Development and production of novel fusion protein biopesticides Researching biological alternatives to chemical pesticides

INTRODUCTION

The use of common chemical pesticides particularly neonicotinoids poses unacceptable eco-agronomical risks due to declines in pollinator populations (Figure 1) and studies on pollinators such as honey (Henry et al, 2012) and bumble bees (Whitehorn et al, 2012) indicate declines in queen numbers, learning and memory defects, increases in numbers that don't return to the hive and ultimately colony-collapse disorder.

FIGURE 1: Economic significance of insect pollination on agricultural production of human food crops worldwide (UNEP 2010; Gallai et al 2009).

Crop category	Average value of a production unit	Economic value (10°€)									
	(€ per metric tonne)	50	100	150	200	250	300	350	400	450	
Vegetables	468									1	
Cereals	139										
Sugar crops	177		-		-						
Edible oil crops	385										
Fruits	452										
Roots and tubers	137										
Pulse	515										
Stimulant crops	1 225									· · ·	
Nuts	1 269						Total economic value				
Spices	1 003						_	lotal eco	nomic va		
All categories pooled toget	ther									16	

The contribution of pollinators to the production of crops used directly for human food has been estimated at €153 billion globally, which is about 9.5% of the total value of human food production worldwide ⁶.

The aim of this project was to expand on research by the Gatehouse group by developing novel GNA fused biopesticides derived from the SFI family of neurotoxic proteins from the venom of the tube web spider *Segestria* florentina (below), and express them in the yeast Pichia pastoris.



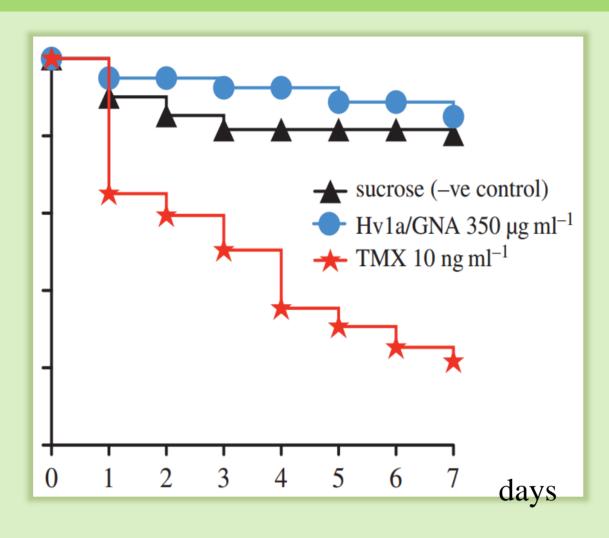
The proteins induce flaccid paralysis in insects and larvae, acting on voltage-dependent calcium (CaV) channels (Fitches et al, 2004).

The insecticidal protein is fused with snowdrop lectin (GNA -Gallanthus *nivalus* agglutinin) so that upon ingestion, the protein is absorbed through the mid-gut and into the insect haemolymph.

A similar biopesticide using ω -hexatoxin-Hv1a from the Australian funnel web spider *Hydronyche versuta* fused to GNA has been shown to have minimal adverse effects on honeybees following injection or chronic consumption (Figure 1; Nakasu et al, 2014) but also exhibits potent toxicity on a range of insect orders including Lepidoptera, Coleoptera, Diptera and Heliptera.

BIBLIOGRAPHY

uin M., Requier F., Rollin O., Odoux J-F., Aupinel P., Tchamitchian S., Decourtye A. 2012. "A Common Pesticide Decreases Foraging



AIMS & METHODS

Project A: Develop and produce biopesticides from the family of eight *S*. florentina insecticidal toxins (SFL1-8).

1. harvested.

2. Expression in *Pichia pastoris:* Digested recombinant pCR2.1 plasmid with restriction enzymes (Not1 and Xho1) \rightarrow Digested pGAPz α vector (yeast integrating plasmid) containing the GNA sequence with same restriction enzymes \rightarrow Toxin sequence and pGAPz α ligated together \rightarrow Transformed into *E.coli* and cloned \rightarrow Transformed into *P. pastoris* for expression of biopesticides.

Project B: Develop two novel synthetic SFI toxins by substituting amino acids from variable regions of one toxin with amino acids from the variable region of another in the same family.

The DNA encoding a synthetic toxin fused with GNA was synthesised by a two-step, 'templateless' PCR assembly of a series of overlapping oligonucleotides.

