

Update on the determination of rust (*Puccinia helianthi*) prevalence and virulence in the Northern Great Plains

Zachary Ittel, Brandt Berghuis, Jessica Scherer, Kaylie Schlecht, Robert Harveson, Febina Mathew, Brent Hulke, Bryan Hanson, and Samuel Markell

Take Home Message & Results

- Rust is important
- High Prevalence:
 - Prevalence = Does rust occur in a field? (Yes/Fields Surveyed)
 - 98% of fields over two years had rust

Take Home Message & Results

- Rust is important
- High Prevalence:
 - Prevalence = Does rust occur in a field? (Yes/Fields Surveyed)
 - 98% of fields over two years had rust
- Virulence Progress:
 - Virulence = Can the pathogen overcome a resistance gene?
 - Over 200 bulk pathogen isolates collected
 - Will be evaluating virulence on many new resistance genes

Objective 1

Determine the prevalence of *P. helianthi* in the Northern Great Plains.

Prevalence: Background

- 1.315 million acres of Sunflower were planted in the USA in 2023.
- Top 5 most destructive sunflower diseases.
- Prevalence (2002-2015)
 - Range approximately 20-80%
 - Mean of about 40-50%

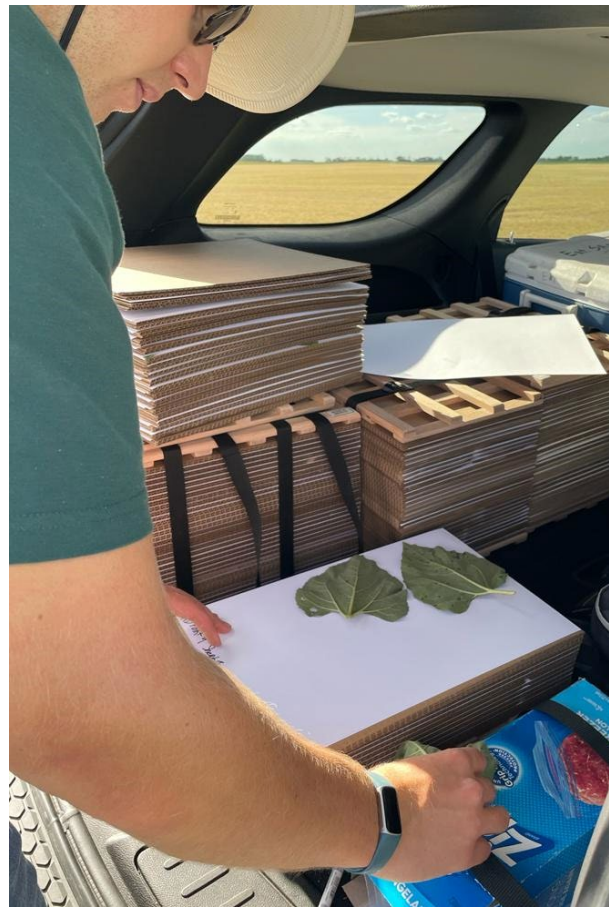


(Photo: Zachary Ittel)

