



Using RNA interference as a biological pesticide for clubroot in canola

RNA Innovation Cohort 1
Clubroot Steering Committee Meeting
April 30, 2020



RNA  **Innovation**

The logo for RNA Innovation. The word "RNA" is in red, and "Innovation" is in black. A stylized black icon of a keyhole or a vertical bar with a circular top and a dashed vertical line inside connects the two words. A thick black horizontal line underlines the entire text.

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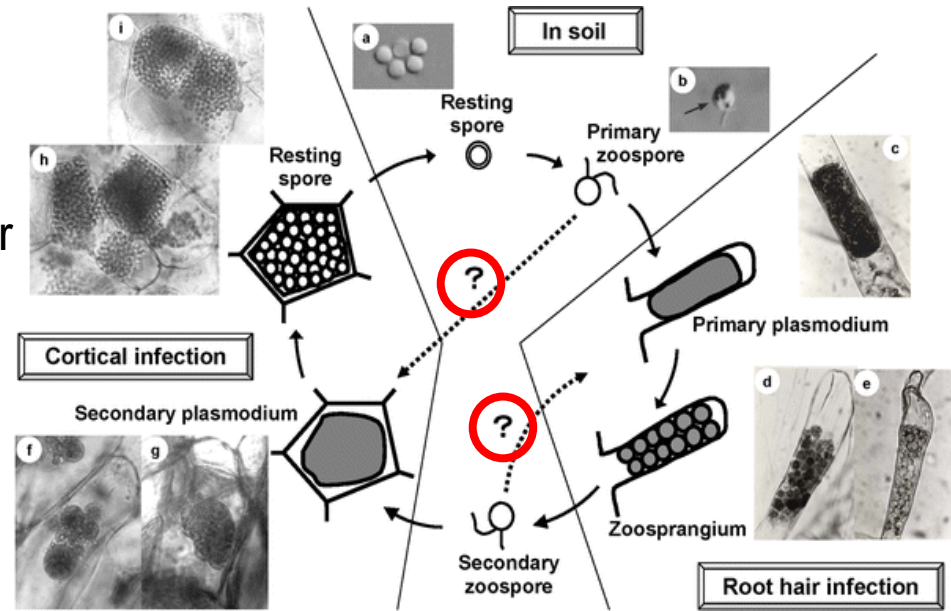
University of
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Bridging the Gap between Academia and Industry



Current challenges in clubroot research

- Limited laboratory cultivation methods.
- Lack of annotated clubroot genome.
- Difficulties with delivery mechanisms for RNAi.
- Unsolved mechanisms in clubroot life cycle.
- Regulatory challenges in marketing GMO products.



Objectives

Using RNAi to tackle clubroot

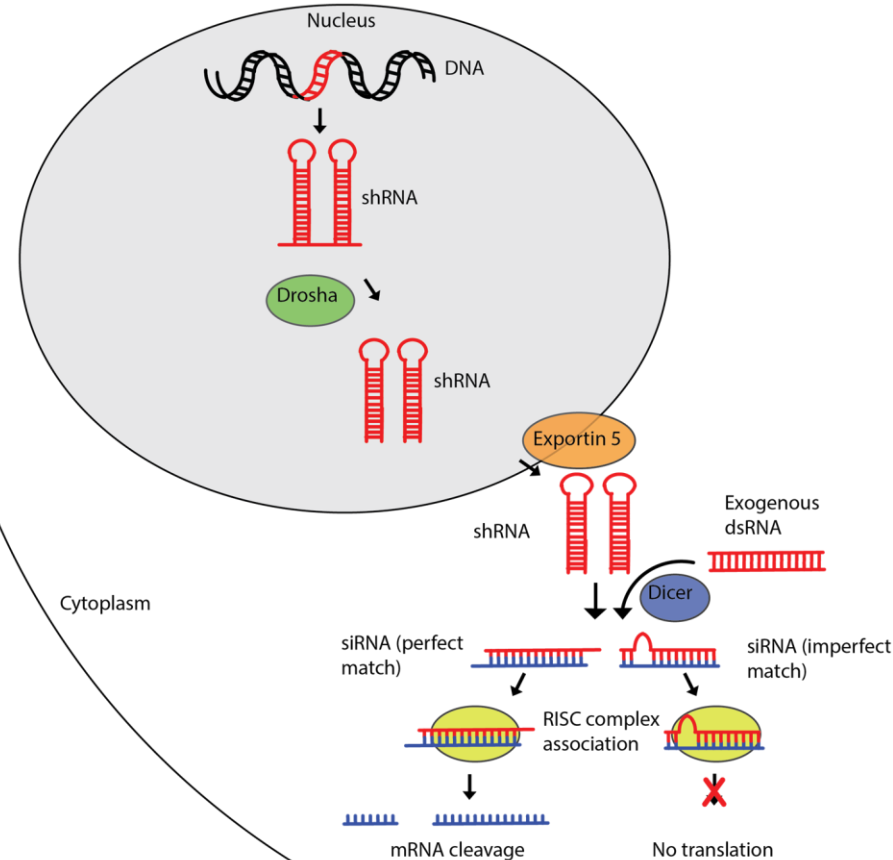
- To reduce the virulence or population of *Plasmodiophora brassicae* in the soil using RNA interference

Design siRNAs against target genes

- Designing and characterization of siRNAs against potential genes which are identified in the Clubroot genome.

