Genetic mapping of resistance to Aphanomyces root rot in alfalfa

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Aphanomyces root rot



Seedling "corpses"

<u>Acute phase</u>



Adult plants lack lateral and fibrous roots, no nodules. Foliage stunted, yellow. <u>Chronic phase</u>.

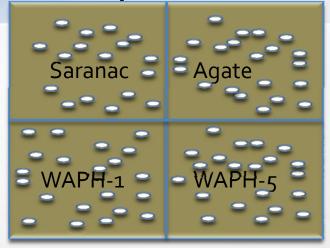
Aphanomyces root rot

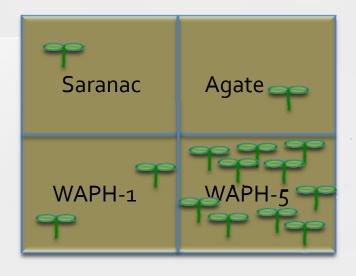


Research questions

- Survey of MN and NY soils for APH and PRR
 - What races are found in alfalfa fields?
- Two races identified by host genetics
 - Can markers be identified linked to resistance for marker assisted breeding?
- Possible failure of race 1+2 resistant cultivars
 - Are there additional races?

Bioassay for APH

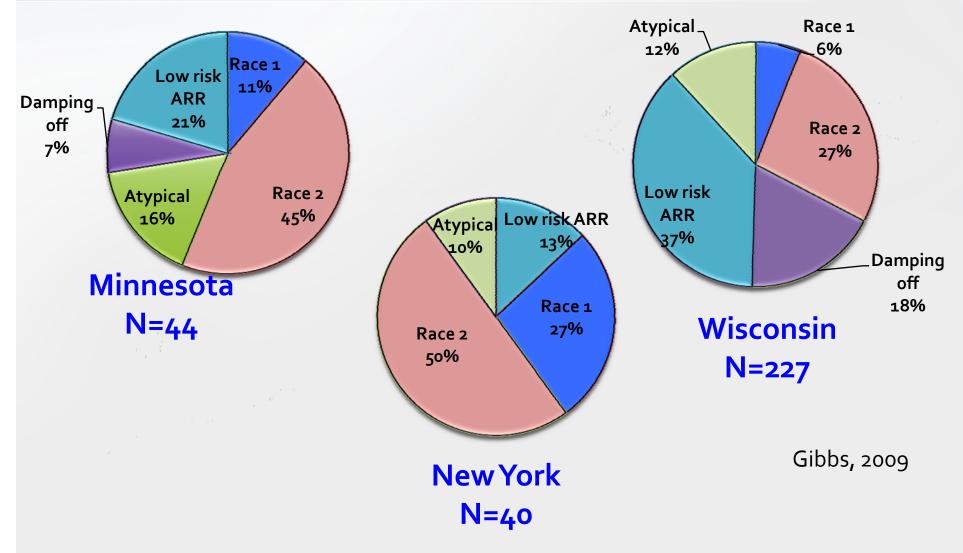




Plant checks in field soil sample. After plants germinate, flood soil for 4 days. Score symptoms at 21 d after planting.

Cultivar	PRR	APH Race 1	APH Race 2
Saranac	S	S	S
Agate	R	S	S
WAPH-1	R	R	S
WAPH-5	R	R	R

