

Keeping a finger on the pulse of pulse crop pathology

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UNIVERSITY OF SASKATCHEWAN
College of Agriculture
and Bioresources
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Government
of
Saskatchewan
Ministry of Agriculture



Chocolate spot of faba bean



Ascochyta pisi on pea



Ascochyta blight of chickpea



Sclerotinia white mould (l) and botrytis grey mould (r) on lentil



Stemphylium blight on lentil

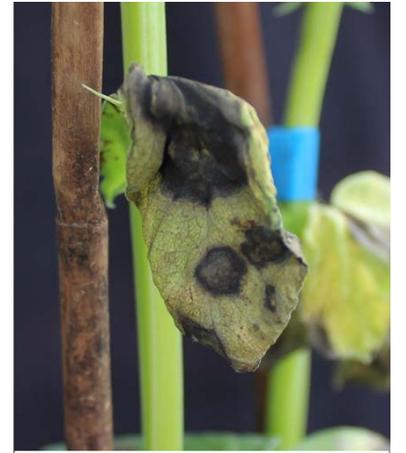
Root rots



Alternaria leaf spot on faba



Mycosphaerella blight of pea



Stemphylium blight on faba

Ascochyta blight of lentil



Anthracnose of bean



Halo blight of bean



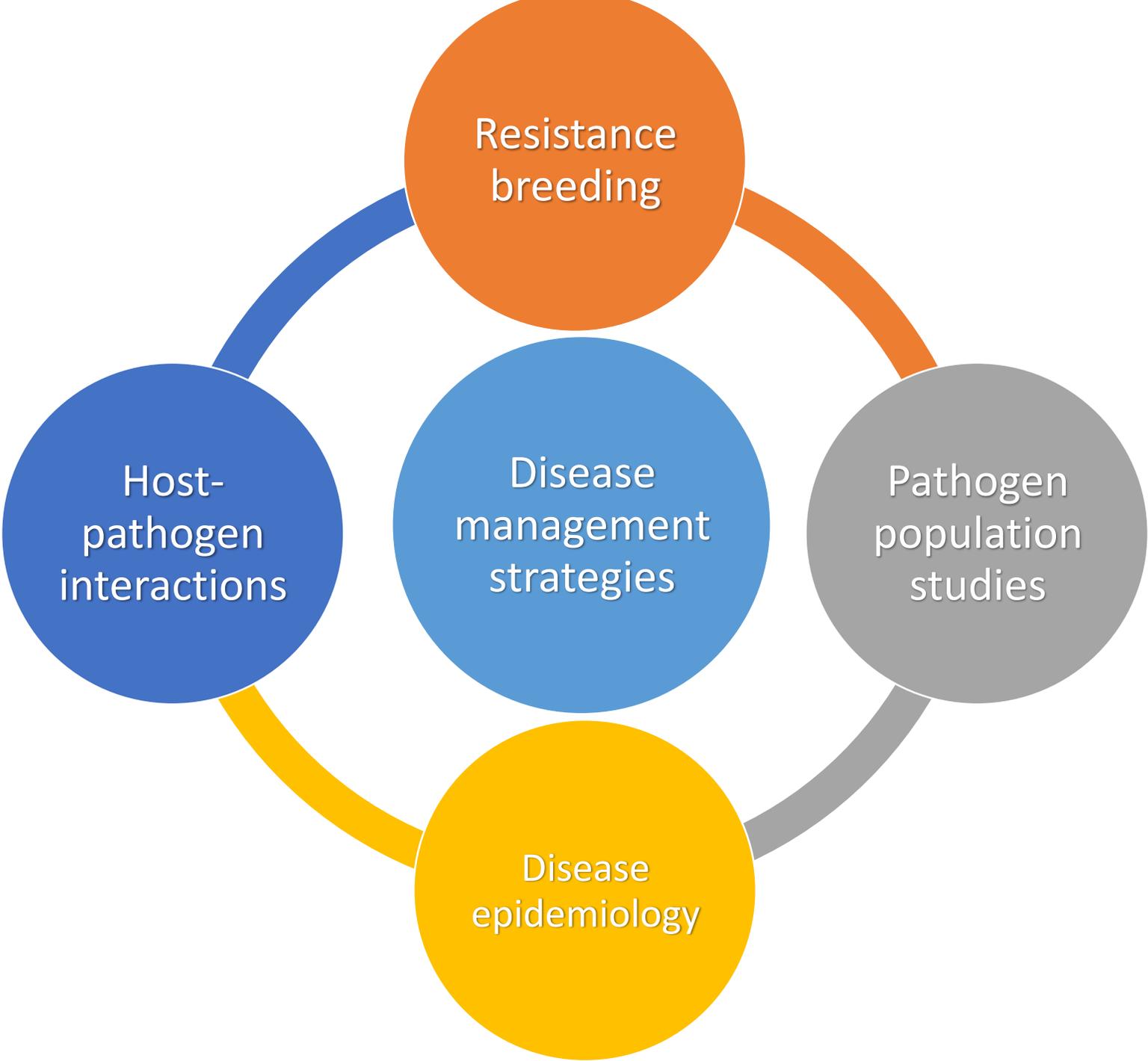
Anthracnose of lentil



Common bacterial blight of bean



Blossom blight





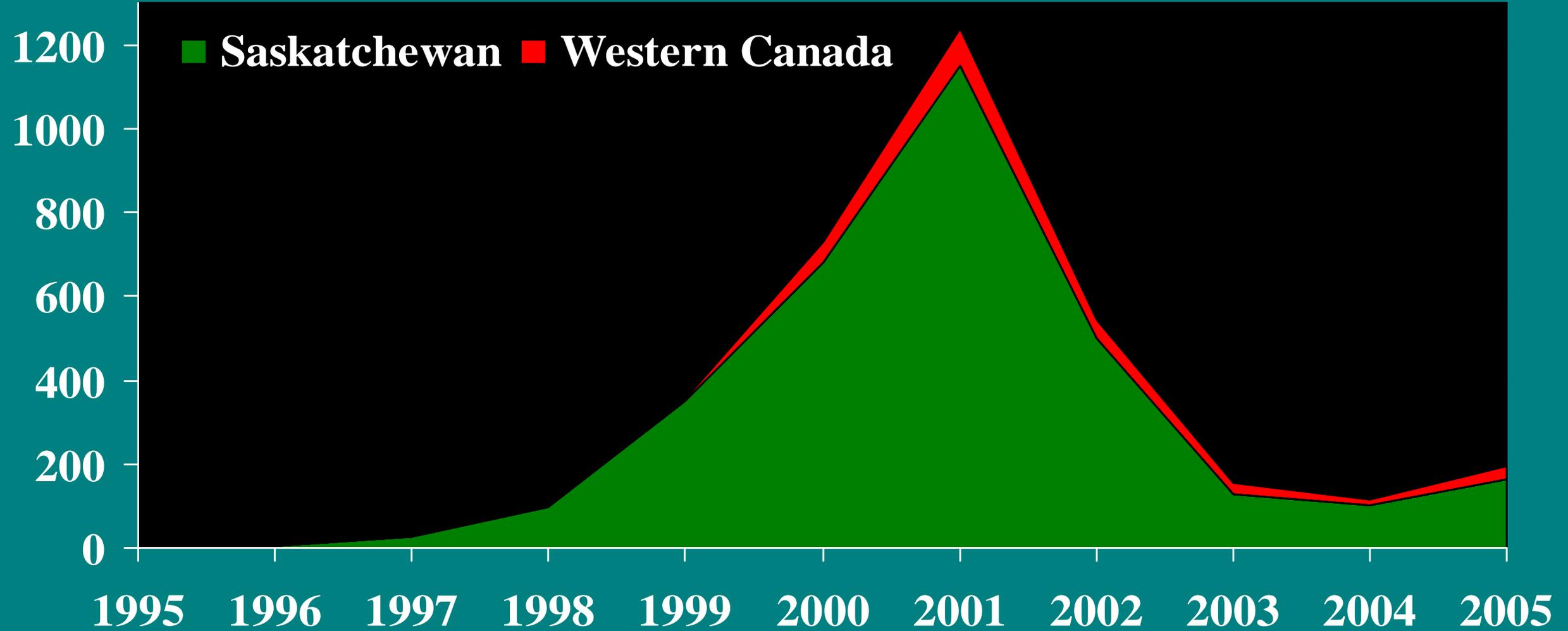
Management of ascochyta blight in chickpea in Canada