The toxin DNA is then transformed, cloned and expressed in much the same way as in Project A (above) with one exception: Since the synthetic sequence encoded GNA, the pCR2.1 and pGAPzα vector were restricted using Xba1 rather than Not1 when preparing to transform into *P. pastoris*.

Nakasu E.Y.T., Williamson S.M., Edwards M.G., Fitches E.C., Gatehouse J.A., Wright G.A., Gatehouse A.M.R. 2014. "Novel biopesticide based on a spider **336:** 351-352.

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FIGURE 2: Chronic consumption of Hv1a/GNA (21.7µg bee ⁻¹day⁻¹) does not affect honeybee survival.

Thiamethoxam/TMX – a neonicotinoid, (0.727ng bee ¹day⁻¹) significantly increases mortality.

(Taken from: Nakasu et al, 2014).

Cloning the toxin DNA: PCR Toxin DNA \rightarrow Ligated into pCR2.1 plasmid vector \rightarrow Transformed into *Escherichia coli* \rightarrow Blue-white selection to select positive transformants \rightarrow Positives grown overnight \rightarrow Plasmid

RESULTS & DISCUSSION

Project A

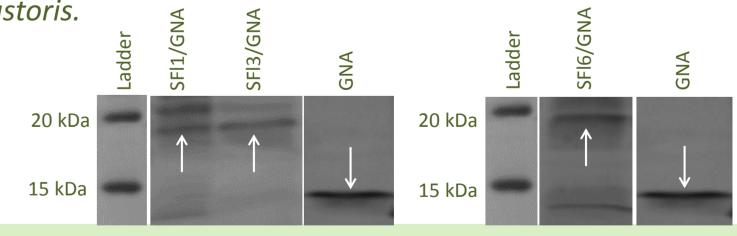
- assumption.

FIGURE 3: SFI1/GNA has acute toxicity on *Lacanobia oleracea* larvae 6 days after ingestion (Taken from: Fitches et al, 2004).

Project B

iGEM, 2009).

Pichia pastoris.

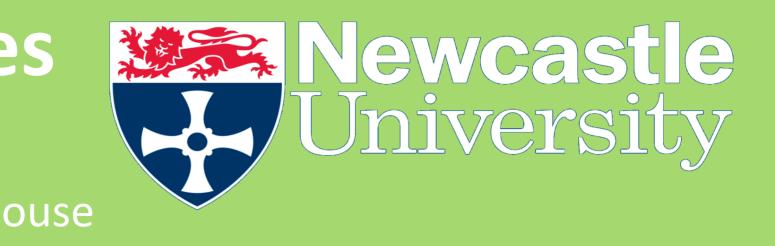


CONCLUSIONS / FURTHER STUDY

- toxin/GNA fusion proteins.

PICTURES

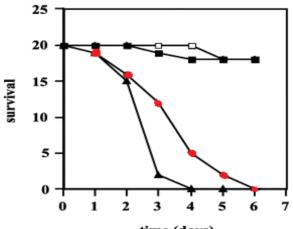
hitehorn P.R., O'Connor S., Wackers F.L., Goulson D. 2012."Neonicotinoid Pesticide Reduces Bumble Bee Colony Growth and Queen Production". Science

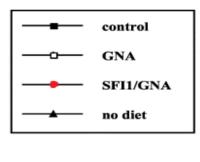


• Successfully cloned all eight SFI toxins in *E.coli* using the pCR2.1 plasmid vector, and transformed four of the toxins into the pGAPz α vector system.

• SDS-PAGE suggested we successfully expressed three different Segestritoxin/GNA fusion proteins in *Pichia pastoris;* SFI1/GNA, SFI3/GNA, and SFI6/GNA (See Figure 4). Further study required to confirm this

 Previous studies indicate SFI toxin/GNA fusion protein genes are potential candidates as biopesticides and show acute toxicity when fed to larvae of tomato moth, Lacanobia oleracea (Figure 3; Fitches et al, 2004).





• The templateless PCR yielded some promising results initially, however we were unable within the timeframe to create the synthetic SFI toxins; with more time we would have been successful since whole synthetic yeast chromosomes have been assembled using a similar protocol (Team JHU,

FIGURE 4: SDS-PAGE showing expression of three SFI/GNA biopesticides in

• New library of biopesticide candidates for expression in *P. pastoris* and testing for toxicity in important plant pests and pollinator species.

• Optimisation of transformation efficiency and protein expression required for improved productivity of fusion proteins.

• More time required to optimise the oligonucleotide assembly of synthetic

