



Update On The Determination Of Rust (*Puccinia helianthi*) Virulence and Associated Rust Projects

Zachary Ittel¹, Brandt Berghuis⁴, Jessica Scherer¹, Kaylie Schlecht¹, Robert Harveson³, Febina Mathew¹, Brent Hulke², Bryan Hansen¹, Samuel Markell¹

¹North Dakota State University – Department of Plant Pathology, Walster Hall, Fargo, ND

²USDA-ARS Sunflower Research Unit, 1307 18th Street N, Fargo, ND

³University of Nebraska – Panhandle Research and Extension Center, 4502 Avenue I, Scottsbluff, NE

⁴University of Wisconsin River Falls – Ag Science Building, 699, 611 S 3rd St, River Falls, WI



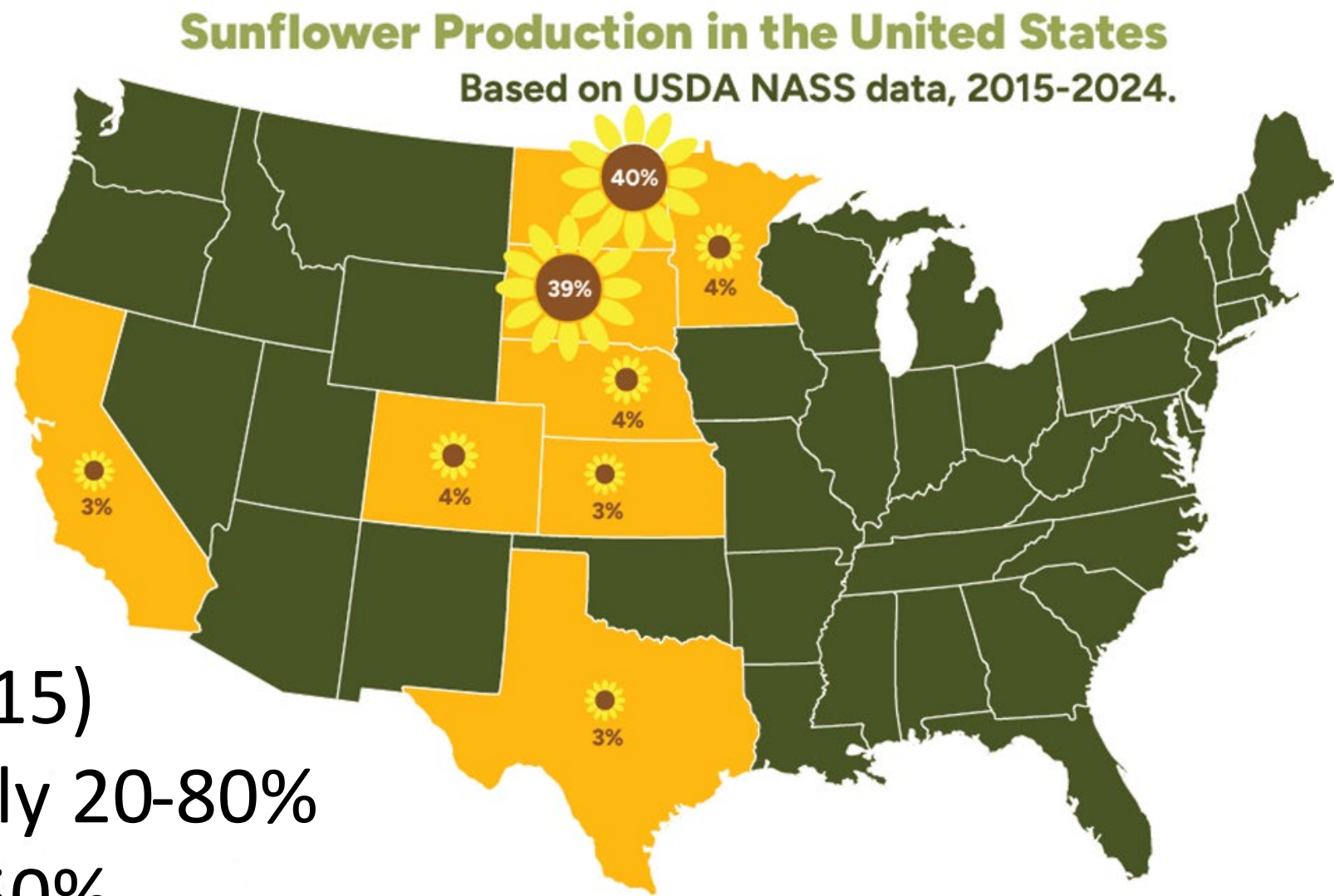
Project Update Outline

- ☀ Prevalence = Does rust occur in a field? (Yes/Fields Surveyed)
- ☀ Virulence = Can the pathogen overcome a resistance gene?
- ☀ Additional/Future objectives



Introduction: Sunflower

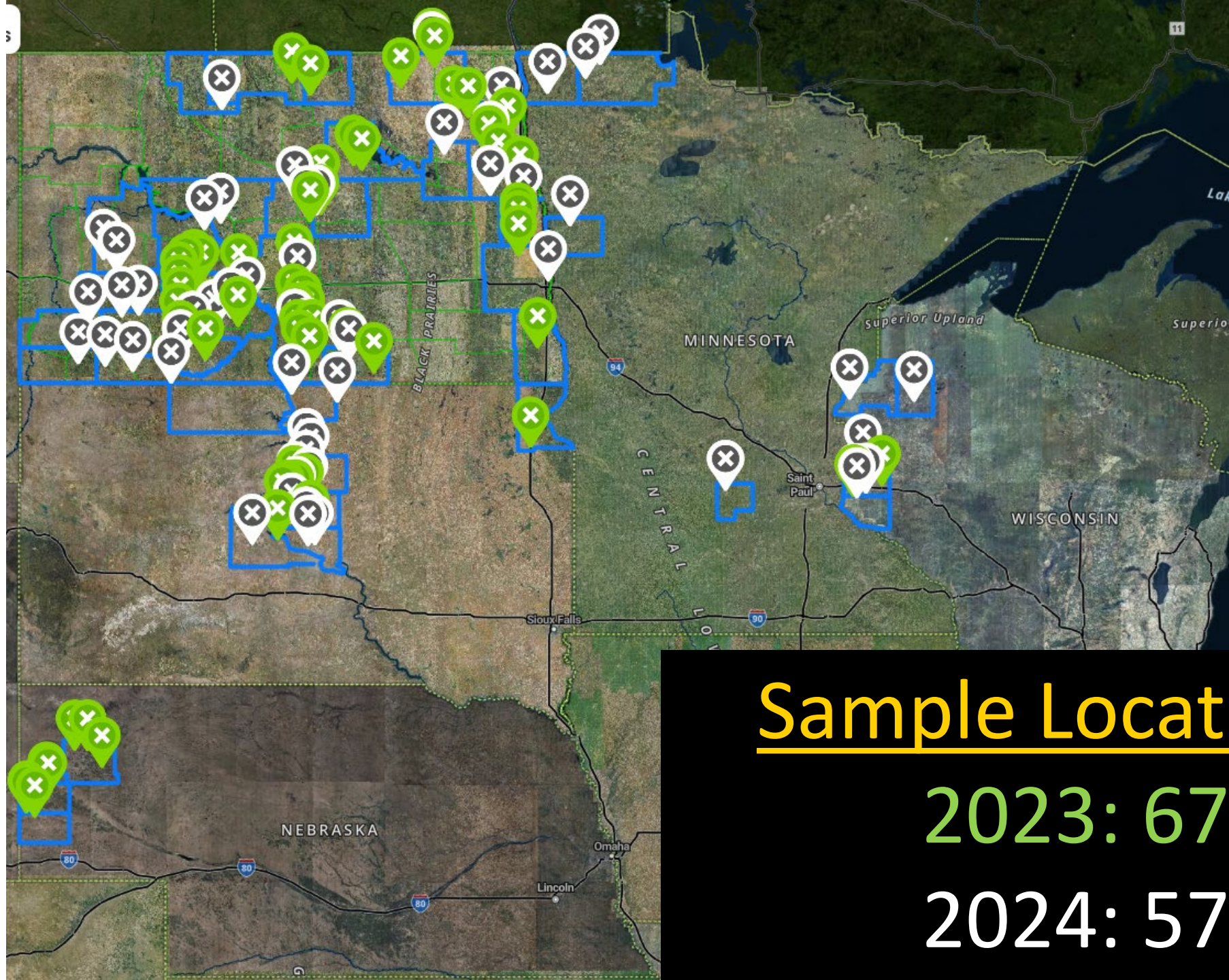
- ☀️ 1.3-2.0 million acres planted annually
- ☀️ Rust is a top 5 sunflower disease.
- ☀️ Prevalence (2002-2015)
 - Range approximately 20-80%
 - Mean of about 40-50%