with Tom Wolf and Yantai Gan, AAFC

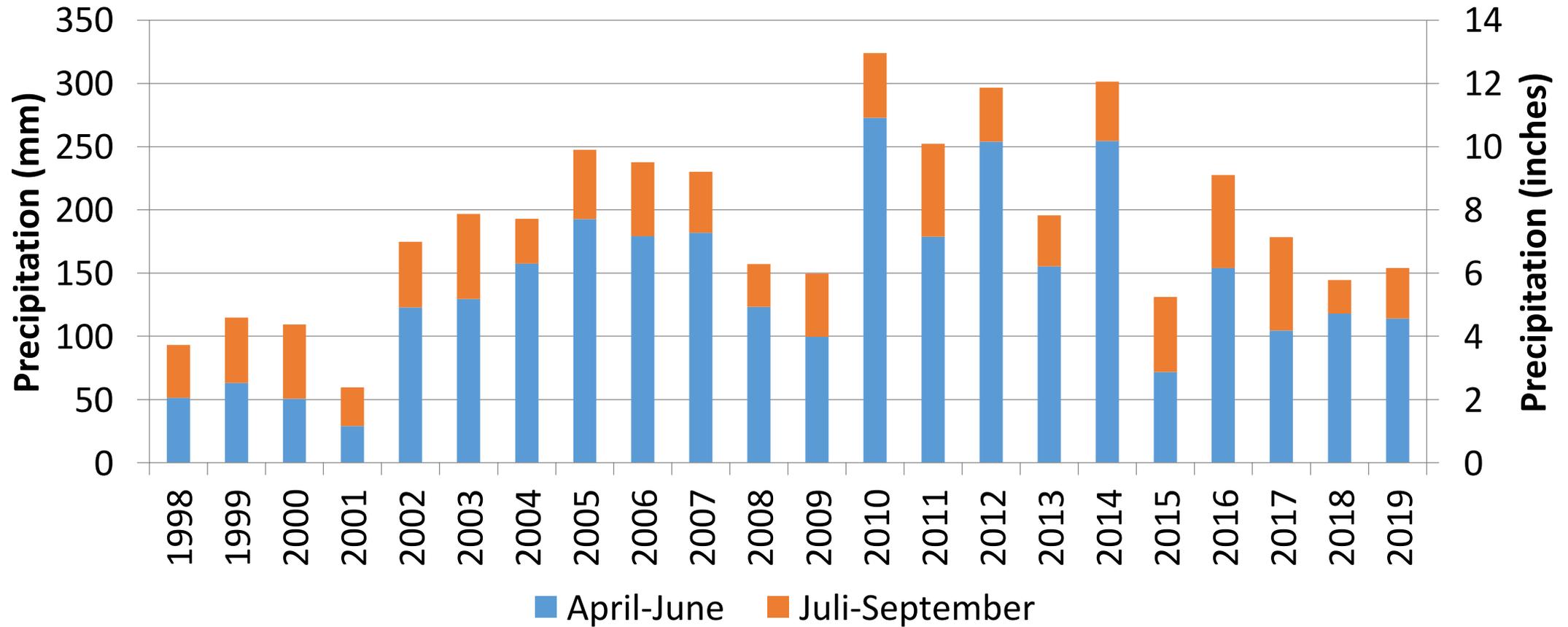
Chickpea area (seeded acres)

Source: Ray McVicar, SAF

Thousands



Epidemics of asochyta blight in chickpea after 2001



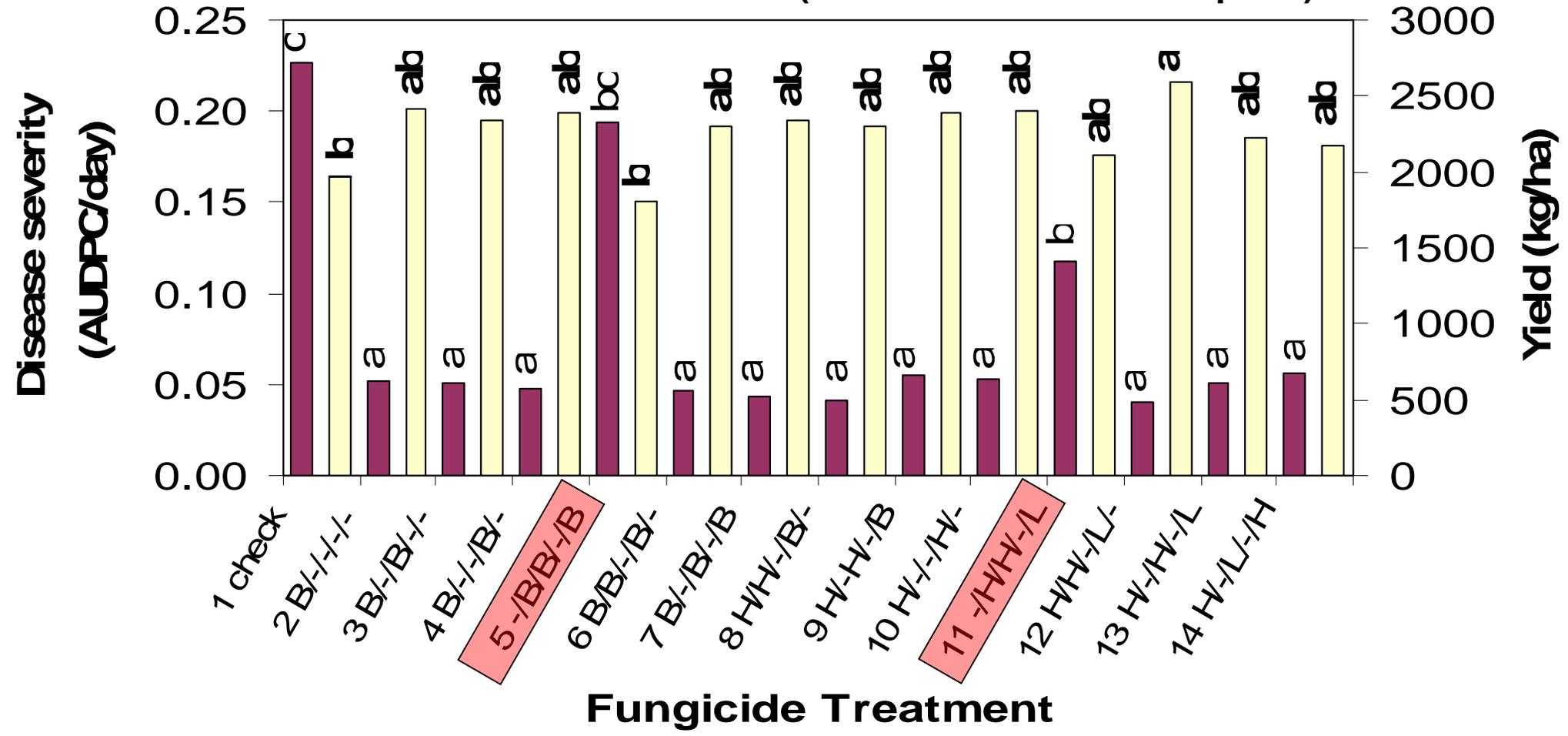


Timing of fungicide applications (Round 1)



■ Disease ■ Yield

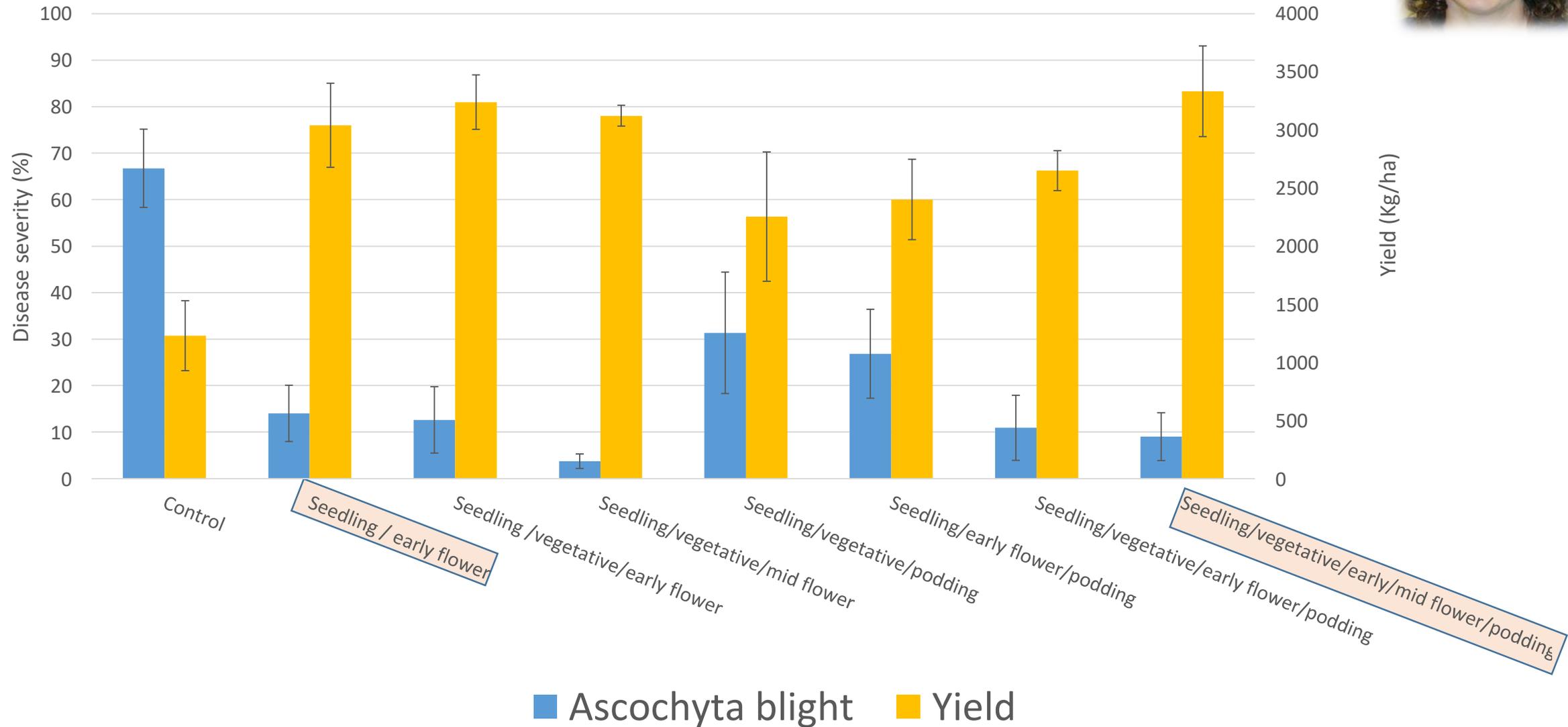
Saskatoon 2003 CDC Yuma (44% disease in control plots)





