ERWINIA OVERCOMES POTATO RESISTANCE BY ATTACKING ITS DEFENCES

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INTRODUCTION

Characterisation of geneX mutant and Real-Time PCR

The role of geneX in virulence was

determined by identifying a Tn5

insertion mutant in this gene. Thereaf-

ter, pathogenicity tests were carried

out on both potato tubers and stems to

compare virulence of *geneX* mutant vs

showed reduced virulence of mutant. GeneX mutant was also complemented

with plasmid carrying geneX and

complementation restored virulence of

geneX is required for virulence

Relative expression levels of TF1 in

leaves infiltrated with Eca or

aeneX mutant

in Eca

mutant to Eca wt levels.

400

s 350 300

200

150 relative

100

50

0

0.5

sion 250

ess

expr

Eca wild type (wt) strain. Results

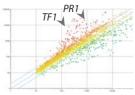
Erwinia carotovora subsp. atroseptica (Eca) is an economically important pathogen of potato, causing tuber soft rot and blackleg. There are no chemical treatments to control Eca in the field. Morever, there are no commercial cultivars with total resistance to Eca. Understanding mechanisms used by *Eca* during infection is crucial in the development of resistant cultivars.

The main weapons used by Eca to infect plants are plant cell wall degrading enzymes. However, the recently identified type III secretion system (T3SS) in *Eca* is also instrumental in injecting effector proteins into the host plant to suppress, manipulate or modulate defences during early stages of infection (Toth and Birch, 2005). The aims of this project were:

- To identify a mutant in geneX, a putative effector, and to assess its role in pathogenicity.
- To identify potato defence pathways modified by this putative effector.
- To use this knowledge for enhanced resistance to Erwinia.

Microarrays

To determine which other genes are up-regulated together with TF1 transcription factor, cDNA from leaves infiltrated with Eca wt vs. geneX mutant 0.5 hours post inoculation (hpi) were hybridised to an Agilent microarray. PR1 gene was also shown to be up-regulated together with the TF1. PR1 is a marker of salicylic acid-dependant pathways.



Results suggest that *geneX* product suppresses SA-dependent potato defence mechanisms

Improved resistance

- •The above results suggest that constitutive expression of TF1 transcription factor can increase resistance to Erwinia.
- Transgenic lines with increased expression of the TF1 transcription factor were generated.
- Pathogenicity assays on Desiree vs. transgenic TF1 lines.
- Lesion measured from 2 to 17 days post inoculation (dpi).



Desiree control (17dpi)



Transgenic TF1 plant (17dpi)

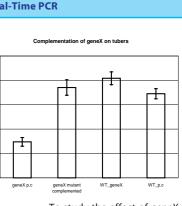
Overexpression of TF1 enhances resistance of of potato cultivar Desiree to Eca

References

Toth IK and Birch PBJ 2005 Rotting softly and stealthily. Current Opinions in Plant Biology 8(4):424-429

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To study the effect of geneX products on potato plants, the expression of transcription factor1 (TF1) was monitored using Real-Time PCR. Initially, low levels of TF1 were observed in leaves infiltrated with Eca wt while a massive increase of the transcription factor occured at 10hpi. On the contrary, TF1 was highly upregulated in leaves infiltrated with geneX mutant within 0.5hpi of inoculation.

This result suggests possible involvement of *geneX* product in suppression of TF1 during early hours of infection by Eca wt

10

geneX

■ WT

Improved resistance

2

4

Time (hpi)

Transgenic lines are significantly resistant to Erwinia compared to control plants