

Light on the horizon: A molecular discovery opens the door for developing cabbage varieties resistant to black rot

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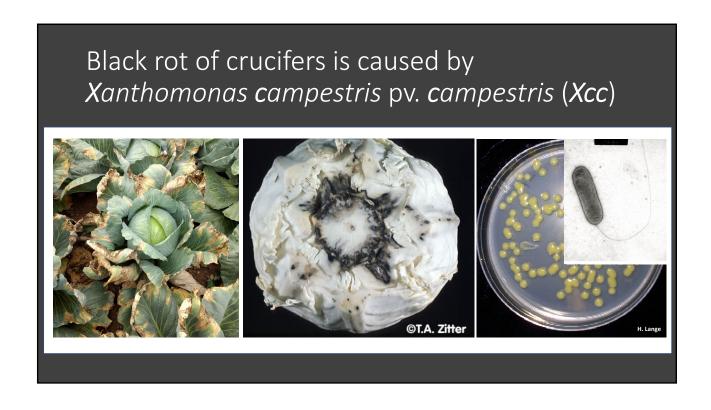
Outline

An overview of Black Rot in New York State

Cabbage resistance and susceptibility to the Black Rot pathogen

Using the Black Rot pathogen to find cabbage weak spots

# Black rot of crucifers is caused by Xanthomonas campestris pv. campestris (Xcc) The second of crucifers is caused by Xanthomonas campestris pv. campestris (Xcc)



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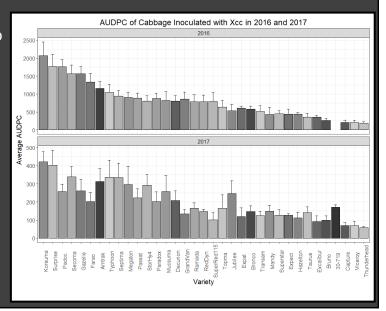
#### Black Rot in New York State

- Smart Lab collected isolates of Xcc for 15 years
  - Genetically different isolates each year
  - Not overwintering in fields
- Xcc primarily spread via seed
- Hot water treatments can limit disease, but...
- 1 in 10,000 infected seed can cause an epidemic



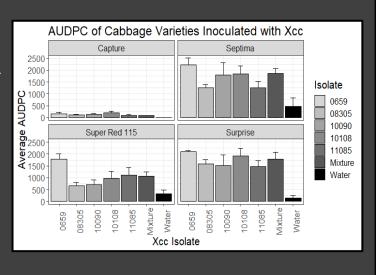
#### Black Rot in New York State

- Susceptibility of 35 varieties to a single NY Xcc isolate
- Tolerant varieties available, but no true resistance!



#### Black Rot in New York State

- Susceptibility of 4 varieties to diverse NY Xcc isolates
- Conclusion: Broad tolerance or susceptibility to the range of Xcc isolates in New York



## Resistance and Susceptibility of Cabbage Varieties to Black Rot

- Tolerant varieties available, but what makes them tolerant is unknown
- Understanding which genes make cabbage tolerant or susceptible to Black Rot -> commercial variety improvement
- Question: What makes cabbage tolerant or susceptible to Black Rot?

## What makes cabbage tolerant or susceptible to Black Rot?

- No major resistance (R) genes for black rot
- Bogdanove lab at Cornell studies Susceptibility (S) genes in rice
- Breeding plants that lack susceptibility genes leads to less disease







• Closely related pathogens to Xcc cause disease in 300+ plant species.

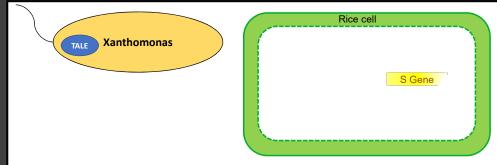


## What do we know about susceptibility genes?

- Closely related pathogens to Xcc cause disease in 300+ plant species.
- Some Xanthomonas pathogens use tools called TAL Effectors to survive in their plant hosts

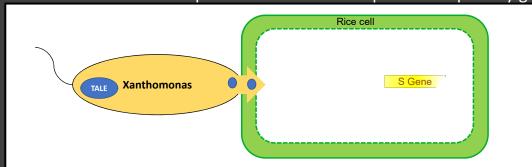


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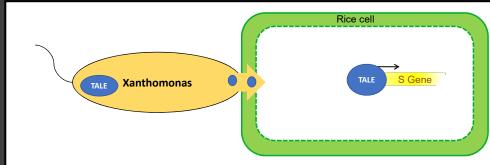