TIMING OF FUNGICIDE APPLICATIONS (Round 2)

CDC Yuma, Saskatoon, SK, 2008

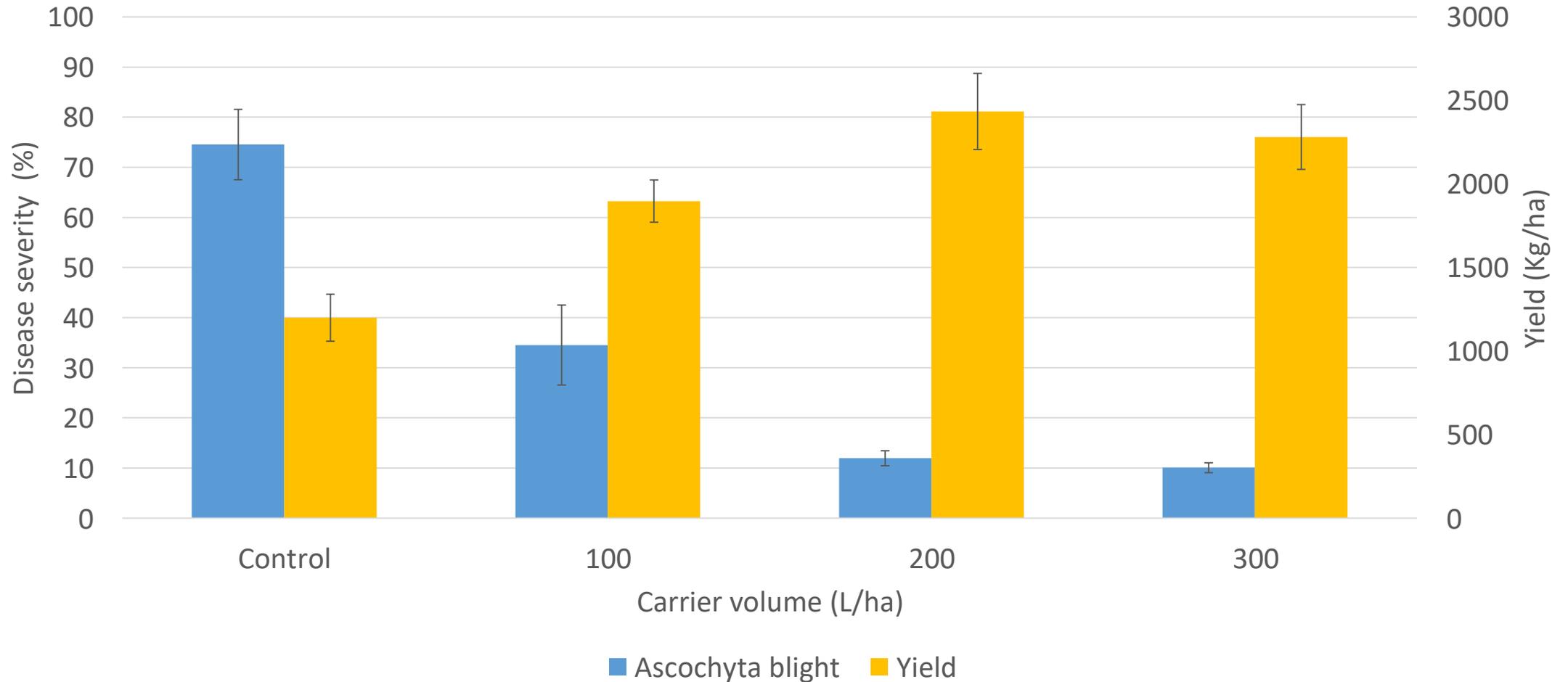




EFFECT OF CARRIER VOLUME

CDC Yuma, Outlook, SK, 2001

Armstrong-Cho et al.: *Crop Protection* 27 (2008), 1020-1030.

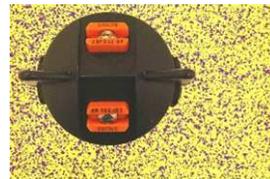
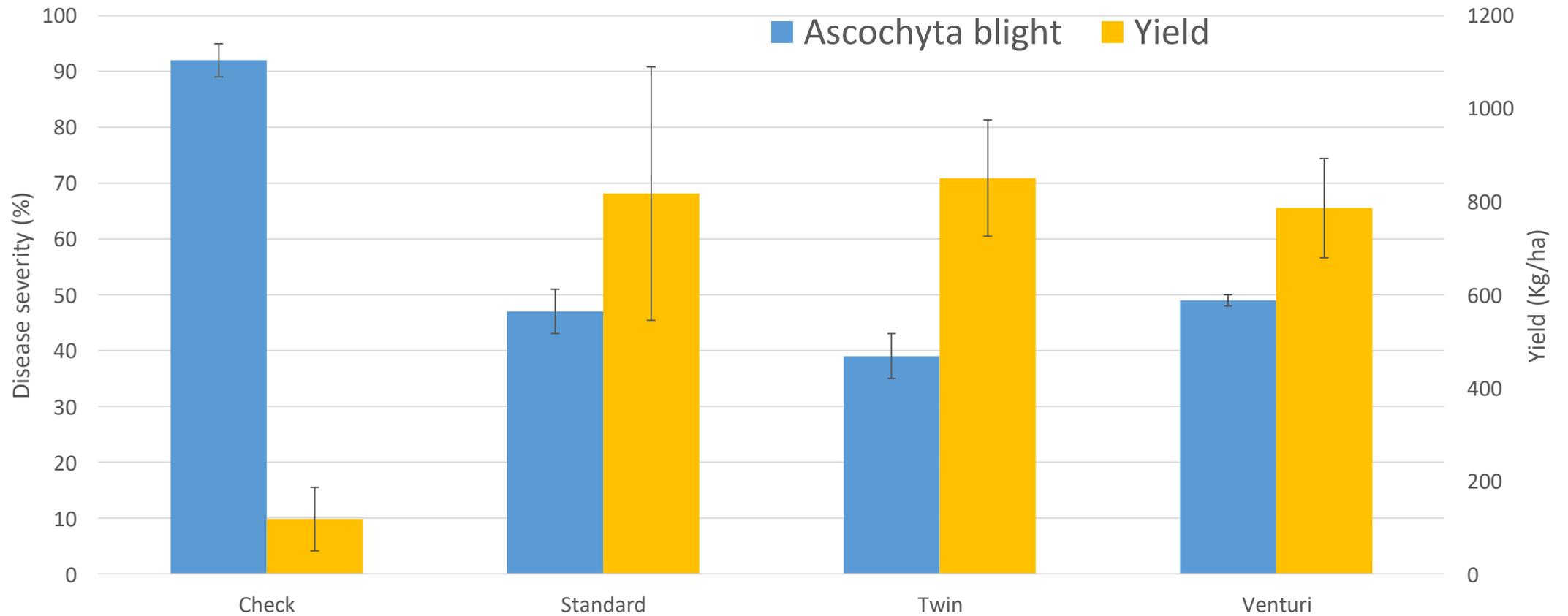




EFFECT OF DROPLET SIZE

Saskatoon, SK, 2002

Armstrong-Cho et al: Crop Protection 27 (2008), 700-709.



Scouting and Management of

Ascochyta Blight in Chickpea

Symptoms

Scouting

Disease Risk Ratings

Disease Cycle & Infection Process

Fungicide & Resistance Management

Fungicide Application Technology

Economic Thresholds

Determining your Risk Rating:

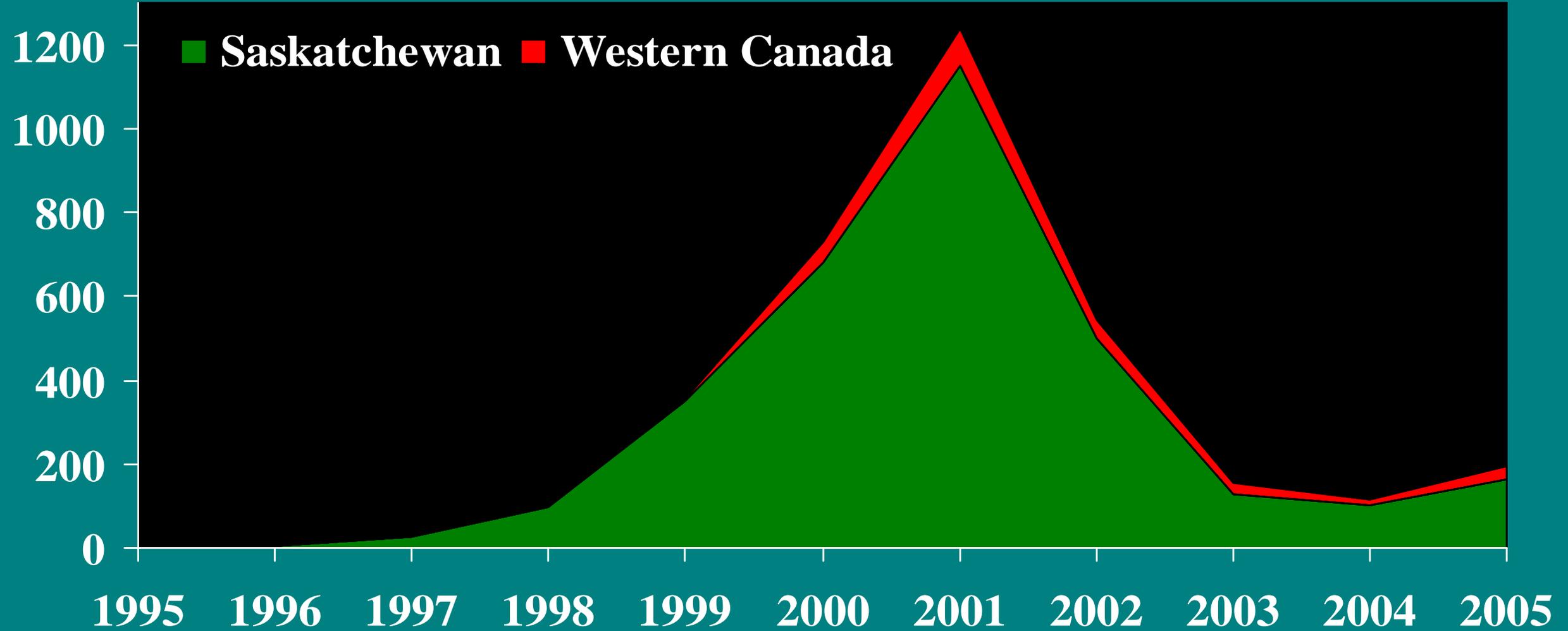
- Review the following six considerations and assign a risk value to each
- Add up the risk values to create a total risk value
- Use the total value to compare to the risk rating scale on the following page