Prevalence: Materials & Methods



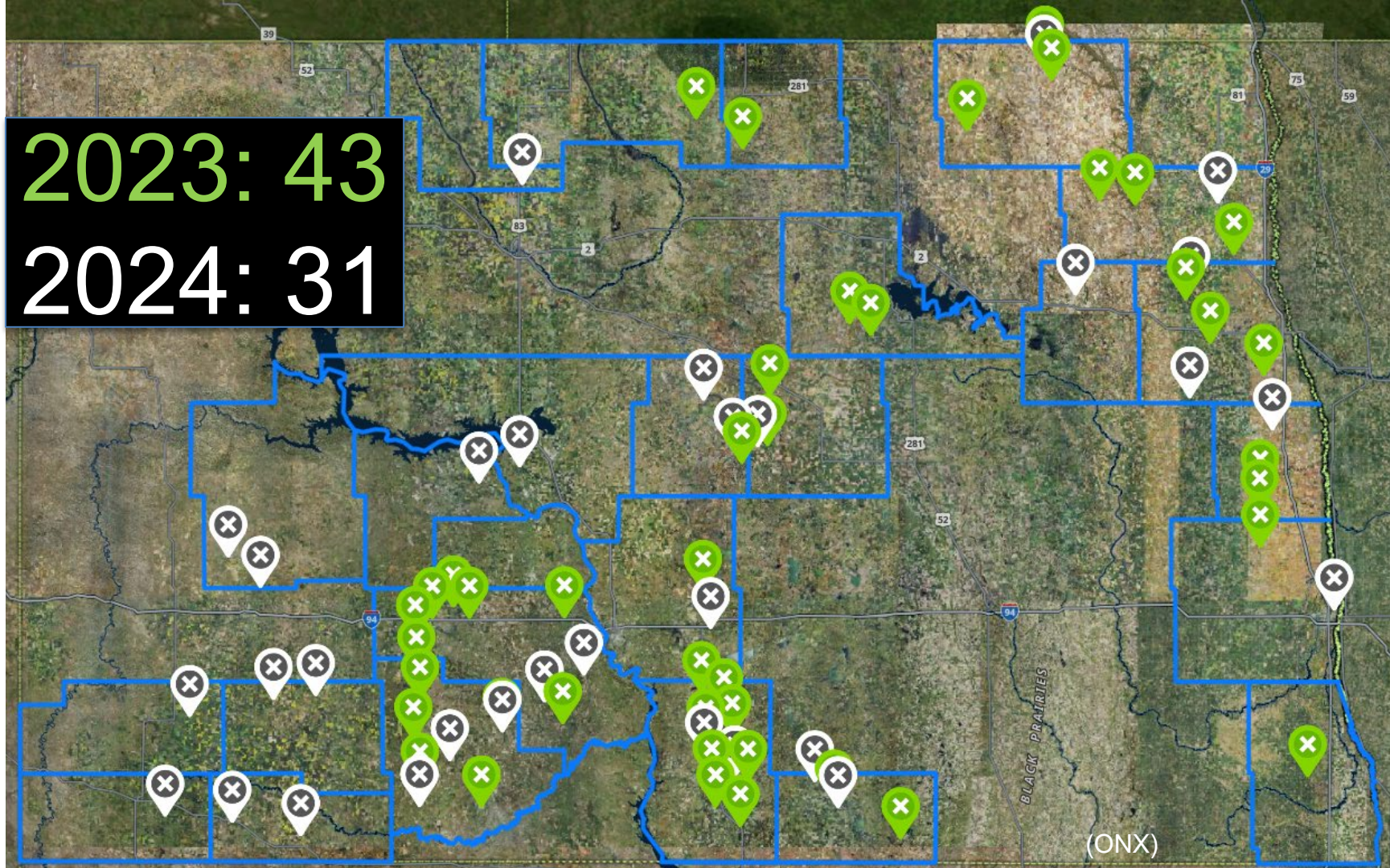
NDSU NORTH DAKOTA
STATE UNIVERSITY



ND Survey Map 2023-2024

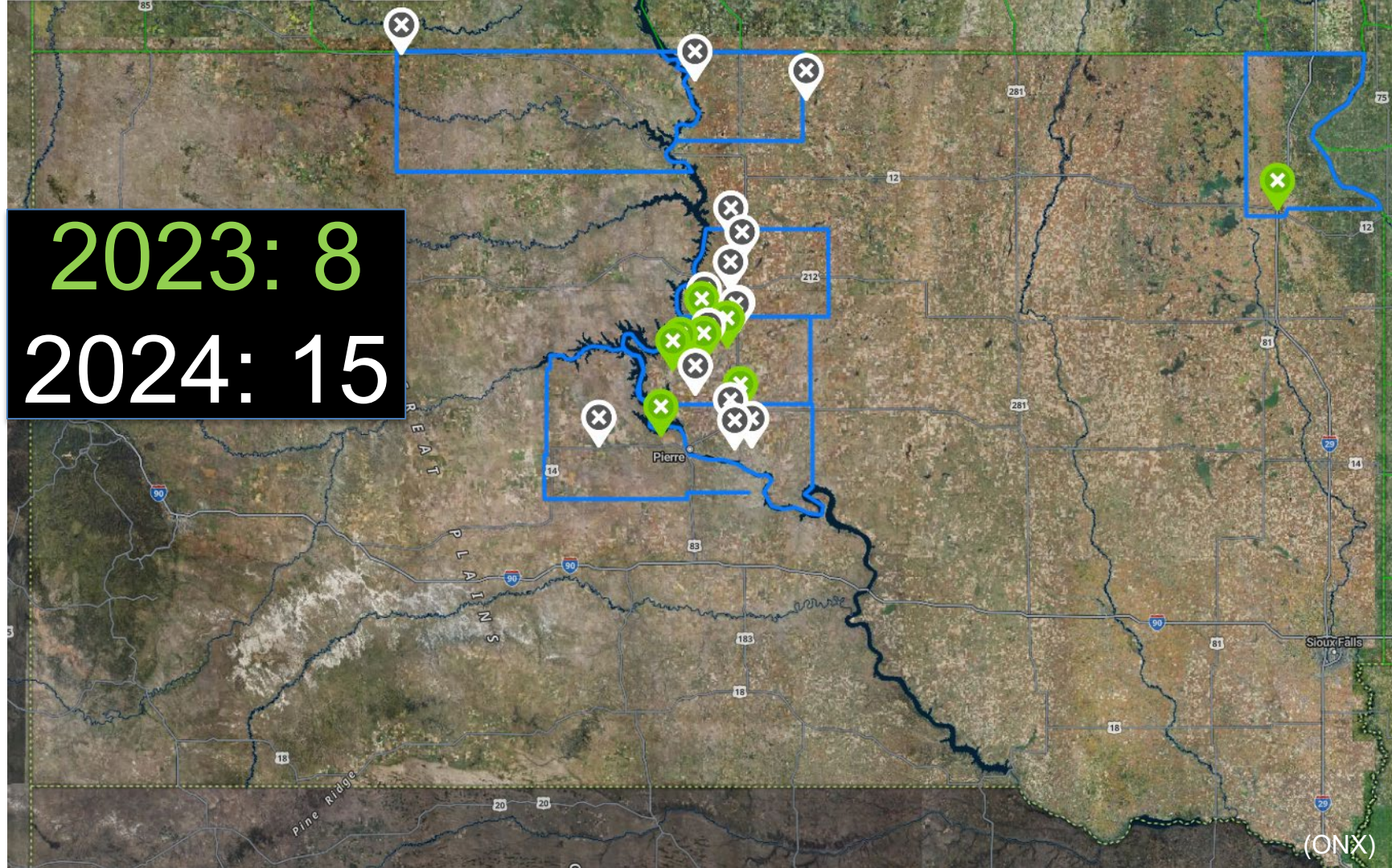
2023: 43

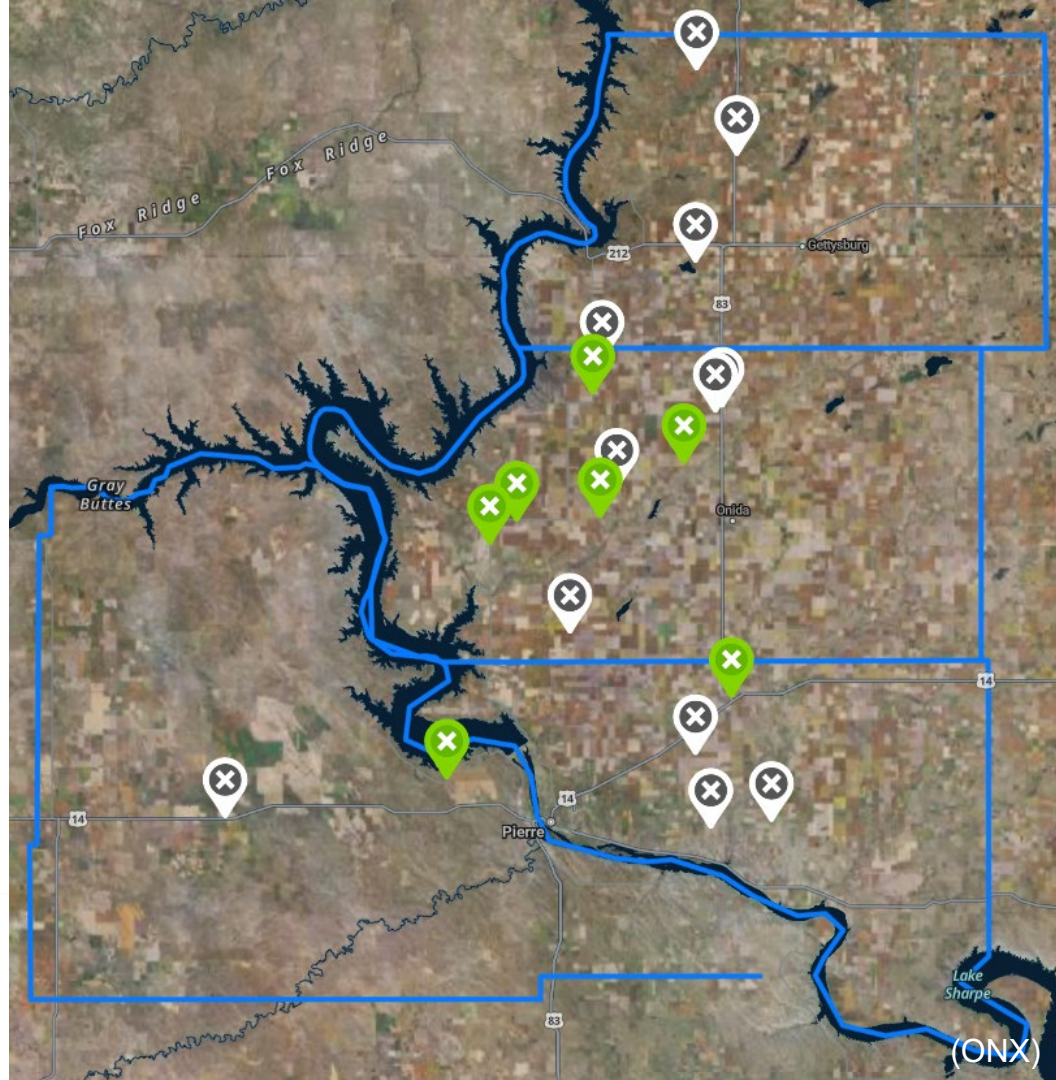
2024: 31



SD Survey Map 2023-2024

2023: 8
2024: 15

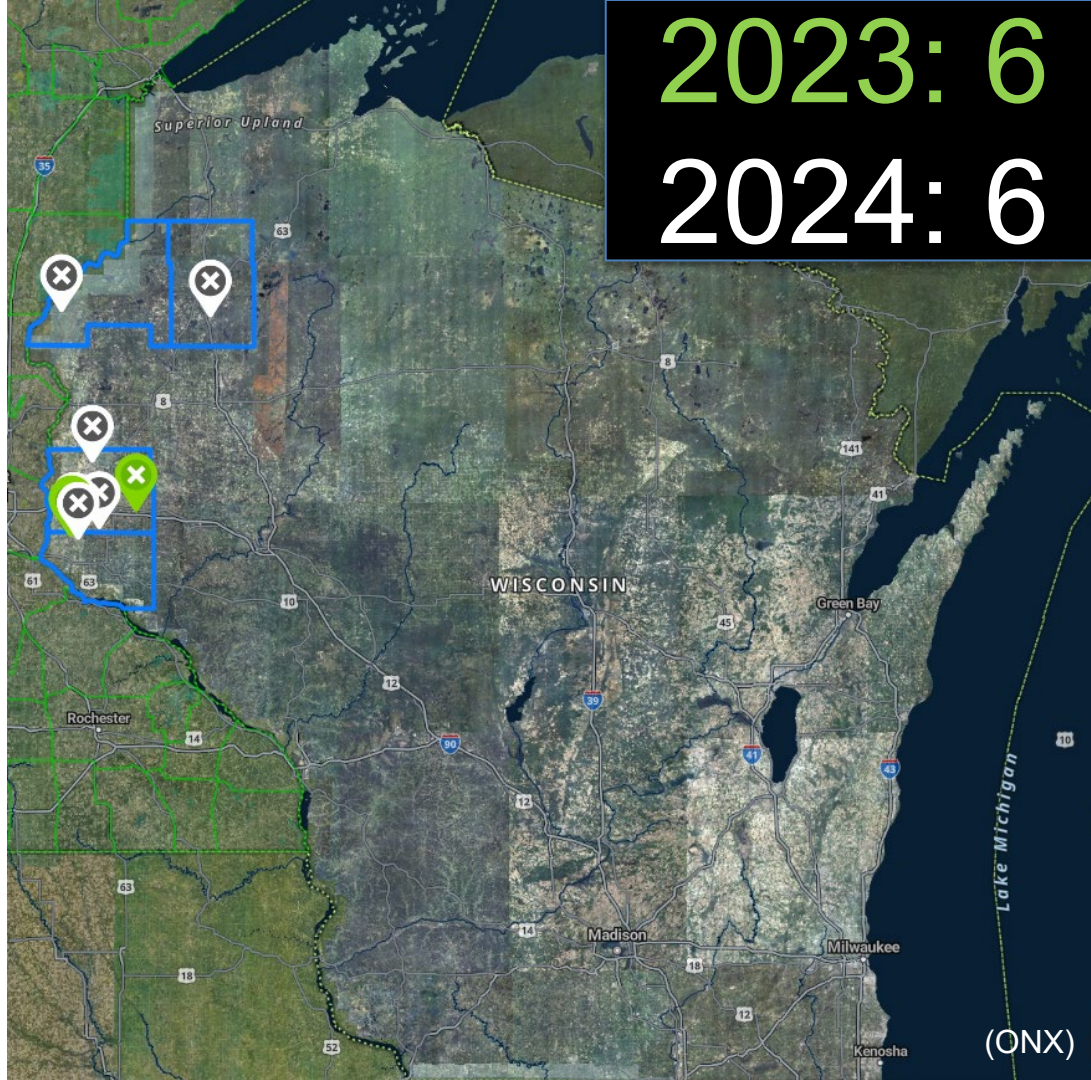


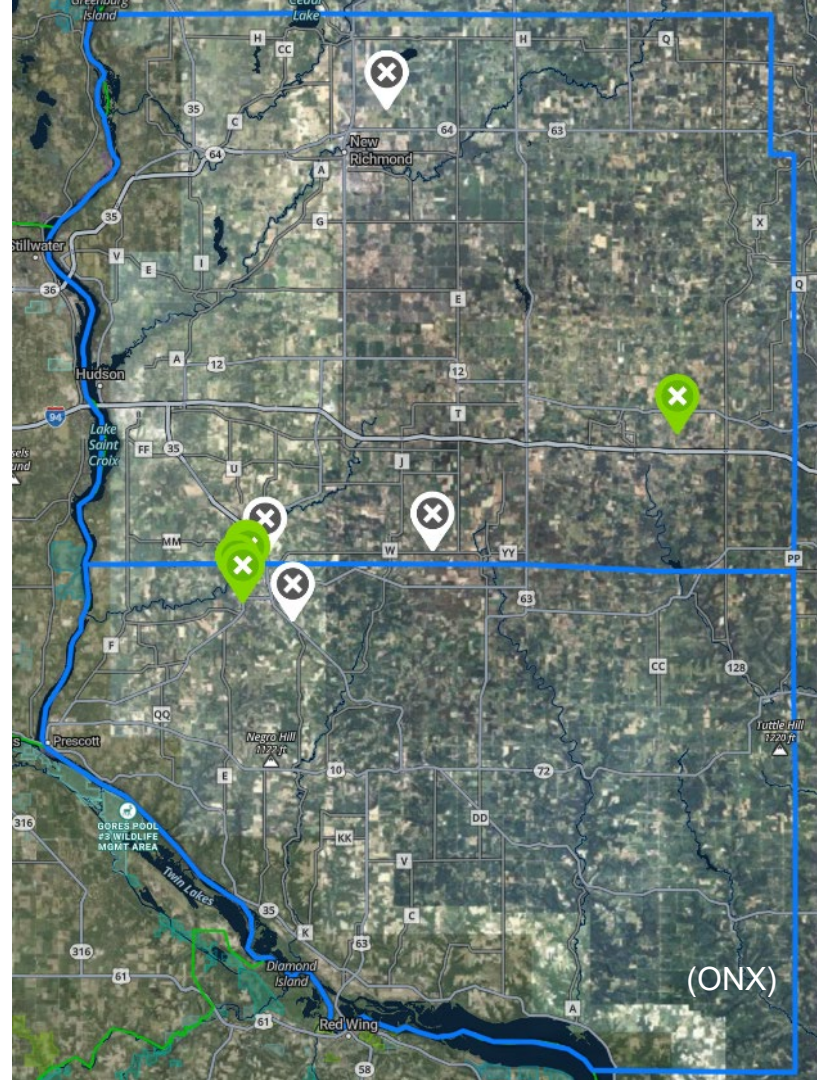


WI Survey Map 2023-2024

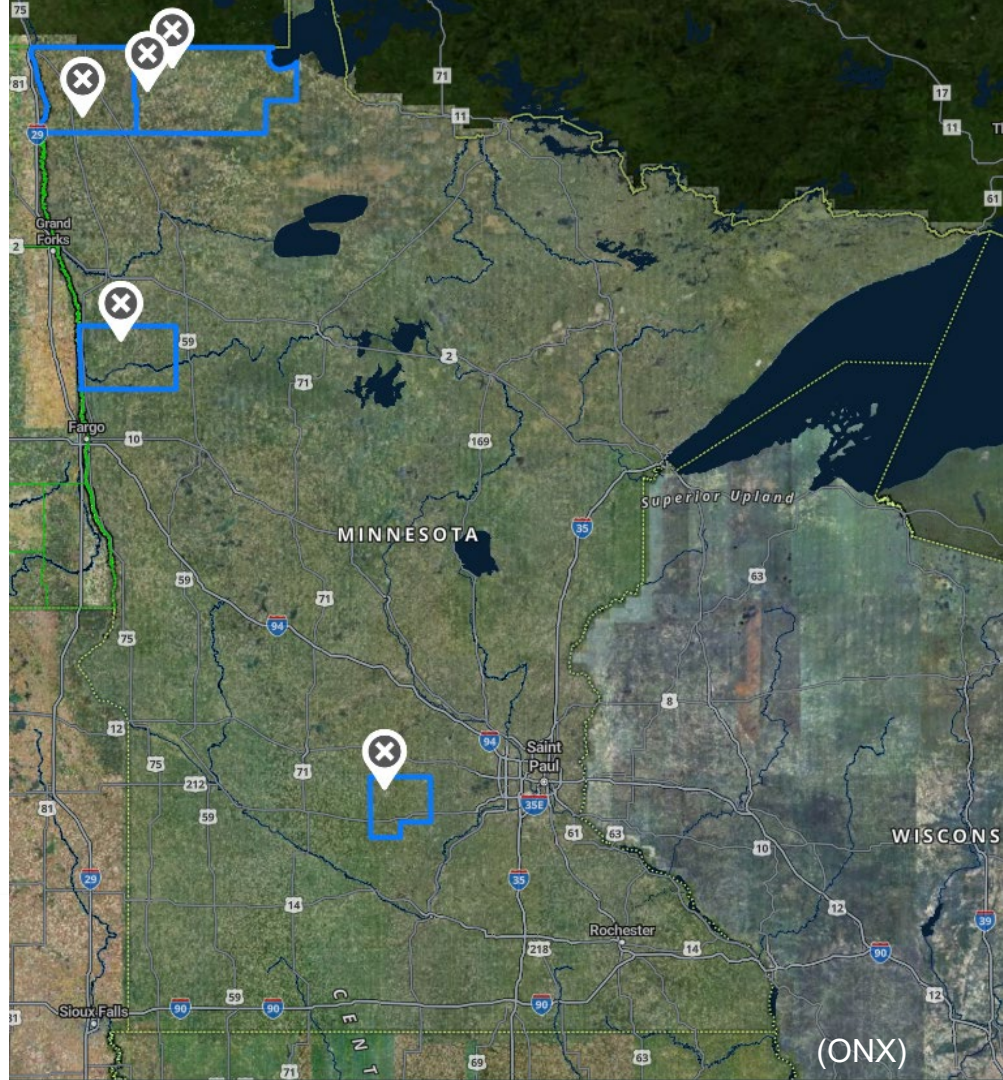