(Courtesy of North Central Integrated Pest Management Center)



(Photos: Zachary Ittel)



Sample Locations:

2023: 67

2024: 57

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.

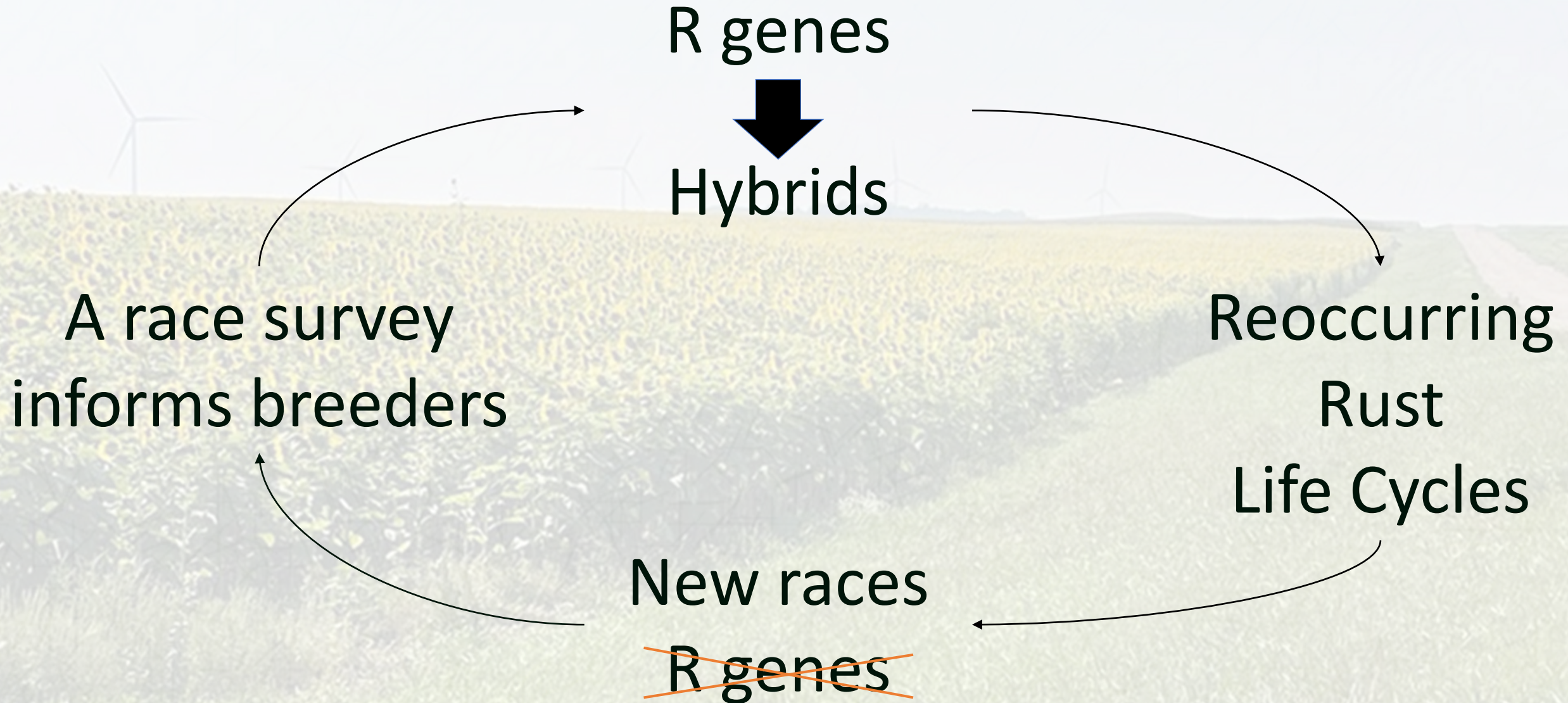
Introduction: Pathogen Virulence

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

Introduction: Pathogen Virulence

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Virulence: Background



(Photo: Zachary Ittel)