	Risk Value
1. Field History and Crop Rotation <ul style="list-style-type: none"> a. Crop is being grown in a region that has never had chickpea production <input type="checkbox"/> 0 b. Crop is planted on land that has not had chickpea for at least 3 years <input type="checkbox"/> 5 c. Crop is planted on land that has had chickpea in the last 2 years; OR is located adjacent to chickpea stubble from the year before <input type="checkbox"/> 10 	
2. Chickpea Variety <ul style="list-style-type: none"> a. Desi variety OR kabuli variety rated as "fair" resistance to ascochyta blight (e.g. B-90 or Amit, CDC Frontier) <input type="checkbox"/> 5 b. Kabuli variety that is rated as "poor" <input type="checkbox"/> 10 c. Kabuli variety that is rated as "very poor" <input type="checkbox"/> 20 	
3. Level of Seed-borne Disease and Use of Seed Treatment <ul style="list-style-type: none"> a. Seed test indicated no seed-borne ascochyta AND used a registered seed treatment for ascochyta blight control <input type="checkbox"/> 0 b. Seed test indicated low levels of ascochyta (< 1%) AND used a registered seed treatment for ascochyta blight control <input type="checkbox"/> 5 c. Seed test indicated significant levels of ascochyta blight (5-10%) AND used a registered seed treatment for ascochyta blight control <input type="checkbox"/> 10 d. The seed quality is unknown, OR I am not using a seed treatment <input type="checkbox"/> 20 	
4. Presence of Disease Symptoms since last Fungicide Application <ul style="list-style-type: none"> a. No new disease lesions have developed since last fungicide application <input type="checkbox"/> 0 b. Disease lesions have developed on new crop growth since last fungicide application <input type="checkbox"/> 10 c. Leaf and/or stem lesions have developed and no fungicide has been applied this season <input type="checkbox"/> 20 	
5. Weather Conditions <ul style="list-style-type: none"> a. No rainfall in the past week and short-term forecast is for continued dry weather <input type="checkbox"/> 5 b. Weather conditions are unknown <input type="checkbox"/> 10 c. Rainfall or heavy dew has occurred during past week <input type="checkbox"/> 20 d. Weather is unsettled; thunderstorms are likely <input type="checkbox"/> 20 	
6. Other Crop Health Considerations <ul style="list-style-type: none"> a. Crop emerged well in the spring and there has been no significant weather/injury to crop <input type="checkbox"/> 0 b. Crop was seeded very early and was slow to emerge <input type="checkbox"/> 5 c. Crop was damaged by early herbicide application or soil-residue <input type="checkbox"/> 10 d. The crop has received a light to moderate hail shower in the past 24 hours <input type="checkbox"/> 10 	
TOTAL RISK VALUE (1 + 2 + 3 + 4 + 5 + 6) =	

Chickpea are (seeded acres)

Source: Ray McVicar, SAF

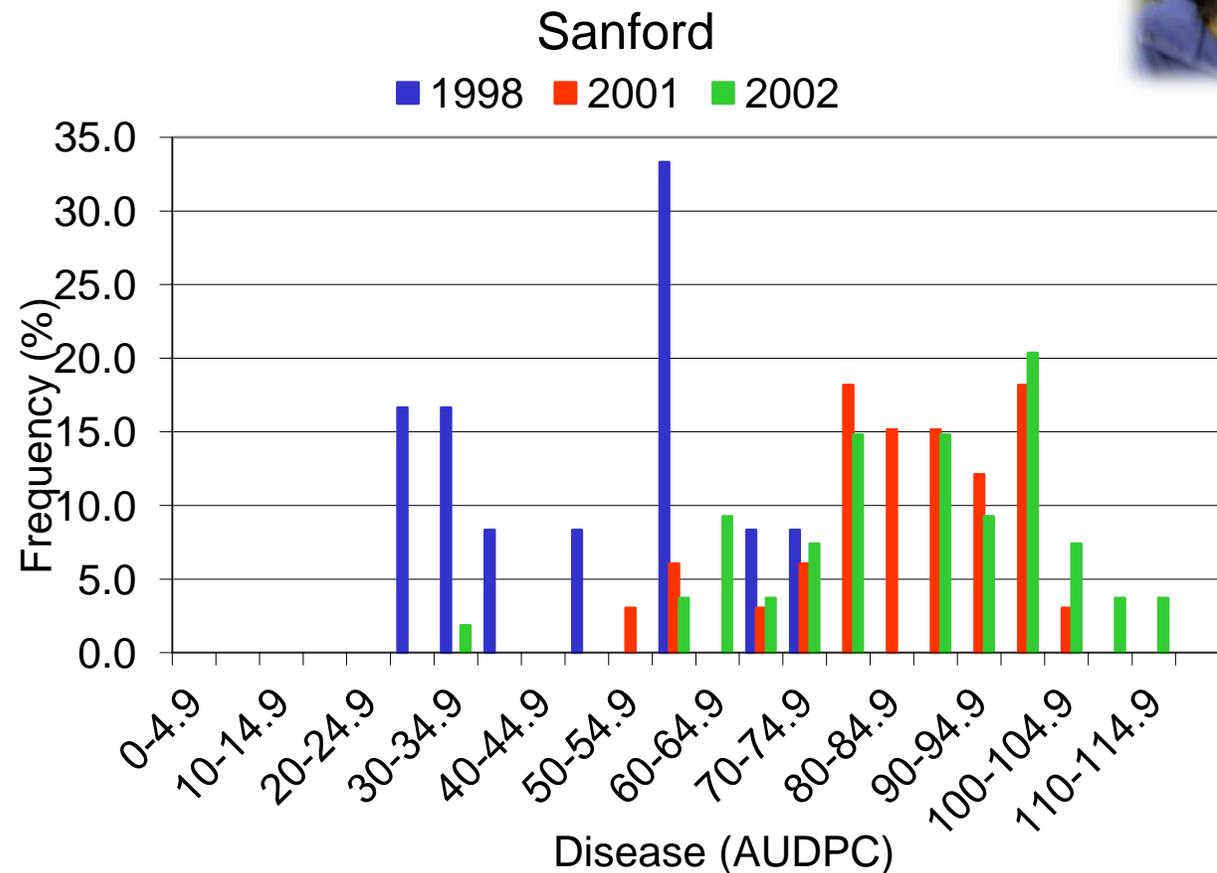
Thousands



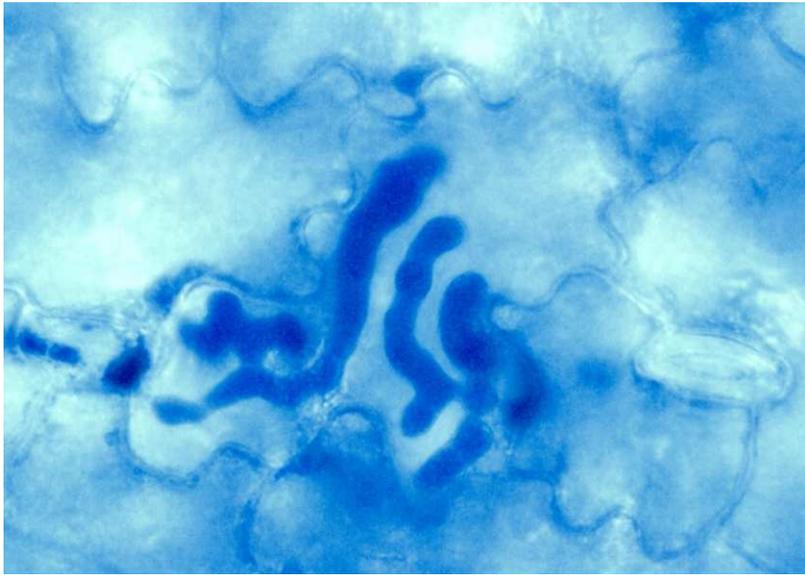


Resistance breakdown in chickpea

- Large kabuli chickpea Sanford considered partially resistant in 1998
- Dominated acreage because of premium price
- Rapid shift in virulence of *Ascochyta rabiei* within 3 years



Vail, S. and Banniza, S., 2008. Structure and pathogenic variability in *Ascochyta rabiei* populations on chickpea in the Canadian prairies. *Plant Pathology*, 57(4), pp.665-673.