2023: 6

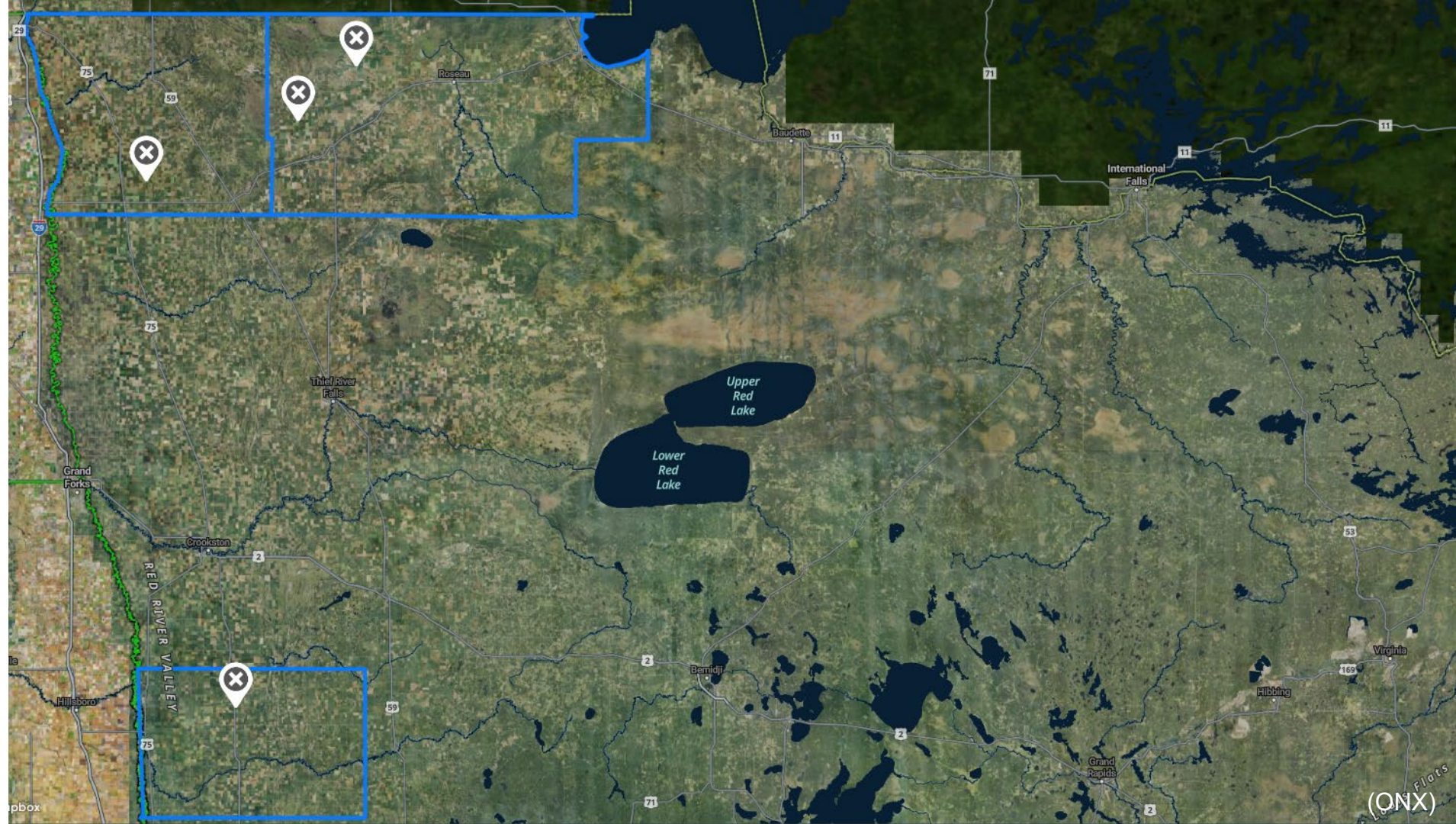
2024: 6





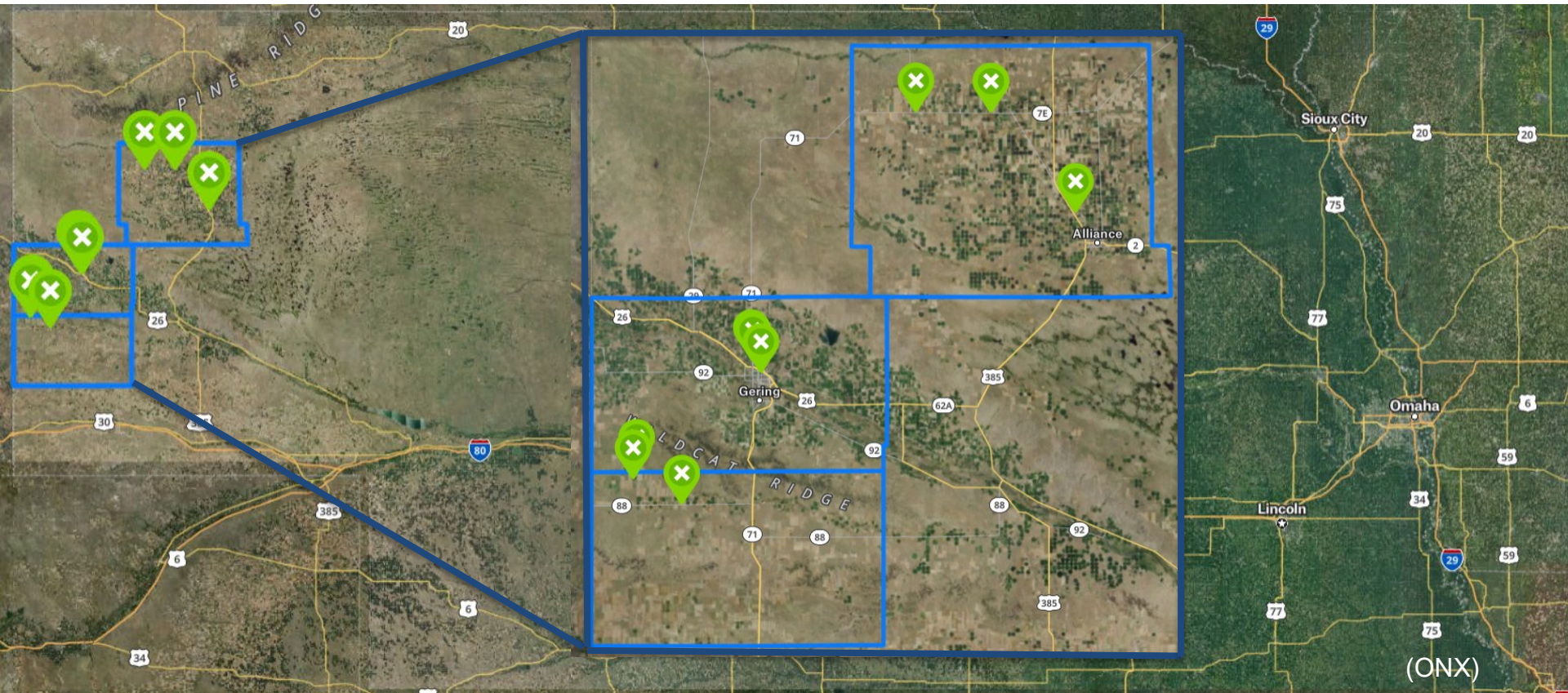
MN Survey Map 2024





(ONX)

NE Survey Map 2023



Prevalence: Results

	2023			
North Dakota	41/43			
South Dakota	8/8			
Wisconsin	6/6			
Nebraska	10/10			
Minnesota	...			
Totals	65/67			

Prevalence: Results

	2023	2024		
North Dakota	41/43	31/31		
South Dakota	8/8	15/15		
Wisconsin	6/6	6/6		