Virulence: Background





P. helianthi Virulence in 2011-2012

- ☀ 238 single pustule isolates
 - 29 races
 - Wide diversity of virulence
- ☀ 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



(Friskop et al. 2015)



P. helianthi Virulence in California during 2017-2018

Production fields:

- ☀️ 2017 – 6 races (11 SPI)
- ☀️ 2018 – 8 races (13 SPI)
- 🐞 CA had rust that could overcome all known R genes
- 🐞 Wide diversity of virulence

Wild sunflower:

- ☀️ 2017 – 6 races (11 SPI)
- ☀️ 2018 – 16 races (20 SPI)



Dr. Brandt Berghuis



Virulence: Methods (Pathogen Recovery)



(Photo: Denis Colombo)



(Photo: Zachary Ittel)



(Photo: Kaylie Schlect)



(Photo: Zachary Ittel)



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	30
South Dakota	8/8	15/15	45	18
Wisconsin	6/6	6/6	33	6
Nebraska	10/10	...	30	0
Minnesota	...	5/5	3	2
Totals	65/67	57/57	234	56

Determining Virulence Phenotype (Race)

A photograph of a large greenhouse with a high, arched glass roof. Inside, numerous rows of young plants in black pots are arranged on long, black metal shelving units. The plants are small and green, with some showing signs of stress or damage. The greenhouse is well-lit by natural light coming through the glass walls and roof. The overall scene suggests a controlled environment for plant research or cultivation.

(Photo: Zachary Ittel)

Infection Type (IT)



0



1 ($<0.2\text{mm}$)



2 ($0.2-0.4\text{mm}$)



3 ($0.4-0.6\text{mm}$)



4 ($0.6-0.8\text{mm}$)



5 (0.8mm)

Universal Differential Set

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

(Friskop et al. 2015; Gulya and Masirevic 1996)

Triplet	Differential	Resistance Gene	Scoring Value	Code
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	HA-R1	R_{4a}	2	
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	HA-R1	R_{4a}	2	
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	HA-R4	R_{4c}	2	
	HA-R5	R_{4d}	4	

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	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	7
	HA-R4	R_{4c}	2	
	HA-R5	R_{4d}	4	

Race Name =

347

Universal Differential Set

Triplet	Differential	Resistance Gene	Scoring value
One	7350	...	1
	MC90	R_1	2
	MC29	$R_2 \dots R_{10}$	4
Two	P386	R_{4a}	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