Overview of RNAi

- Process used to inhibit gene expression
- Commercially available SmartStax[®] Pro seeds contain RNAi
- Minimal work reported on RNAi use for clubroot
- RISC complex is present in clubroot genome
 - Based on bioinformatics work conducted by our group



Screening of target genes using bioinformatics

- Selected 7 genes from *P. brassicae* as potential targets
- 2 genes are related to zoospore primary infection (Fei *et al.*, 2016).
- 5 genes are related to secondary infection of clubroot (Pérez-López *et al.* 2020).

SSPbP03



SSPbP53



SSPbP94



SSPbP22

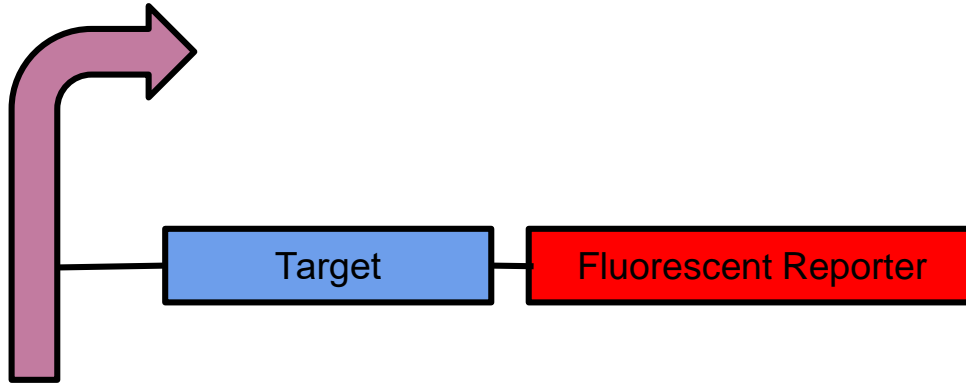


SSPbP02



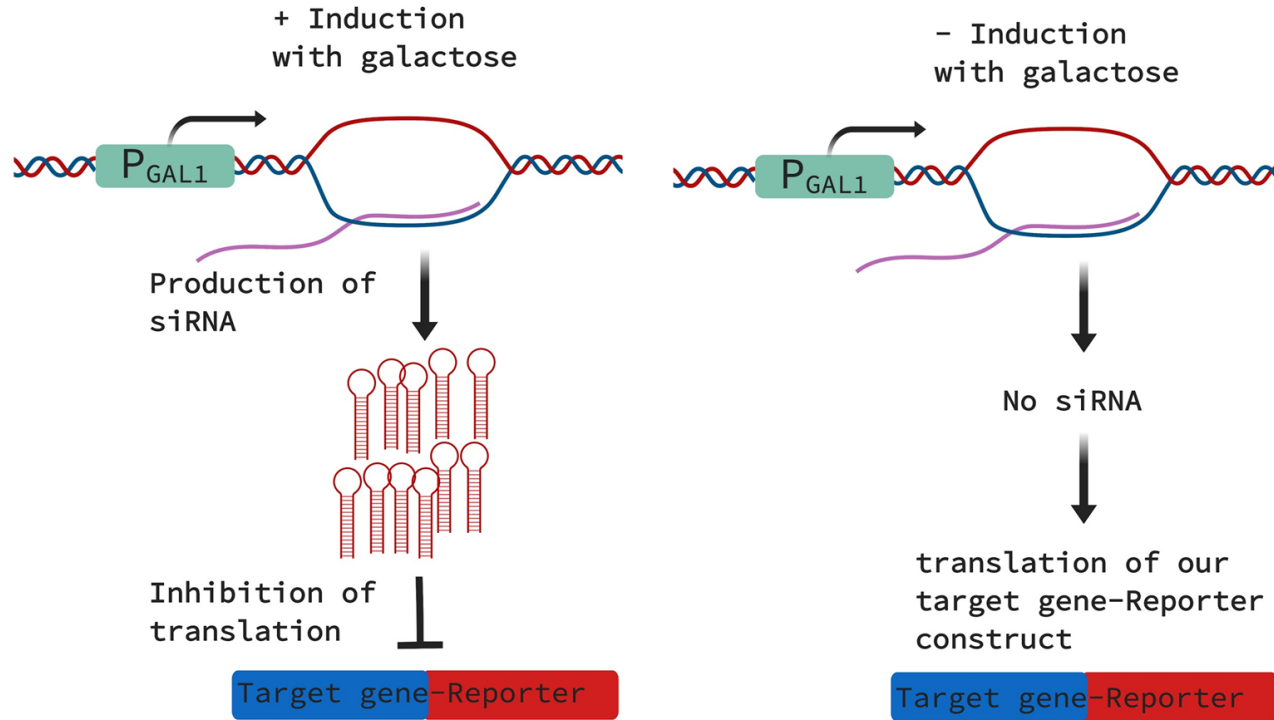
S. cerevisiae as a model organism

- Single celled eukaryotic organism and easy to manipulate yeast genetics.
- Obtained engineered yeast strains which contain the RISC complex.



- Towards this project we are creating new yeast strain with *P. brassicae* gene in its genome by using homologous recombination.
- For that we constructed the target gene with fluorescent reporter protein.
- Upon successful construction of yeast strain, we would like to use it for our future experiments.

Work plan



- Conditional expression of siRNA in yeast
- Screen for potential siRNA molecule against our target gene

Future Directions

- Insertion of target genes in yeast genome.
- Screening of potential siRNAs agonistic in *P. brassicae*.
- Upon successful knockdown in yeast, expand the study to *P. brassicae*



<https://www.canolacouncil.org/canola-encyclopedia/diseases/clubroot/about-clubroot/>



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References

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