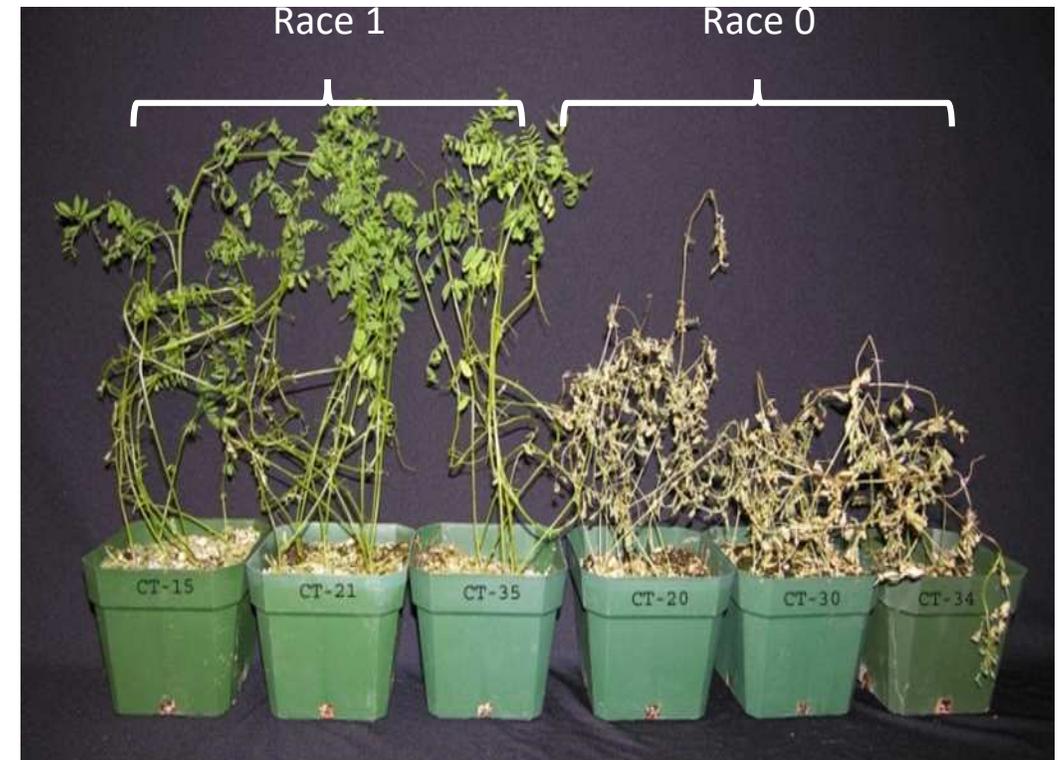
Understanding virulence in *Colletotrichum lentis* and anthracnose resistance in lentil

Colletotrichum lentis, the hemibiotrophic pathogen



- 2 **pathogenic** races previously described (Buchwaldt et al 2004)
- Both races undergo biotrophic phase before switching to necrotrophic phase
- Quantitative differences in the infection process between the two races

Armstrong-Cho, C., Wang, J., Wei, Y. and Banniza, S., 2012. The infection process of two pathogenic races of *Colletotrichum truncatum* on lentil. *Canadian Journal of Plant Pathology*, 34(1), pp.58-67.

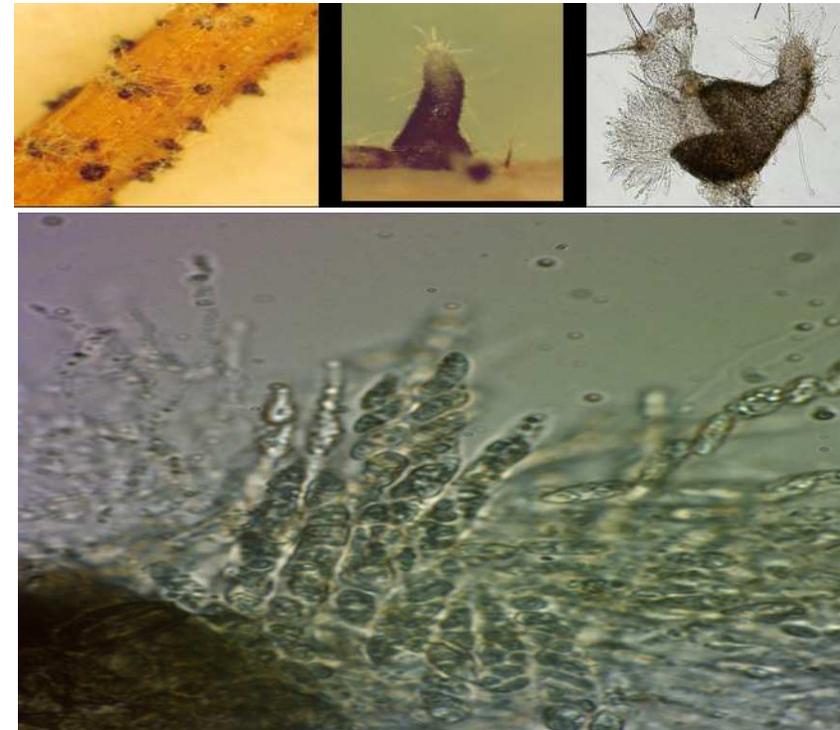




Sexual phase of *Colletotrichum lentis*



- Sexual phase induced in the laboratory
- Two sexual incompatibility groups (IG)
- F₁ segregates in a 1:1 ratio for IG
- IG-2 more common in the field
- Both IGs on the same plant

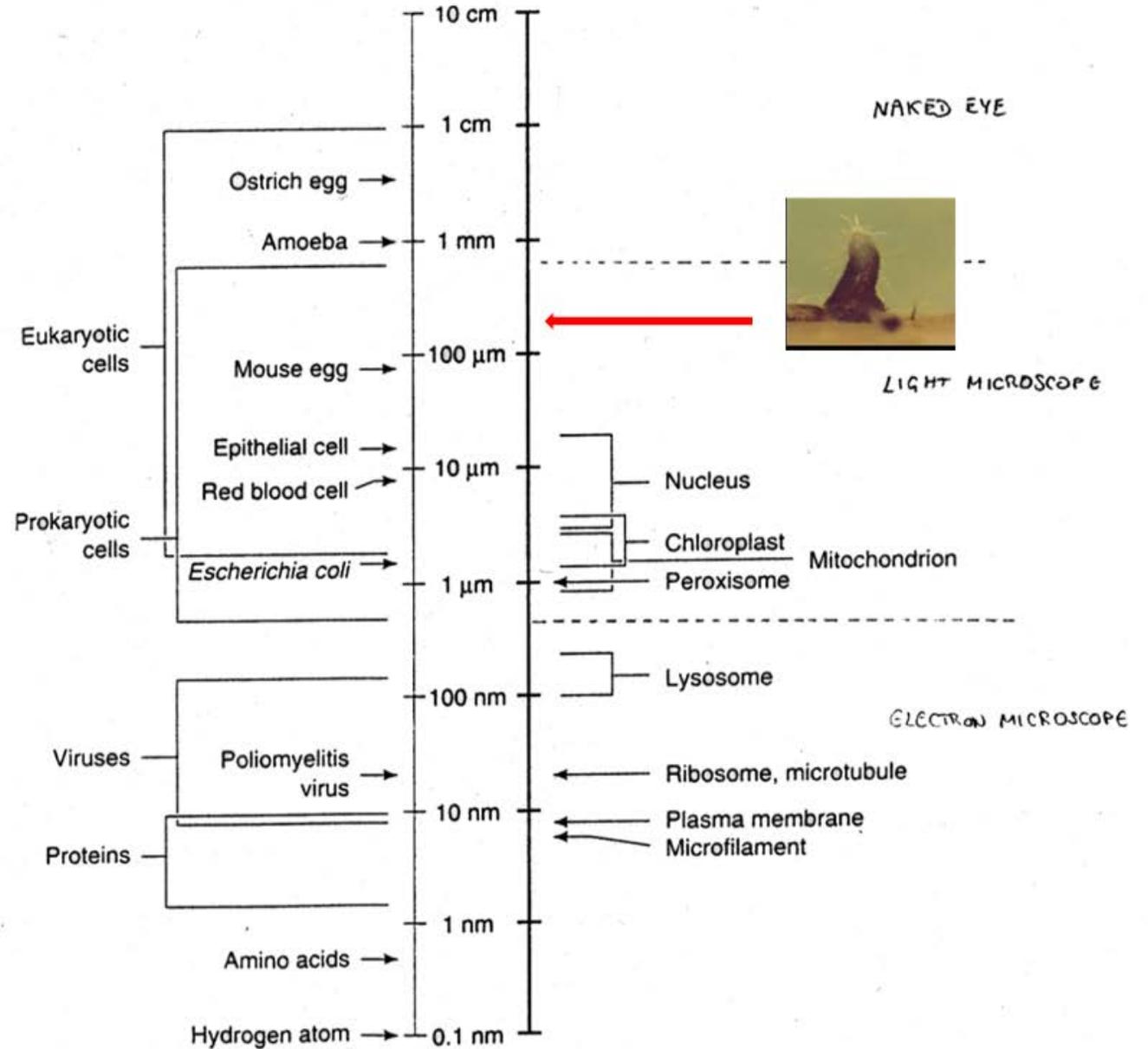


Menat et al. 2015. Fungal Ecology, in press.
Menat & Banniza 2012. Mycologia 104, 641-649.
Menat 2012. PhD thesis, University of Saskatchewan.
Armstrong-Cho & Banniza 2006. Mycological Research 110, 951-956.