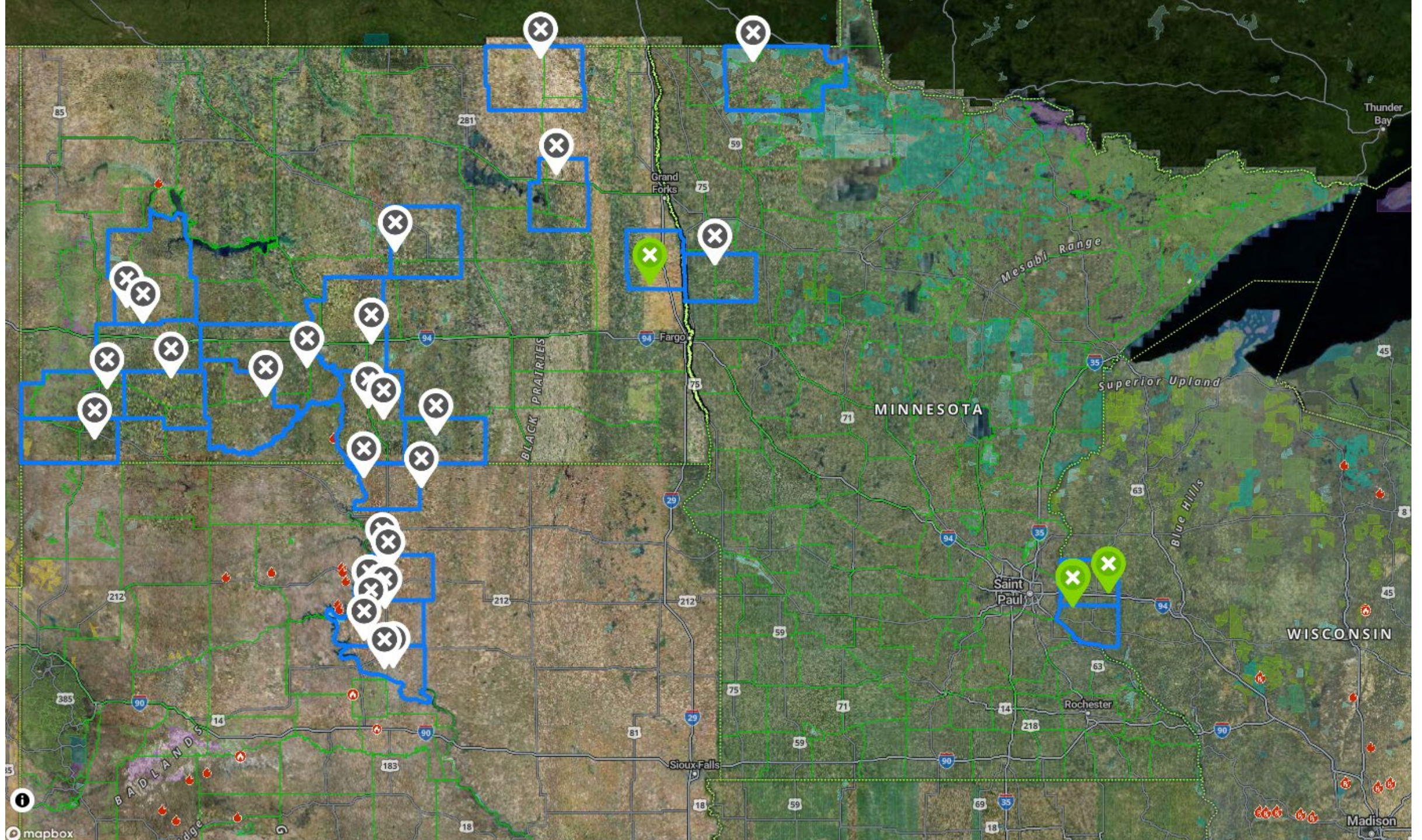
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	HA-R1	R_{4a}	2	
	HA-R2	R_5	4	
Three	HA-R3	R_{4b}	1	
	HA-R4	R_{4c}	2	
	HA-R5	R_{4d}	4	
Four	?	?	?	?
	?	?	?	
	?	?	?	

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

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Three	HA-R3	R_{4b}	1
	HA-R4	R_{4c}	2
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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



Differential	Resistance Gene	Virulent/Screened	% Virulent
7350	...		
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MC29	R_2+R_{10}		
P386	R_{4e}		
HA-R1	R_{4a}		
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RHA 464	R_{12}		
HAR-6	R_{13a}		
HAR-8	R_{15}		
HAR-18	R_{17}		
HAR-19	R_{18}		

Differential	Resistance Gene	Virulent/Screened	% Virulent
7350	...	33 / 33	
MC90	R_1	33 / 33	
MC29	R_2+R_{10}	2 / 28	
P386	R_{4e}	16 / 31	
HA-R1	R_{4a}	22 / 33	
HA-R2	R_5	9 / 29	
HA-R3	R_{4b}	3 / 27	
HA-R4	R_{4c}	7 / 23	
HA-R5	R_{4d}	31 / 33	
HAR-9	R_{11}	4 / 14	
RHA 464	R_{12}	5 / 25	
HAR-6	R_{13a}	0 / 32	
HAR-8	R_{15}	1 / 33	
HAR-18	R_{17}	3 / 32	
HAR-19	R_{18}	14 / 32	