Nebraska	10/10	...		
Minnesota	...	5/5		
Totals	65/67	57/57		

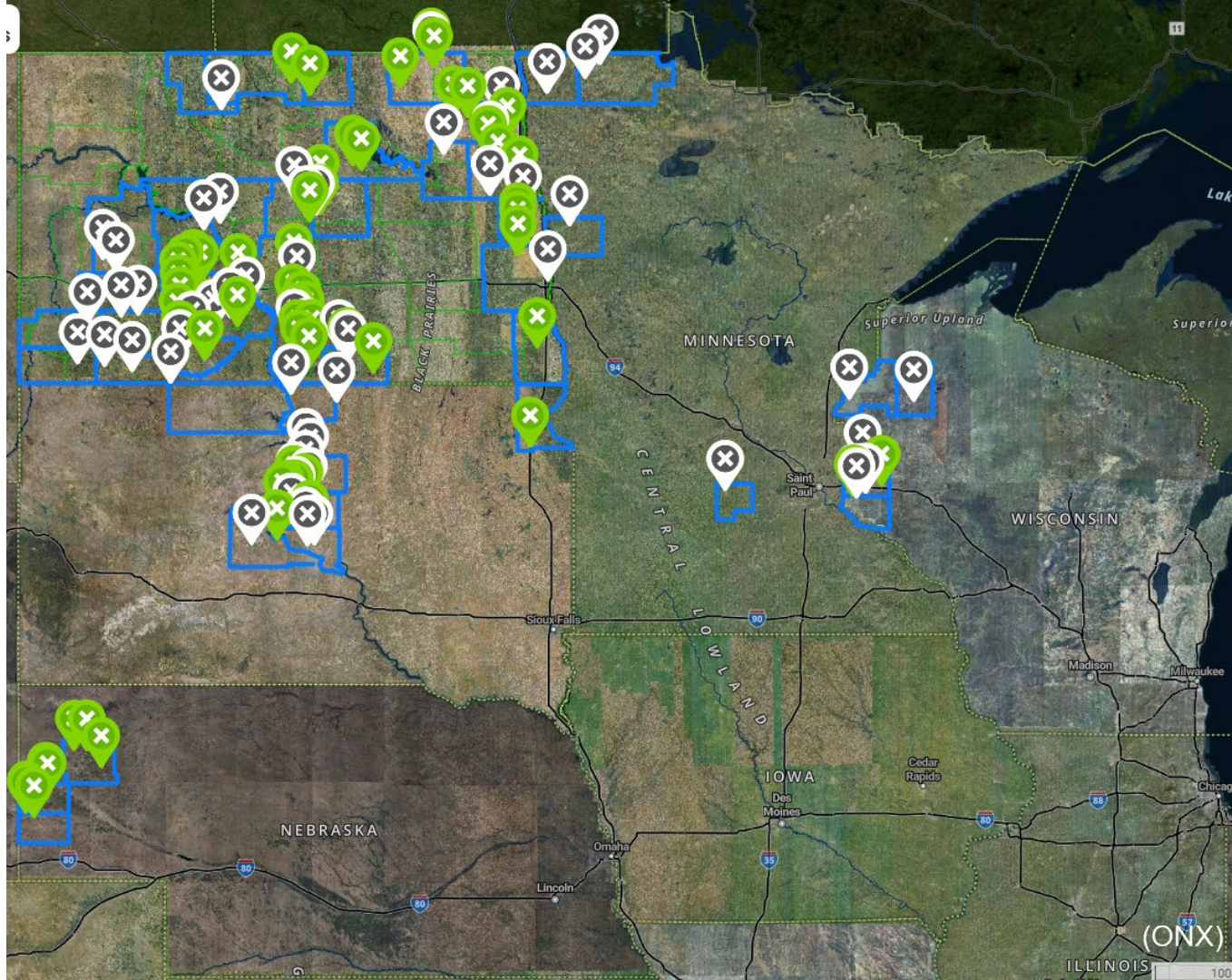
Rust Prevalence 2023-2024

122/124
Fields

>98%

Objective 2

Determine the virulence phenotype of *P. helianthi* in the Northern Great Plains.



Prevalence: Results

	2023	2024	Bulk Isolates	
North Dakota	41/43	31/31	123	
South Dakota	8/8	15/15	45	
Wisconsin	6/6	6/6	33	
Nebraska	10/10	...	30	
Minnesota	...	5/5	3	
Totals	65/67	57/57	234	

Prevalence: Results

	2023	2024	Bulk Isolates	SPI's
North Dakota	41/43	31/31	123	3
South Dakota	8/8	15/15	45	22
Wisconsin	6/6	6/6	33	6
Nebraska	10/10	...	30	0
Minnesota	...	5/5	3	0
Totals	65/67	57/57	234	31

Virulence: Background

- New races can overcome genetic resistance
- Management:
 - Genetic Resistance
 - Fungicides
 - Removal of volunteer plants



(Photo: Zachary Ittel)