Race 2 is most common



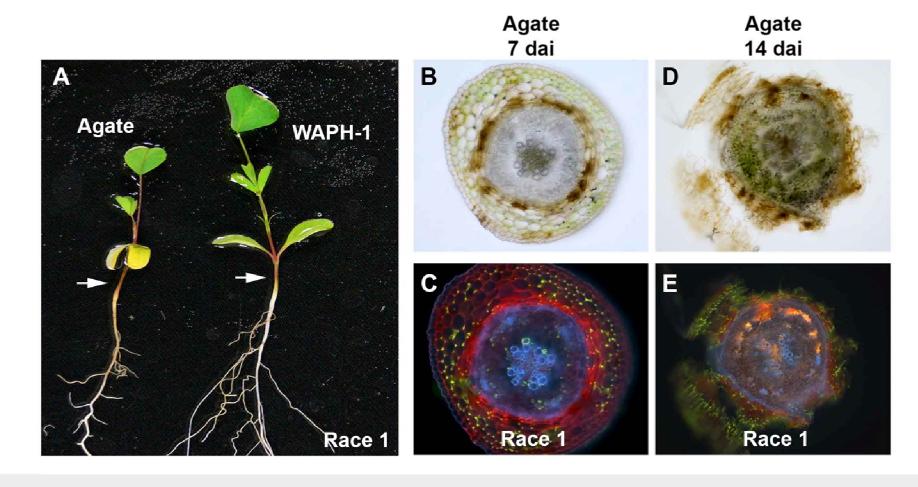
Results from atypical soils

State	Soil	WAPH-1 resistant plants (%)	WAPH-5 resistant plants (%)	Predominant race
MN	GRE	6	15	Undetermined
MN	JIM	0	5	Undetermined
MN	WAS	14	28	Undetermined
NY	CAR	13	32	Undetermined
NY	ROY	0	21	Undetermined

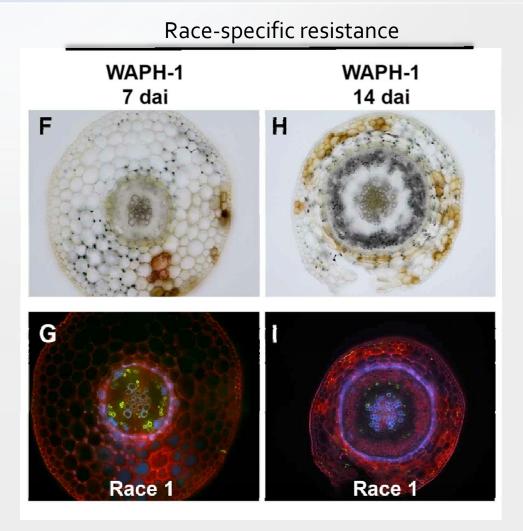
60 pure Aphanomyces strains tested

- 4 check cultivars, 4-6 reps, 30 seeds/rep
- All strains read as either race 1 or race 2
- Resistance in WAPH-5 was not overcome
- What is the nature of resistance?

APH Susceptible Reaction



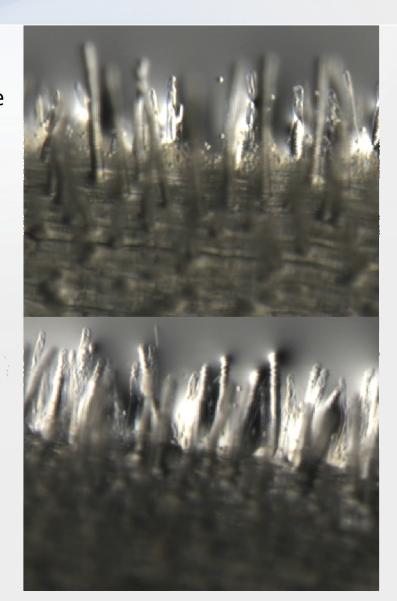
APH Resistant Reaction



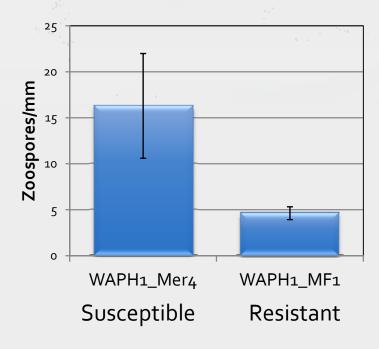
Hypersensitive response indicates defense is triggered by NBS-LRR type resistance gene.

Pre-infection resistance?