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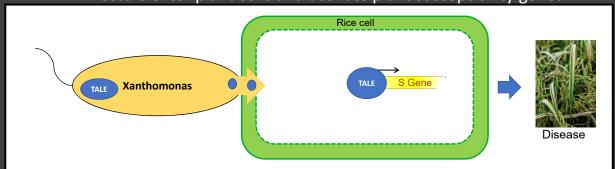


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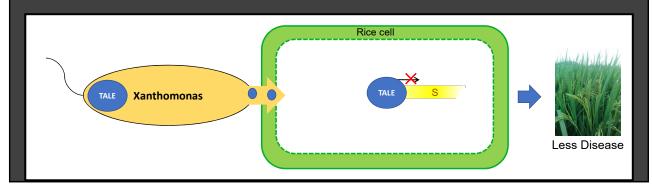


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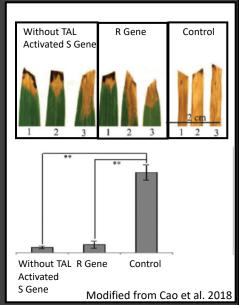


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# Loss of TAL Effector activated susceptibility genes results in resistance in rice

- TAL effector activated Susceptibility genes have been bred out of rice to control Bacterial Blight
  - Susceptibility genes are often Sugar Transporters
- Successfully used in the field
- Can we do this in cruciferous crops?

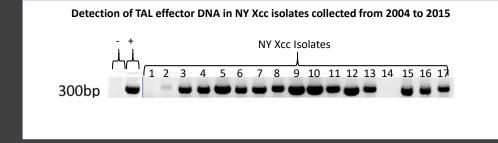


- 1. Does the pathogen have TAL effectors?
- 2. Are TAL effectors using genes important to susceptibility?
- 3. What genes are the TAL effectors using?
- 4. Modify or remove susceptibility genes and determine if plants are less susceptible to disease

## Will this strategy work to reduce susceptibility of cabbage to Black Rot?

- Does Xcc have TAL effectors?
  - Lab test to see if TAL effector DNA is present

• Does Xcc have TAL effectors?



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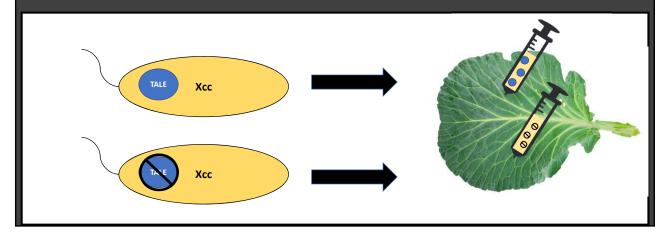
• Does Xcc have TAL effectors?

Detection of TAL effector DNA in NY Xcc isolates collected from 2004 to 2015

- Does Xcc have TAL effectors?
- Are Xcc TAL effectors using genes important to susceptibility?

# Are Xcc TAL effectors using genes important to susceptibility?

- Remove TALEs from Xcc
- Xcc without TAL effectors less virulent than Xcc with TAL effectors?



## Are TAL Effectors important in black rot? Yes!



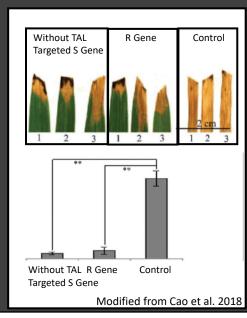
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  - Removal of several TAL effectors from NY Xcc strains, results in reduced disease
- What genes are the TAL effectors using?
  - Experiments in progress
  - Initial data: Sugar transporter genes (the same susceptibility genes from rice, cotton, and cassava)



#### Conclusions

- Finding susceptibility genes that can be bred out of cabbage = strategy to reduce disease
- TAL effectors can be used to find susceptibility genes in cabbage
- Successful in other crops like rice
- Still a lot to do!



### Acknowledgments

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  - Smart Lab-Cornell
    - Chris Smart
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    - Laurent Noel
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