Differential	Resistance Gene	Virulent/Screened	% Virulent
7350	...	33 / 33	100%
MC90	R_1	33 / 33	100%
MC29	R_2+R_{10}	2 / 28	7%
P386	R_{4e}	16 / 31	52%
HA-R1	R_{4a}	22 / 33	67%
HA-R2	R_5	9 / 29	31%
HA-R3	R_{4b}	3 / 27	11%
HA-R4	R_{4c}	7 / 23	30%
HA-R5	R_{4d}	31 / 33	94%
HAR-9	R_{11}	4 / 14	29%
RHA 464	R_{12}	5 / 25	20%
HAR-6	R_{13a}	0 / 32	0%
HAR-8	R_{15}	1 / 33	3%
HAR-18	R_{17}	3 / 32	9%
HAR-19	R_{18}	14 / 32	44%

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Race Comparison

31406

32400

32406

32424

33400

33440

33440

33700

37400

37410

37737

74010



Race Comparison

31406

32400

32406

32424

33400

33440

33440

33700

37400

37410

37737

74010



Race Comparison

31406

32400

32406

32424

33400

33440

33440

33700

37400

37410

37737

74010

Differential	Resistance Gene	2011-2012	2017-2018	2023-2024
7350	...	100%	100%	100%
MC90	R_1	100%	92%	100%
MC29	R_2+R_{10}	16%	28%	7%
P386	R_{4e}	26%	78%	52%
HA-R1	R_{4a}	40%	84%	67%
HA-R2	R_5	21%	2%	31%
HA-R3	R_{4b}	4%	42%	11%
HA-R4	R_{4c}	33%	72%	30%
HA-R5	R_{4d}	98%	74%	94%
HAR-9	R_{11}	-	-	29%
RHA 464	R_{12}	-	-	20%
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Race Comparison

2011-2012

- ☀️ 238 isolates
- ☀️ 29 races
- ☀️ Virulence on 9/9 differentials
- ☀️ Most common:
 - 🐞 300 & 304

2017-2018

- ☀️ 55 isolates
- ☀️ 22 races
- ☀️ Virulence on 9/9 differentials
- ☀️ Most common:
 - 336

2023-2024*

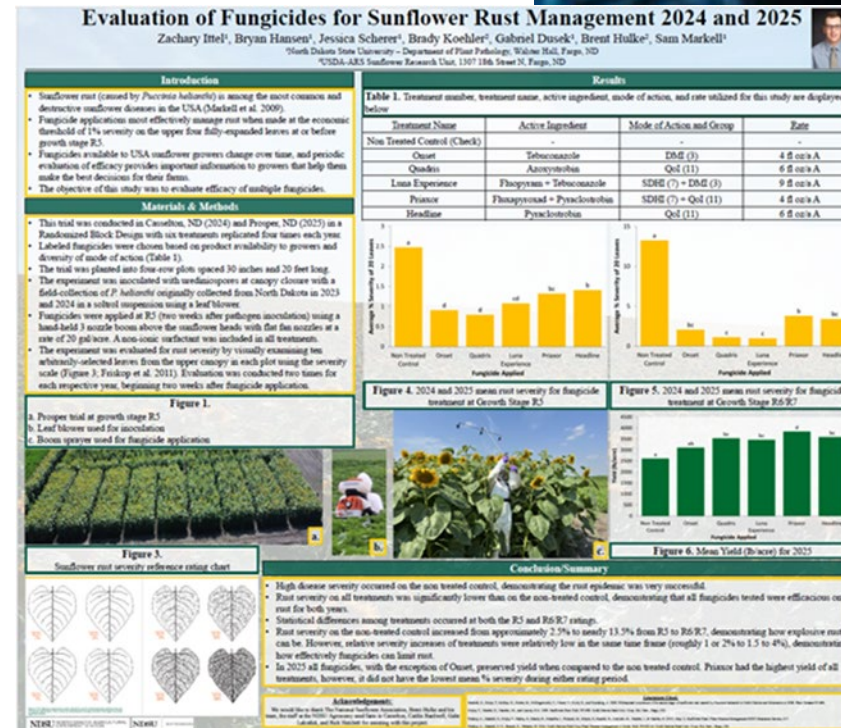
- ☀️ 31 isolates
- ☀️ Virulence on 14/15 differentials
- Most common:
 - 33400