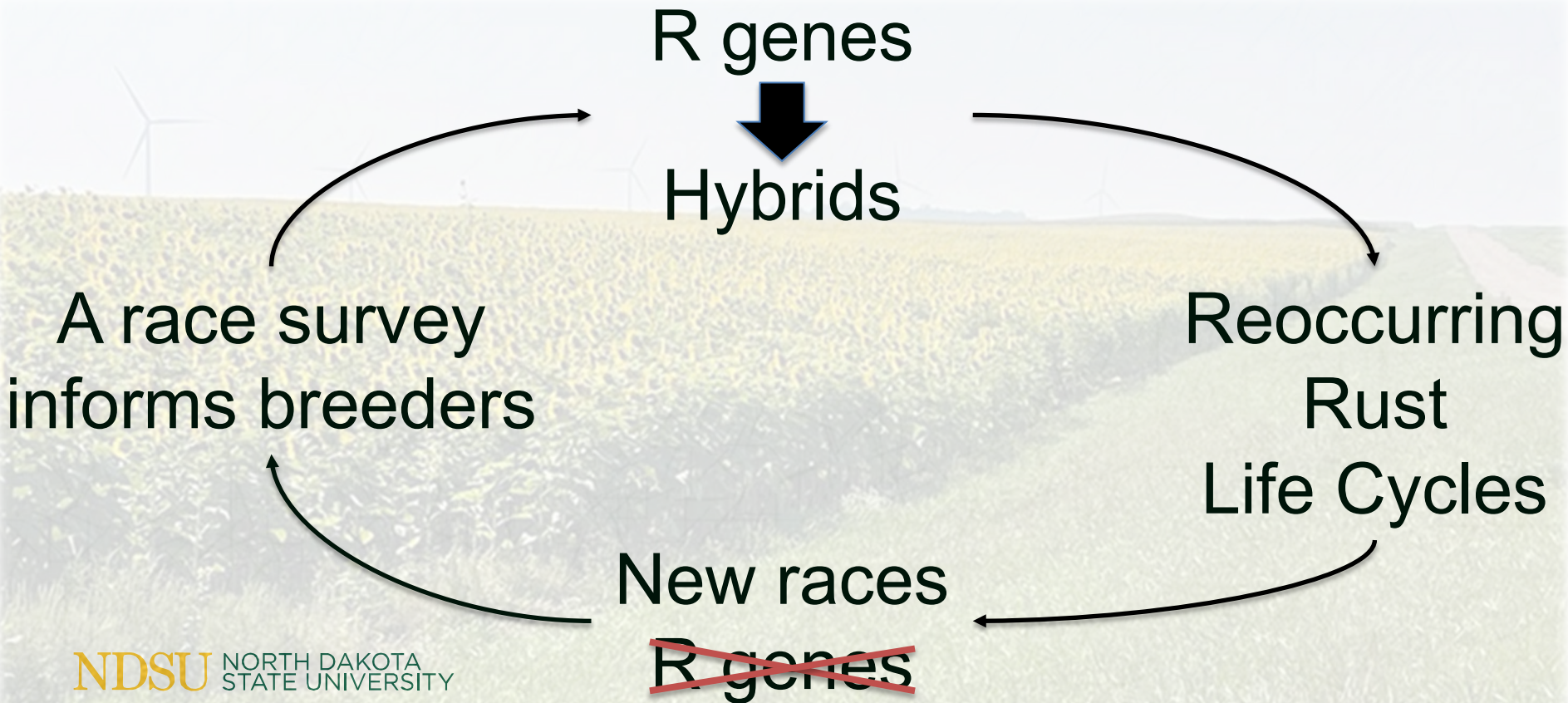
Virulence: Background

- New races can overcome genetic resistance
- Management:
 - Genetic Resistance
 - Fungicides
 - Removal of volunteer plants



(Photo: Zachary Ittel)

Virulence: Background



Virulence: Background



P. helianthi Virulence in 2011-2012

- 238 single pustule isolates
 - 29 races
- States: NE, ND, SD, CA, IA, MN, TX, and MB.
- NE, ND, and MB each had rust that could overcome all known R genes



Dr. Andrew Friskop

P. helianthi Virulence in CA in 2017-2018

- Production fields:
 - 2017 – 6 races (11 SPI)
 - 2018 – 8 races (13 SPI)
- Wild sunflower:
 - 2017 – 6 races (11 SPI)
 - 2018 – 16 races (20 SPI)
- CA had rust that could overcome all known R genes



Virulence: Methods (Pathogen Recovery)



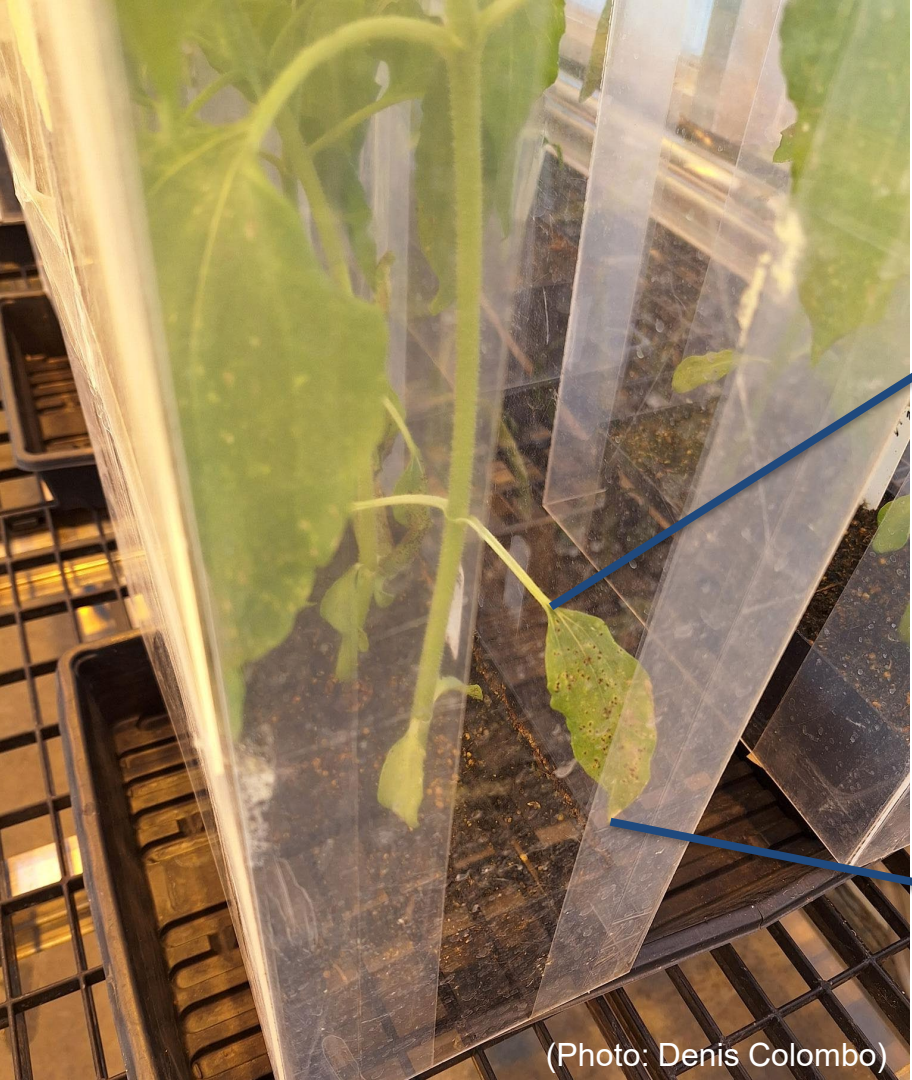
(Photo: Denis Colombo)