Does *Colletotrichum lentis* have sexual reproduction in the field?



mashable.com

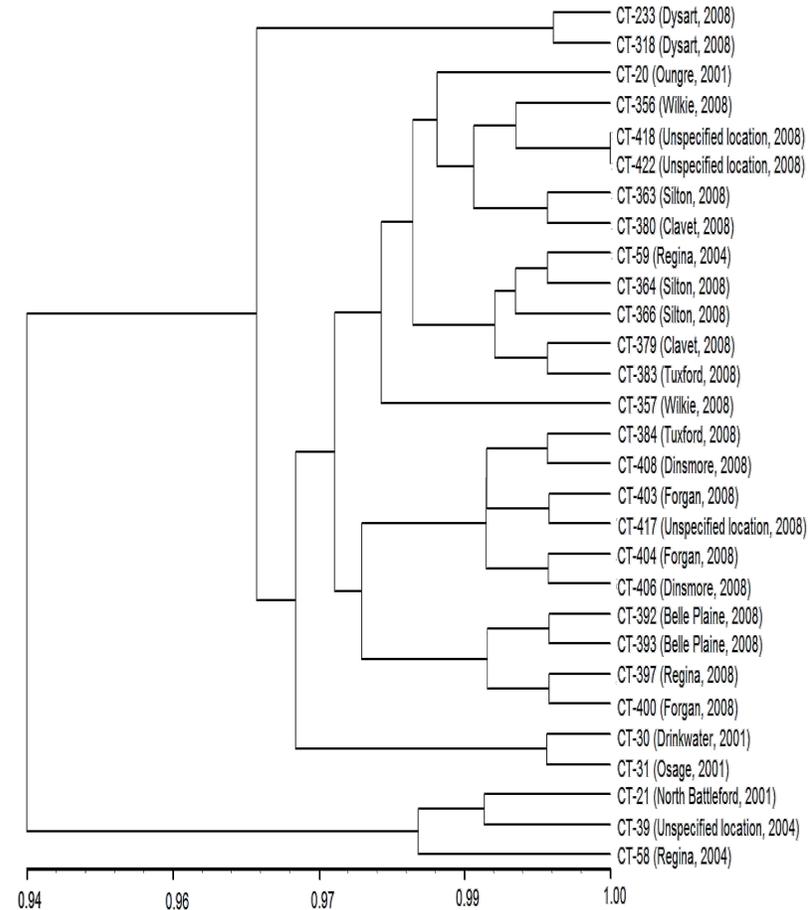




Does *Colletotrichum lentis* have sexual reproduction in the field?

- High levels of genetic similarity ($\geq 94\%$ based on AFLP markers) in field population
- Same haplotype (n=131) in one lentil field
- Indication of linkage disequilibrium

Menat, J., Armstrong-Cho, C. and Banniza, S., 2016. Lack of evidence for sexual reproduction in field populations of *Colletotrichum lentis*. *Fungal Ecology*, 20, pp.66-74..



Tree based on Dice genetic distance among 29 field isolates of *Colletotrichum* sp. representing distinct haplotypes obtained after clone correction.



Does *Colletotrichum lentis* have sexual reproduction in the field?

- Field isolates with combinations of
 - IG-1/race 0
 - IG-2/race 0
 - IG-2/race 1
 - never IG1/race 1
- Ascospore-derived F_1 isolates have all combinations at equal frequencies



Menat, J., Armstrong-Cho, C. and Banniza, S., 2016. Lack of evidence for sexual reproduction in field populations of *Colletotrichum lentis*. *Fungal Ecology*, 20, pp.66-74..



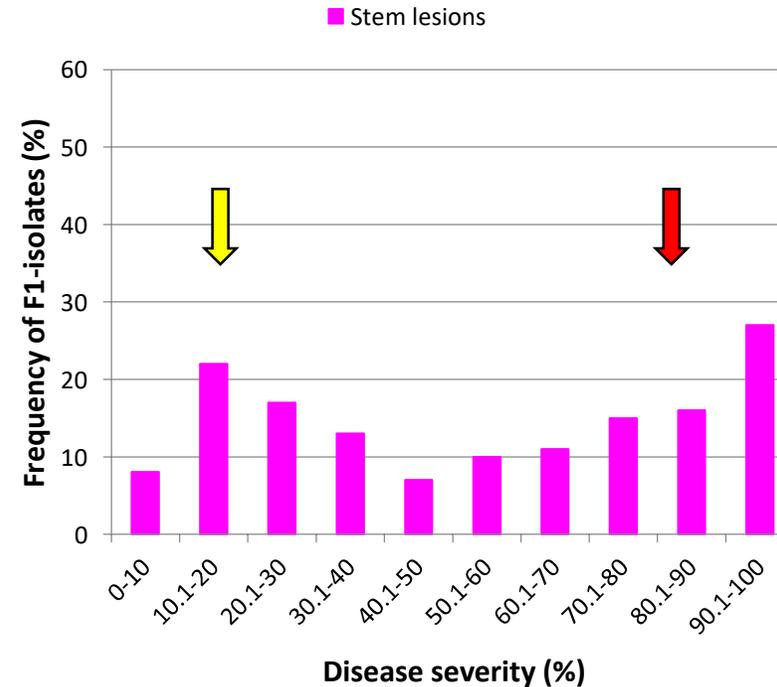
Breeding fungi: *Colletotrichum lentis* race 1 x race 0 population



- CT-21 (race 1/IG2) × CT-30 (race 0/IG1)
- Phenotyping of 142 asco-spore derived F₁-isolates and parental isolates
- Potential major gene / QTL effect

Bhadoria, V., MacLachlan, R., Pozniak, C., Cohen-Skalie, A., Li, L., Halliday, J. and Banniza, S., 2019. Genetic map-guided genome assembly reveals a virulence-governing minichromosome in the lentil anthracnose pathogen *Colletotrichum lentis*. *New Phytologist*, 221(1), pp.431-445

Virulence of F₁ isolates of
Colletotrichum lentis on
CDC Robin

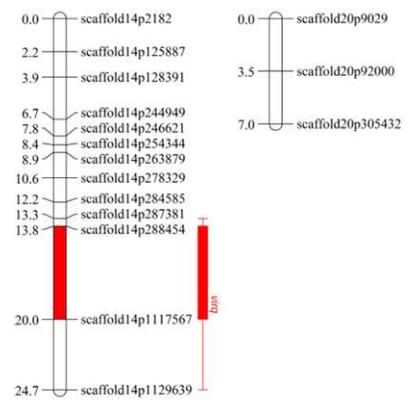
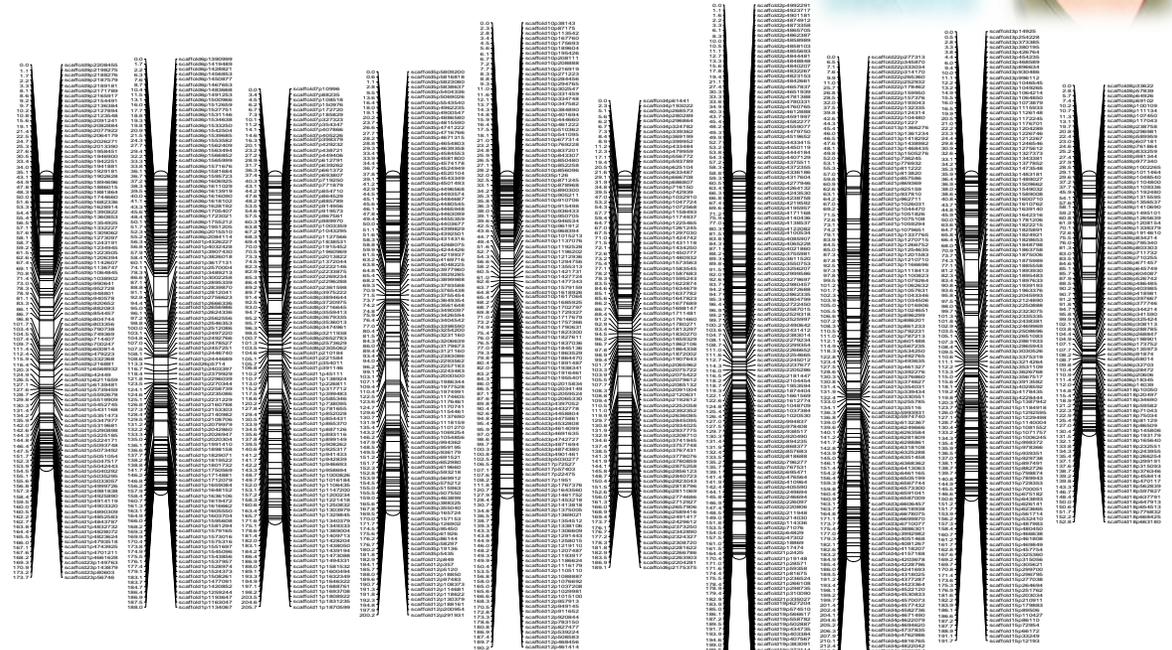


↓ race 1 parent ↓ race 0 parent

Sequencing and mapping of *Colletotrichum lentis*



- 56.10 Mbp genome assembly
- SNP-based map with 10 core chromosomes and 2 dispensable smaller chromosomes 98.27% of the genome
- QTL for virulence on LG/Chromosome 11