Additional/Future Objectives



Puccinia helianthi population genetic analysis

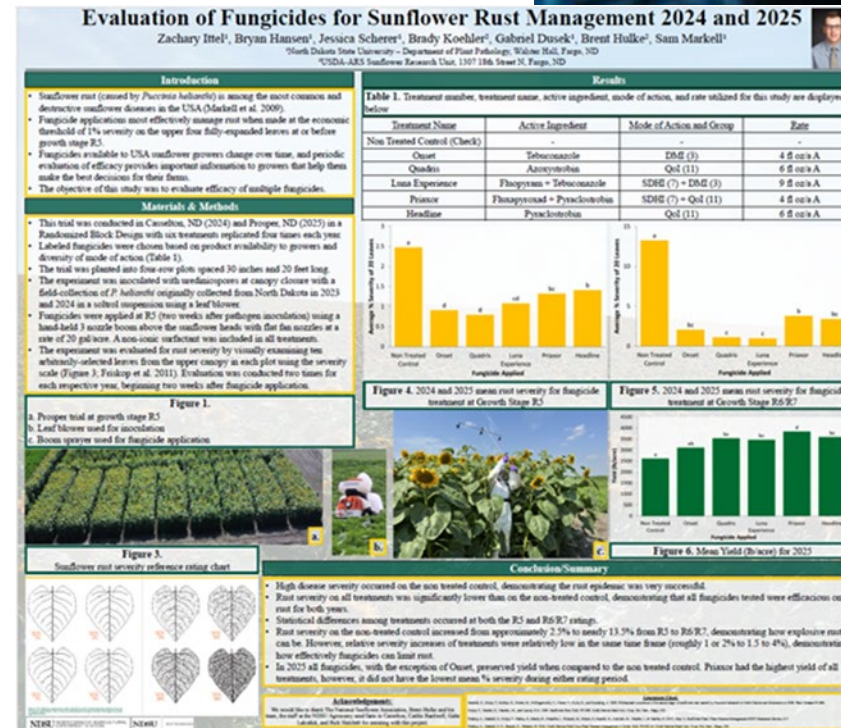




Additional/Future Objectives

☀️ *Puccinia helianthi*
population genetic
analysis

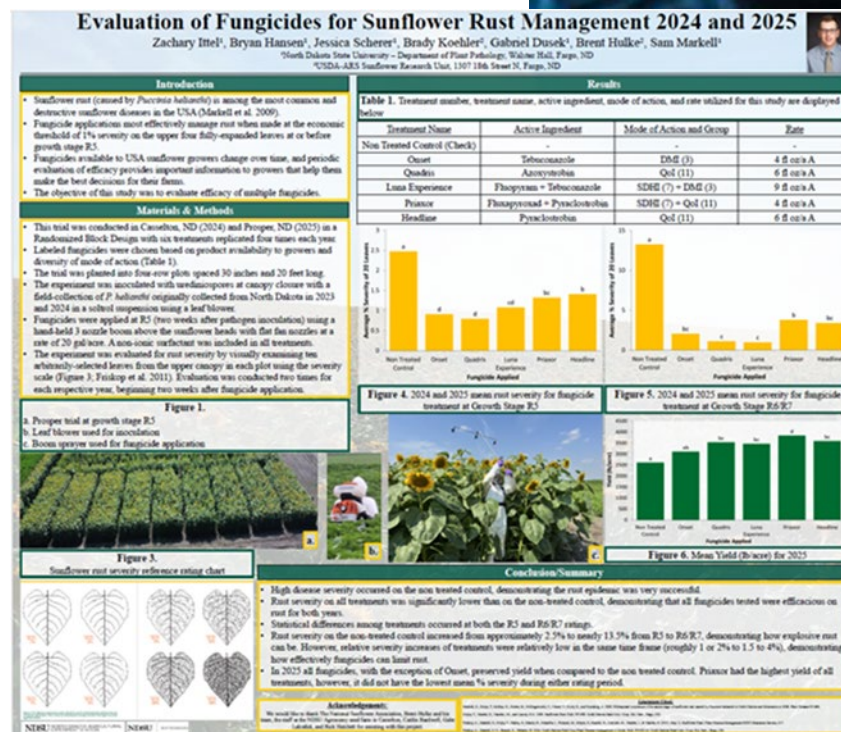
☀️ Whole Genome
Pathogen Sequencing





Additional/Future Objectives

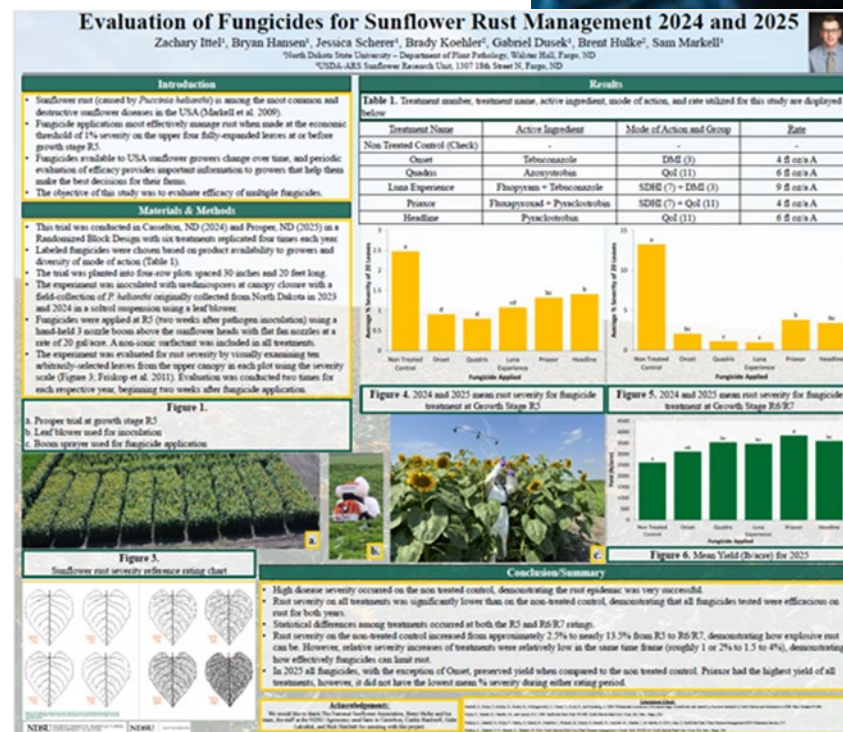
- ☀️ *Puccinia helianthi* population genetic analysis
- ☀️ Whole Genome Pathogen Sequencing
- ☀️ SAM pop experiment





Additional/Future Objectives

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





Project Update Summary

- ☀ Prevalence = Does rust occur in a field? (Yes/Fields Surveyed)
- ☀ Virulence = Can the pathogen overcome a resistance gene?
- ☀ Future directions



Acknowledgements

Survey Help:

- ❖ **ND** (County Agents, NSA Board members, and Farmers)
- ❖ **SD** (Mr. Weber, NSA Board members, Farmers, and Dr. Febina Mathew's Team)
- ❖ **NE** (Dr. Bob Harveson and his team)
- ❖ **WI** (Dr. Brandt Berghuis and his team)
- ❖ **MN** (Farmers)

Other Pivotal Help:

- ❖ Dr. Bill Underwood
- ❖ Scott Radi
- ❖ John Sandbakken
- ❖ Kaylie Schlecht
- ❖ Caitlin Bardwell
- ❖ Gabe Lakoduk
- ❖ Rick Hatchet
- ❖ Jessica Scherer
- ❖ Bryan Hansen



A man wearing a tan sun hat with a logo, sunglasses, and a dark green shirt is smiling. He is holding a large sunflower in front of him. The background is a field of sunflowers under a cloudy sky.

Thank you

Questions?