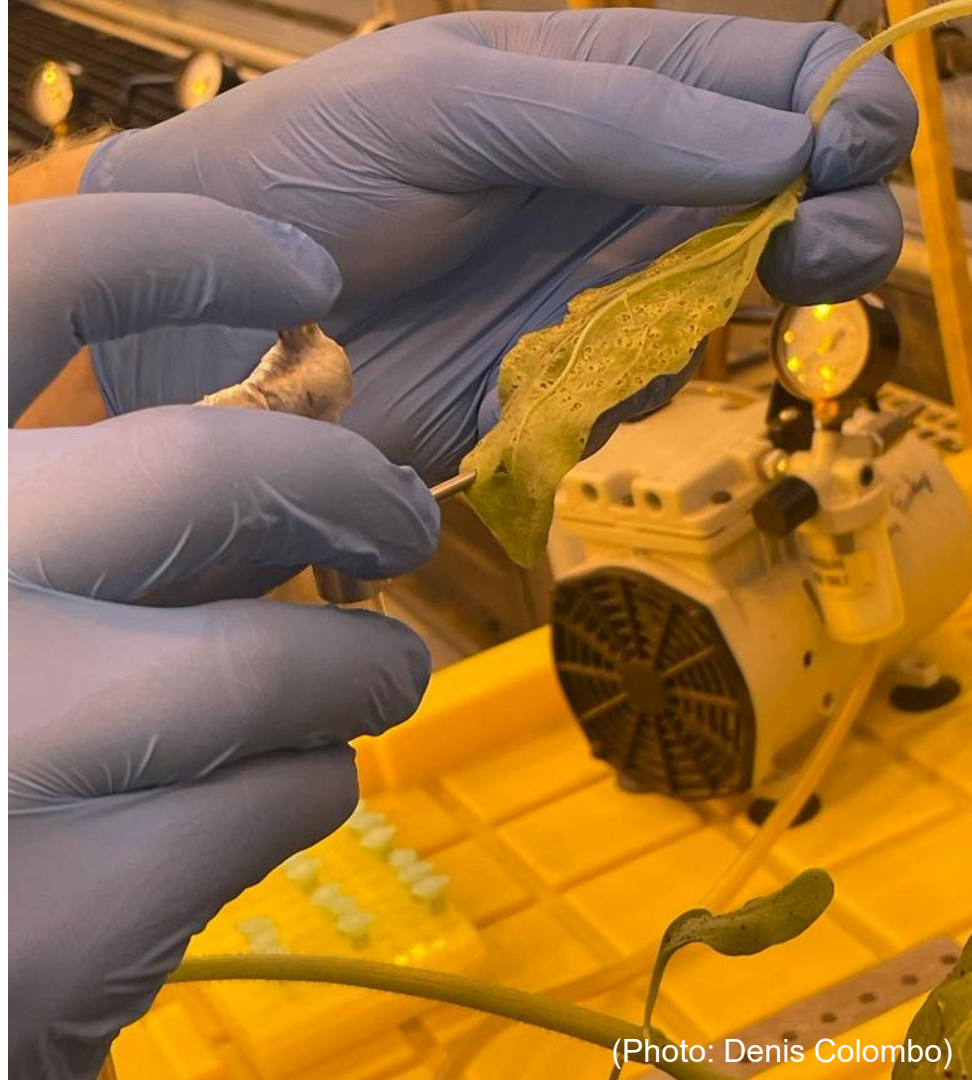
(Photo: Zachary Ittel)



(Photo: Zachary Ittel)



(Photo: Denis Colombo)



(Photo: Denis Colombo)



(Photo: Denis Colombo)



(Photo: Zachary Ittel)

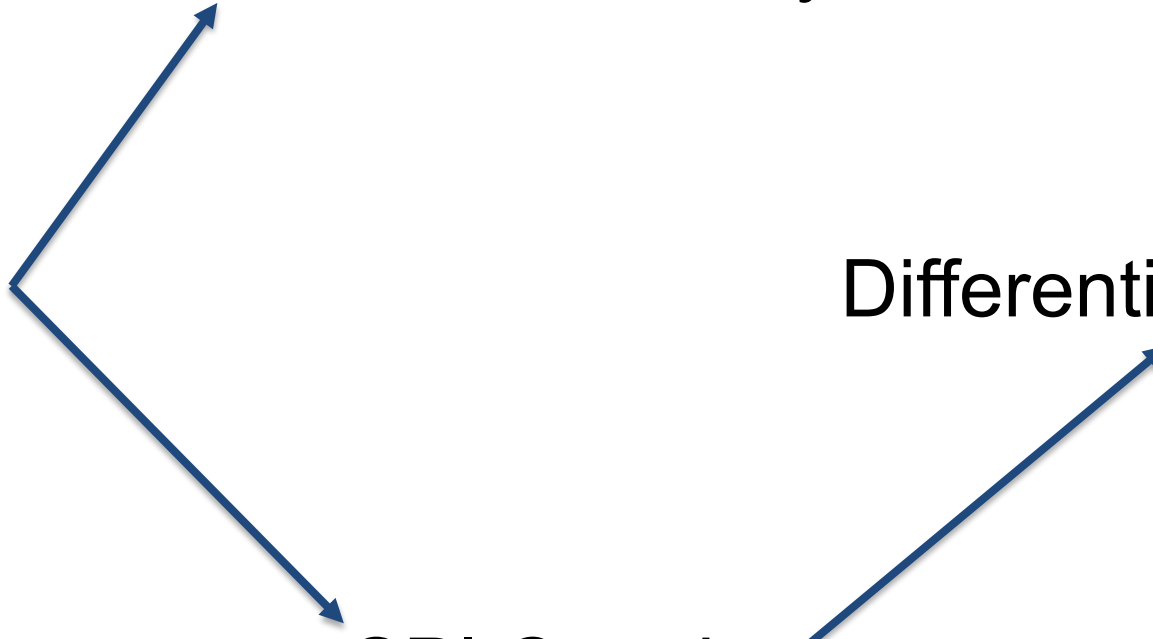
Virulence: Next Steps

Bulk Recovery

Recovery

Differential Testing

SPI Creation



Determining Virulence Phenotype (Race)

Universal Differential Set

Triplet	Differential	Resistance Gene	com g value
One	7350	...	...
	MC90	R_{10}	2
	MC29	$R_2 + R_{10}$	4
Two	P386	R_{4e}	1
	HA-R1	R_{4a}	2
	HA-R2	R_5	4
Three	HA-R3	R_{4b}	1
	HA-R4	R_{4c}	2
	HA-R5	R_{4d}	4

Triplet	Differential	Resistance Gene	Scoring Value	Code
One	7350	...	1	
	MC90	R_1	2	
	MC29	R_2+R_{10}	4	

Triplet	Differential	Resistance Gene	Scoring Value	Code
One	7350	...	1	3
	MC90	R_1	2	
	MC29	R_2+R_{10}	4	

Triplet	Differential	Resistance Gene	Scoring Value	Code
One	7350	...	1	3
	MC90	R_1	2	
	MC29	R_2+R_{10}	4	
Two	P386	R_{4e}	1	
	HA-R1	R_{4a}	2	
	HA-R2	R_5	4	

Triplet	Differential	Resistance Gene	Scoring Value	Code
One	7350	...	1	3
	MC90	R_1	2	
	MC29	R_2+R_{10}	4	
Two	P386	R_{4e}	1	4
	HA-R1	R_{4a}	2	
	HA-R2	R_5	4	

Triplet	Differential	Resistance Gene	Scoring Value	Code
One	7350	...	1	3
	MC90	R_1	2	
	MC29	R_2+R_{10}	4	
Two	P386	R_{4e}	1	4
	HA-R1	R_{4a}	2	
	HA-R2	R_5	4	
Three	HA-R3	R_{4b}	1	
	HA-R4	R_{4c}	2	
	HA-R5	R_{4d}	4	

Triplet	Differential	Resistance Gene	Scoring Value	Code
One	7350	...	1	3
	MC90	R_1	2	
	MC29	R_2+R_{10}	4	
Two	P386	R_{4e}	1	4
	HA-R1	R_{4a}	2	
	HA-R2	R_5	4	
Three	HA-R3	R_{4b}	1	6
	HA-R4	R_{4c}	2	
	HA-R5	R_{4d}	4	

Race Name =
346

Triplet	Differential	Resistance Gene	Scoring Value	Code
One	7350	...	1	
	MC90	R_1	2	
	MC29	R_2+R_{10}	4	
Two	P386	R_{4e}	1	
	HA-R1	R_{4a}	2	
	HA-R2	R_5	4	
Three	HA-R3	R_{4b}	1	
	HA-R4	R_{4c}	2	
	HA-R5	R_{4d}	4	
Four	?	?	?	?
	?	?	?	
	?	?	?	