Bhadauria, V., MacLachlan, R., Pozniak, C., Cohen-Skalie, A., Li, L., Halliday, J. and Banniza, S., 2019. Genetic map-guided genome assembly reveals a virulence-governing minichromosome in the lentil anthracnose pathogen *Colletotrichum lentis*. *New Phytologist*, 221(1), pp.431-445



**NSERC
CRSNG**

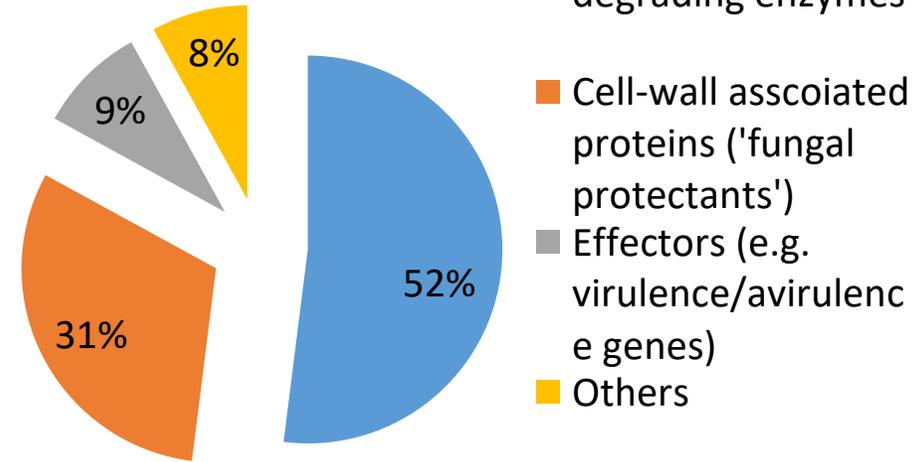




Virulence of *C. lentis* races

Bhadauria et al 2013. Eukaryotic Cell 12, 31
Bhadauria et al 2011. BMC Genomics 12, 327
Bhadauria et al 2015 BMC Genomics 16, 628

- Expressed sequence tag libraries
- *CtNudix* likely elicits switch from biotrophy to necrotrophy by inducing host cell death
- Knock-down of virulence effector *ClToxB* reduced virulence on lentil



Finding the receptor for ClToxB in lentil

- *In vitro* pull-down assay identified 43 putative interactors with ClToxB of interest
- Infiltration of purified ClToxB induces necrosis in lentil



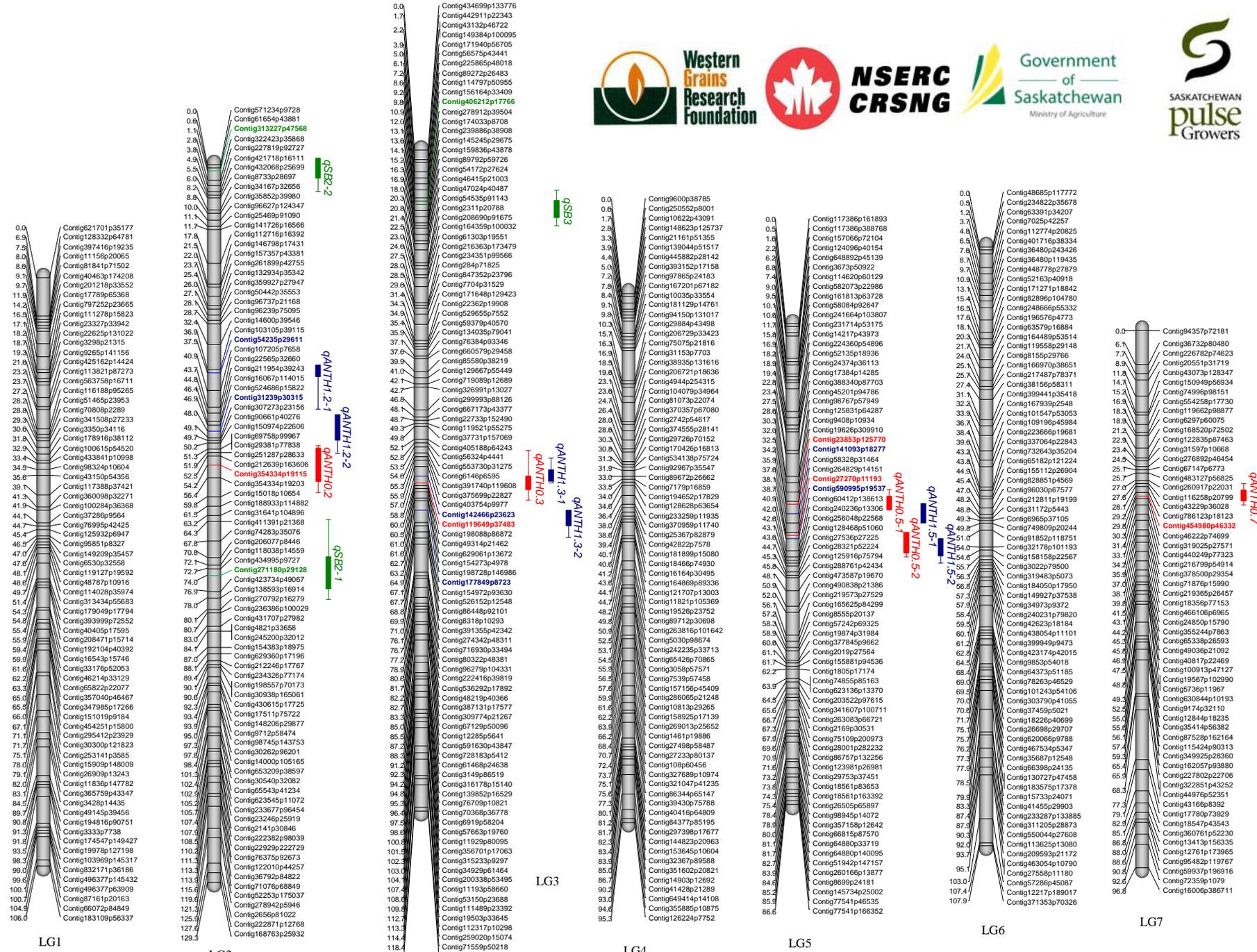


Resistance to anthracnose in *Lens ervoides* population LR-66

Green: stemphylium blight resistance QTL

Blue: anthracnose race 1 resistance QTL

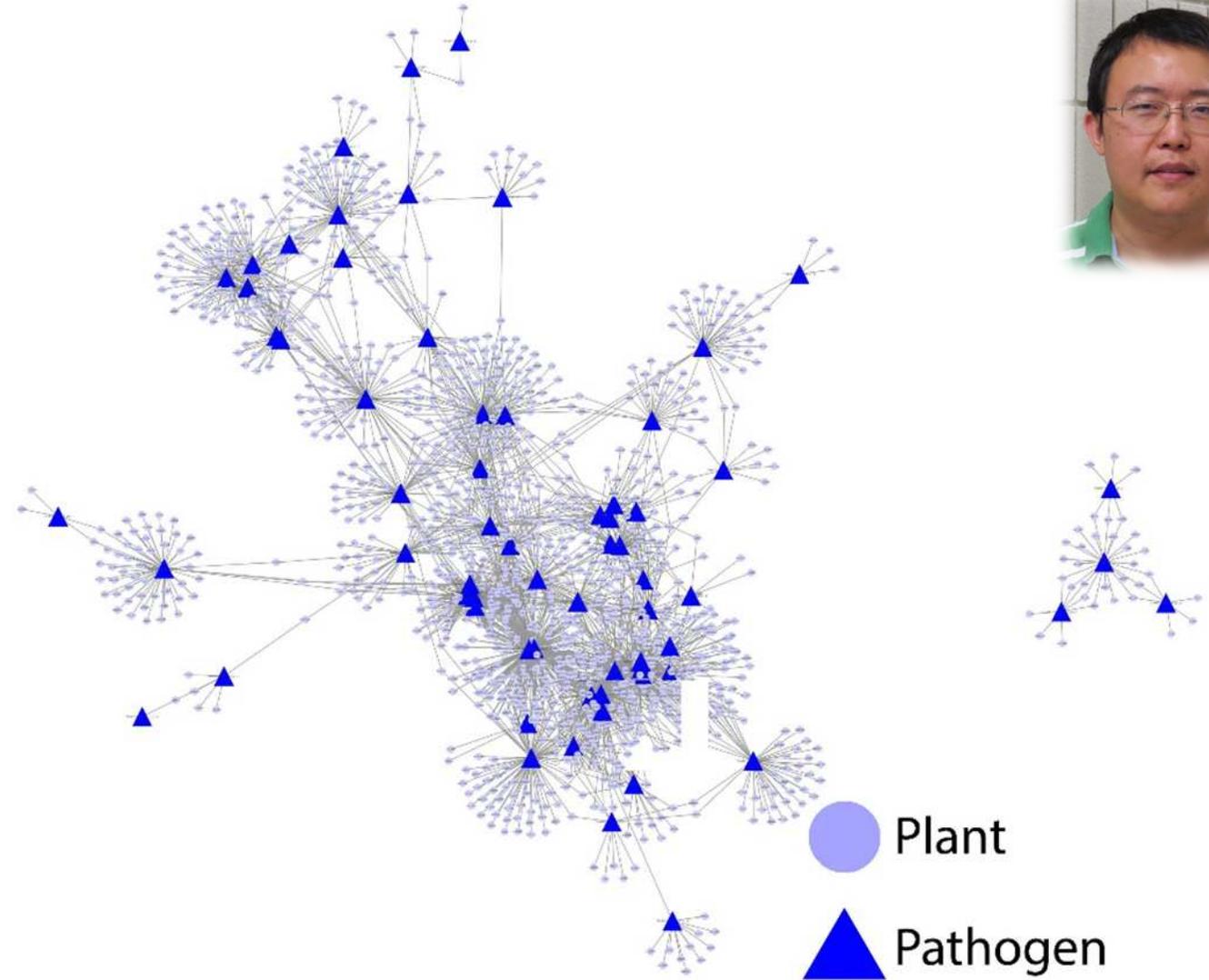
Red: anthracnose race 0 resistance QTL



Can we now just edit genes to make lentil resistant?