Susceptible



Resistant



Developing MAS for APH

- Selected resistant and susceptible seedlings
 - WAPH-1 (race 1), WAPH-5 (race 1+2), 53V54 (race 1+2)
 - Retested with vegetative cuttings
- Symptoms correlated with oospores counts and Aphanomyces DNA content

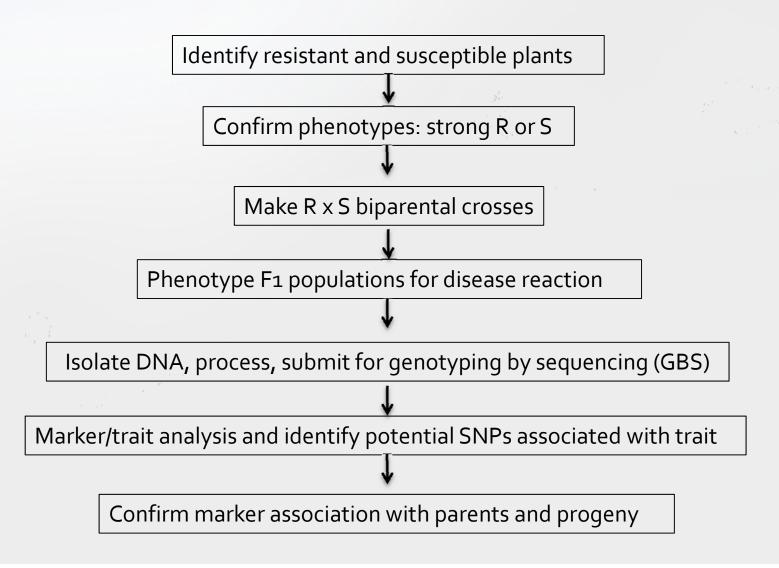


Resistant

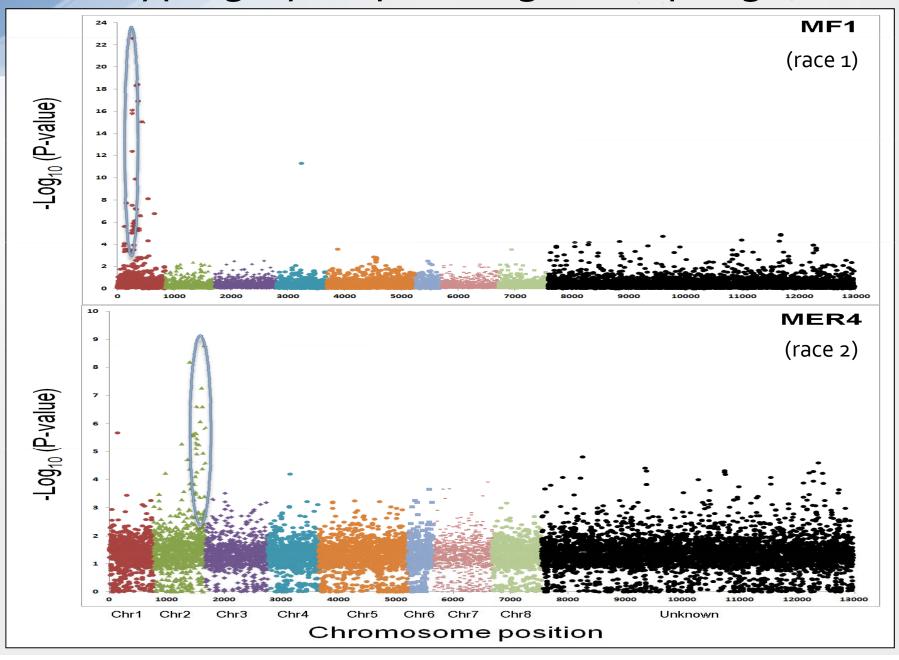


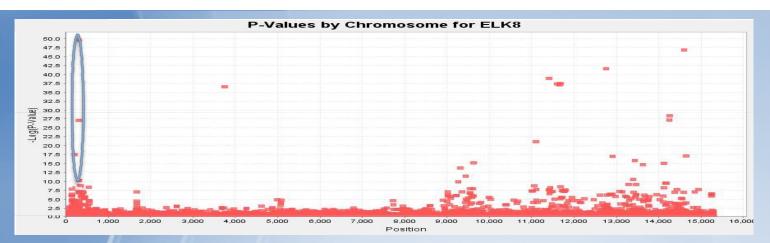
Susceptible

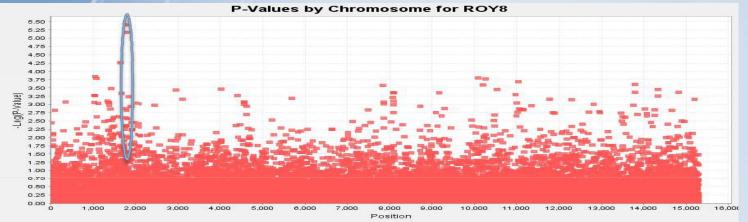
Identify DNA markers associated with racespecific resistance to APH

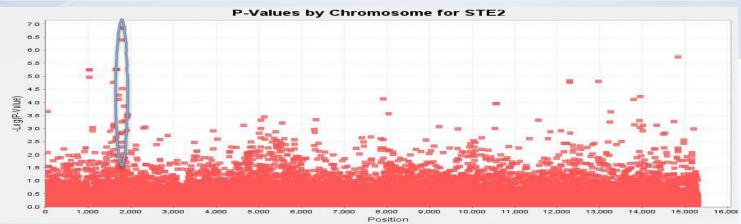


Genotyping by sequencing to map R genes









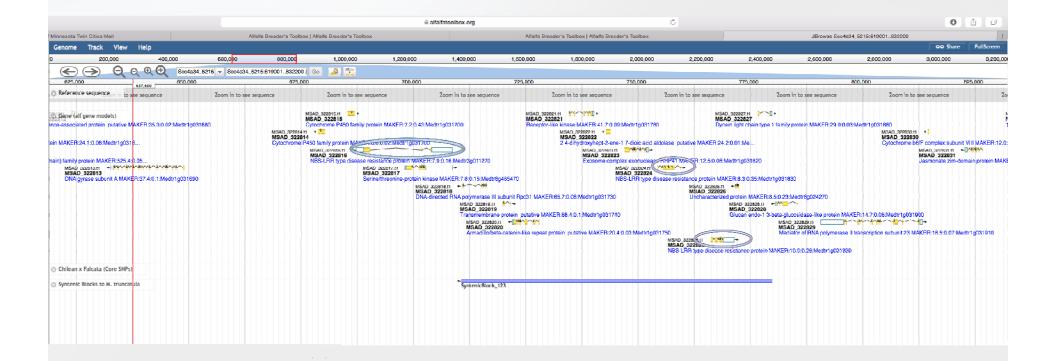
Reaction of strains on F1 population (L85)

MER4	GRE ₅	ROY8	STE ₂	JIM1	Number of plants (of 269)
R	R	R	R	R	2
S	S	S	S	S	16
R	S	S	S	S	6
S	R	S	S	S	2
S	S	R	S	S	5
S	S	S	R	S	2
S	S	S	S	R	3

Suggests that there are multiple R genes in WAPH-5.

Intercross plants with resistance to a single strain to develop new checks.

Genes near MF-1 (race 1) marker



Pennsylvania field soil Flooded for 4 days (6-9 days after planting) Photo at 12 days after planting

Commercial race 1 resistant variety

WAPH-5 race 1+2 resistant check



Susceptible checks

Commercial race 1+2 resistant variety

WAPH-1 race 1 resistant check

Conclusions

- Resistance appears to be conditioned by dominant R-genes with additional pre- and post-infection defense mechanisms
- R-genes may be clustered on chromosome 1 and 2
- WAPH-5 may have multiple R-genes for APH
- Strains of *A. euteiches* with diverse virulences are found in field soil
- Developers of resistant varieties should use diverse isolates to have sufficient field resistance to Aphanomyces root rot