Differential Additions

Differential Name	Pedigree	Gene	Year Released
HAR-9	Selection of Rf ANN 1742	R_{11}	2013
RHA 464	Selection of PI 413047	R_{12}	2013
HAR-6	(R702CLPlus/HAR6-1)F2	R_{13a}	2013
HAR-8	Selection of PI 432512	R_{15}	2018
HAR-18	Selection of South Africa line KP193	R_{17}	2020
HAR-19	Selection of South Africa line KP193	R_{18}	2020

Triplet	Differential	Resistance Gene	Scoring Value
One	7350	...	1
	MC90	R_1	2
	MC29	R_2+R_{10}	4
Two	P386	R_{4e}	1
	HA-R1	R_{4a}	2
	HA-R2	R_5	4
Three	HA-R3	R_{4b}	1
	HA-R4	R_{4c}	2
	HA-R5	R_{4d}	4
Four	HAR-9	R_{11}	1
	RHA 464	R_{12}	2
	HAR-6	R_{13a}	4
Five	HAR-8	R_{15}	1
	HAR-18	R_{17}	2
	HAR-19	R_{18}	4

Take Home Message & Results

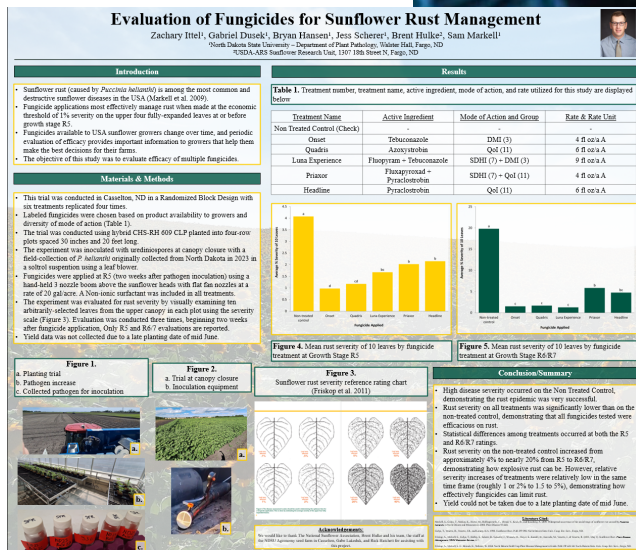
- Rust is important
- High Prevalence:
 - Prevalence = Does rust occur in a field? (Yes/Fields Surveyed)
 - 98% of fields over two years had rust

Take Home Message & Results

- Rust is important
- High Prevalence:
 - Prevalence = Does rust occur in a field? (Yes/Fields Surveyed)
 - 98% of fields over two years had rust
- Virulence Progress:
 - Virulence = Can the pathogen overcome a resistance gene?
 - Over 200 bulk pathogen isolates collected
 - Will be evaluating virulence on many new resistance genes

Future Objectives

- Whole Genome Pathogen Sequencing
- *Puccinia helianthi* population genetic analysis
- SAM pop experiment
- Fungicide evaluation



Acknowledgements

Survey Help:

- Advisor: Sam Markell
 - His team
- Co-advisor: Upinder Gill
- Brent Hulke
- ND (County Agents, NSA Board members, and Farmers)
- SD (Mr. Weber, NSA Board members, Farmers, and Febina Mathew's Team)
- NE (Bob Harveson and his team)
- WI (Brandt Burghuis and his team)
- MN (Farmers)
- Scott Radi
- Bill Underwood
- Kaylie Schlecht
- John Sandbakken



References

- Berghuis, B. (2021). Phenotypic diversity of *Puccinia helianthi* on sunflower in California [Doctoral dissertation, North Dakota State University].
- Bulos, M., Ramos, M. L., Altieri, E., & Sala, C. A. (2013). Molecular mapping of a sunflower rust resistance gene from HAR6. *Breeding science*, 63(1), 141–146. <https://doi.org/10.1270/jsbbs.63.141>
- Friskop, A. J., Gulya, T. J., Harveson, R. M., Humann, R. M., Acevedo, M., and Markell, S. G. (2015). Phenotypic diversity of *Puccinia helianthi* (sunflower rust) in the United States from 2011 and 2012. *Plant Dis.* 99:1604-1609.
- Gong, L., Hulke, B. S., Gulya, T. J., Markell, S. G., & Qi, L. L. (2013). Molecular tagging of a novel rust resistance gene R(12) in sunflower (*Helianthus annuus* L.). *TAG. Theoretical and applied genetics. Theoretische und angewandte Genetik*, 126(1), 93–99. <https://doi.org/10.1007/s00122-012-1962-z>
- Gulya, T., Harveson, R., Mathew, F., Block, C., Thompson, S., Kandel, H., Berglund, D., Sandbakken, J., Kleingartner, L., and Markell, S. (2019). Comprehensive disease survey of U.S. sunflower: Disease trends, research priorities and unanticipated impacts. *Plant Disease* 103:601-618.
- Gulya, T., and Masirevic, S. (1996). Inoculation and evaluation methods for sunflower rust. In: Proc. 18th Sunflower Research Workshop. National Sunflower Association, Bismarck, ND.
- Harveson, R., Markell, S., Block, C., and Gulya, T. (2016). Compendium of Sunflower Diseases and Pests. American Phytopathology Press, St. Paul, MN. 140 Pp.

References

- Meyer, W. B., Boshoff, W. H. P., Minnaar-Ontong, A., Young, A. J., Kong, G., Thompson, S., Pretorius, Z. A., Visser B. Phenotypic and Genotypic Variation of *Puccinia helianthi* in South Africa. Plant Disease 2021 May;105(5):1482-1489.
- Ma, G. J., Song, Q. J., Markell, S. G., & Qi, L. L. (2018). High-throughput genotyping-by-sequencing facilitates molecular tagging of a novel rust resistance gene, R 15 , in sunflower (*Helianthus annuus* L.). *TAG. Theoretical and applied genetics. Theoretische und angewandte Genetik*, 131(7), 1423–1432. <https://doi.org/10.1007/s00122-018-3087-5>
- ONX. Retrieved January, 2025, from <https://www.onxmaps.com/>
- Qi, L., Ma, G., Seiler, G.J. (2021). Registration of HA-R14, HA-R15, HA-R16, and HA-R17 oilseed sunflower germplasms with broad resistance to rust and downy mildew. *Journal of Plant Registrations*. 16:137–146. <https://doi.org/10.1002/plr2.20187>.
- Qi, L., Gong, L., Markell, S.G., Seiler, G.J., Gulya Jr, T.J., Hulke, B.S. (2014). Registration of two confection sunflower germplasm Lines, HA-R10 and HA-R11, Resistant to sunflower rust. *Journal of Plant Registrations*. 8:329-333. DOI: 10.3198/jpr2014.02.0010crg.
- Thorson, T. (2024). Crop Production 2023 Summary. United States Department of Agriculture National Agricultural Statistics Service, 48. <https://downloads.usda.library.cornell.edu/usdaesmis/files/k3569432s/ns065v292/8910md644/cropan24.pdf>

A man wearing a tan bucket hat with a logo, sunglasses, and a dark green shirt is smiling and holding a large sunflower. He is standing in a field of sunflowers under a cloudy sky. The text "Thank you" is overlaid in yellow and "Questions?" is overlaid in dark green.

Thank you
Questions?