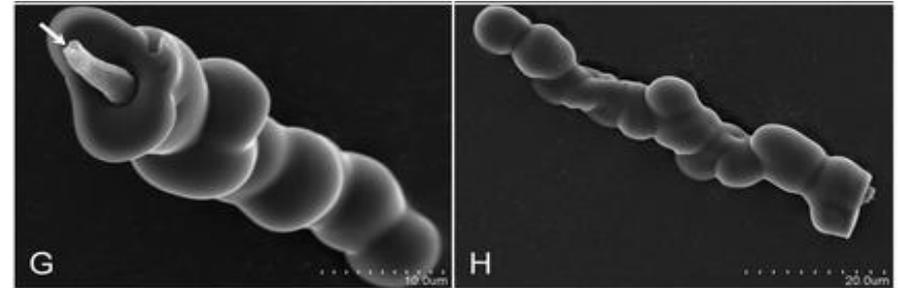
Eltayb Abdellatef



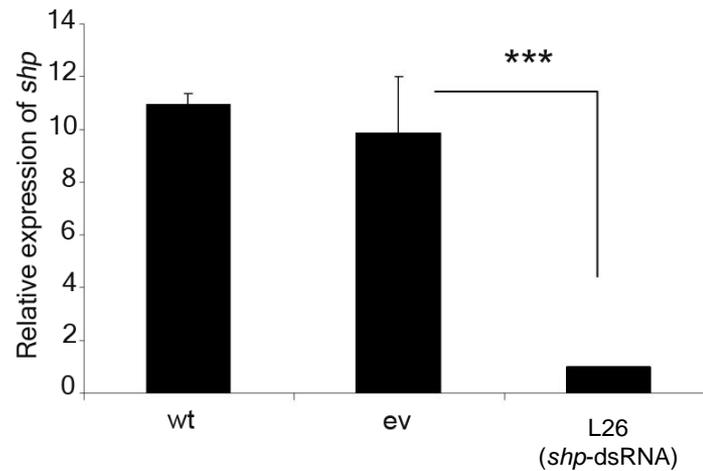
*Collaboration with A. Vilcinskis and T. Will, Inst. f. Entomology, JLU Gießen

Silencing of Salivary Sheath Protein SHP in *S. avenae*

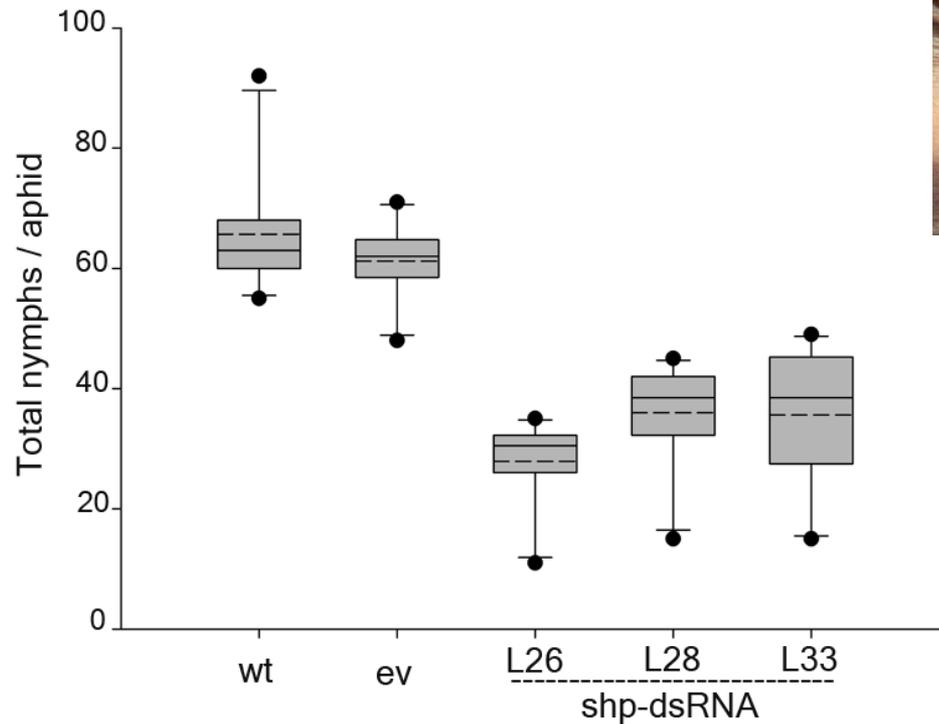
- Aphids feed on sap suck from sieve tube of vascular plants. During this process aphids secrete gel saliva that forms a sheath to enclose the stylet.
- The stylet sheath is built up by different proteins, though SHP seem essential because it forms the structural backbone of the sheath.



Reduced expression of *shp* in aphids fed on transgenic barley



Reproduction rate, growth development, and survival rate was negatively affected



Silencing of *shp* is transmitted transgenerationally