- NOT YET
- Quantitative host-pathogen systems are highly complex
- Gene editing requires
 - Knowledge of gene function
 - Identification of major switches for resistance or susceptibility

TO BE CONTINUED



Cao, Zhe, and Sabine Banniza. "Cross-Kingdom Gene Coexpression Analysis Using a *Stemphylium botryosum*–*Lens ervoides* System Revealed Plasticity of Intercommunication Between the Pathogen Secretome and the Host Immune Systems." *Molecular Plant-Microbe Interactions* 34, no. 12 (2021): 1365-1377.



Root rots in pulse crops

Aphanomyces root rot

CROPS: Pea and lentil
LOCATION: Saskatchewan

NAMES AND AGENCIES:

S. Banniza¹, V. Bhadauria¹, C.O. Peluola², C. Armstrong-Cho¹, and R.A.A. Morrall³

¹Crop Development Centre, University of Saskatchewan, 51 Campus Drive, Saskatoon, SK S7N 5A8

Telephone: 306-966-2619, **Facsimile:** 306-966-5015, **E-mail:** sabine.banniza@usask.ca

²Crop Protection Laboratory, 346 McDonald Street, Regina, SK S4N 6P6

³Department of Biology, University of Saskatchewan, 112 Science Place, Saskatoon, SK S7N 5E2

TITLE: FIRST REPORT OF APHANOMYCES EUTEICHES IN SASKATCHEWAN

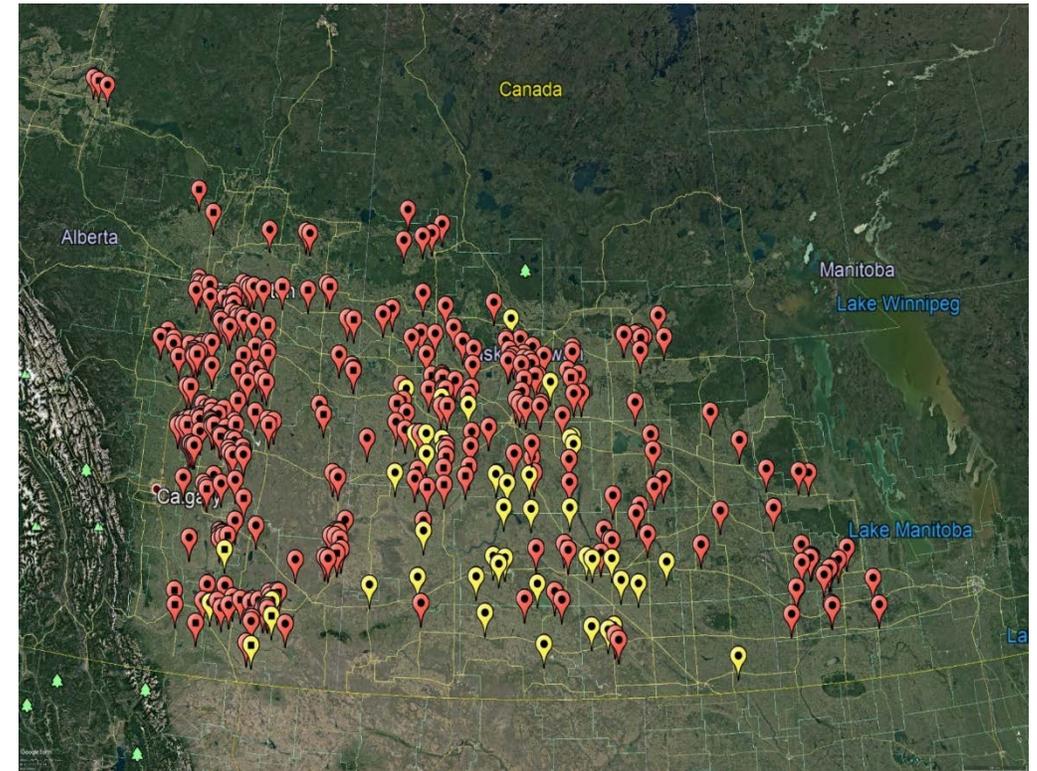
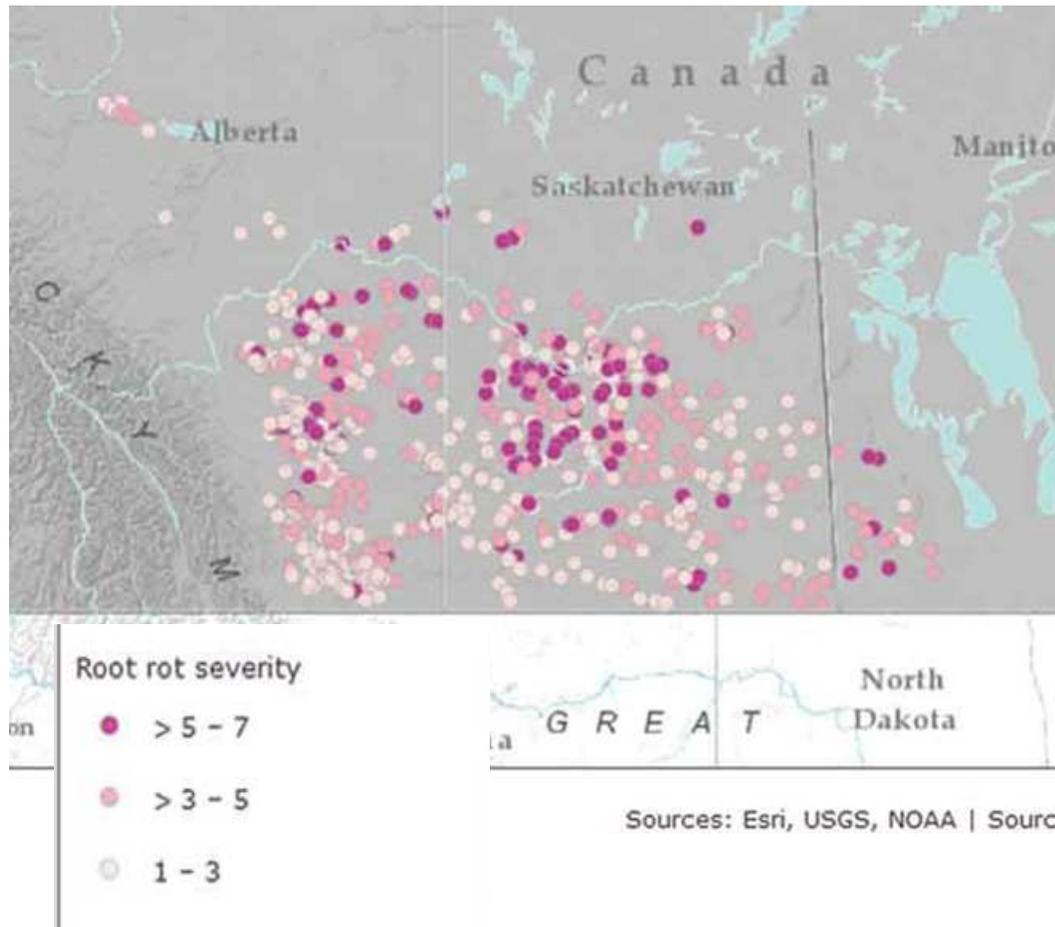
ABSTRACT: *Aphanomyces euteiches* was identified in samples from three diverse Saskatchewan locations using probe-based quantitative PCR. Wet conditions in several areas in 2012 and in preceding seasons have favored root rot and exacerbated symptom development. *Aphanomyces euteiches* is probably widespread and has been present for a long time, with recent conditions enabling its detection.

INTRODUCTION AND METHODS: Since 1948 there have been regular reports in the Canadian Plant Disease Survey (e.g. Vol.28 pp. 45; Vol. 34 pp. 69; Vol. 42 pp. 97; Vol. 56 pp. 23; Vol. 78. pp.12, 98; Vol.79, pp. 70) and elsewhere (Tu, 1985) of *Aphanomyces euteiches* or of "black root" diseases on pea and other legumes. These reports come from seven of Canada's ten provinces. Although it appears that *A. euteiches* is widespread in Canada and it is known to cause severe damage to pea crops in the U.S.A.

[/cgi-bin/rbaccess/rbunxcgi?F6=1&F7=IB&F21=IB&F22=IB&REQUEST=ClientSignin&LANGUAGE=ENGLISH&_ga=2.263379208.583740149.1538261856-1740749555.15382618](#)

Banniza, S., Bhadauria, V., Peluola, C.O., Armstrong-Cho, C. and Morrall, R.A.A., 2013. First report of *Aphanomyces euteiches* in Saskatchewan. *Can. Plant Dis. Surv.*, 93, pp.163-164.

Pea and lentil root rot surveys 2014-2017



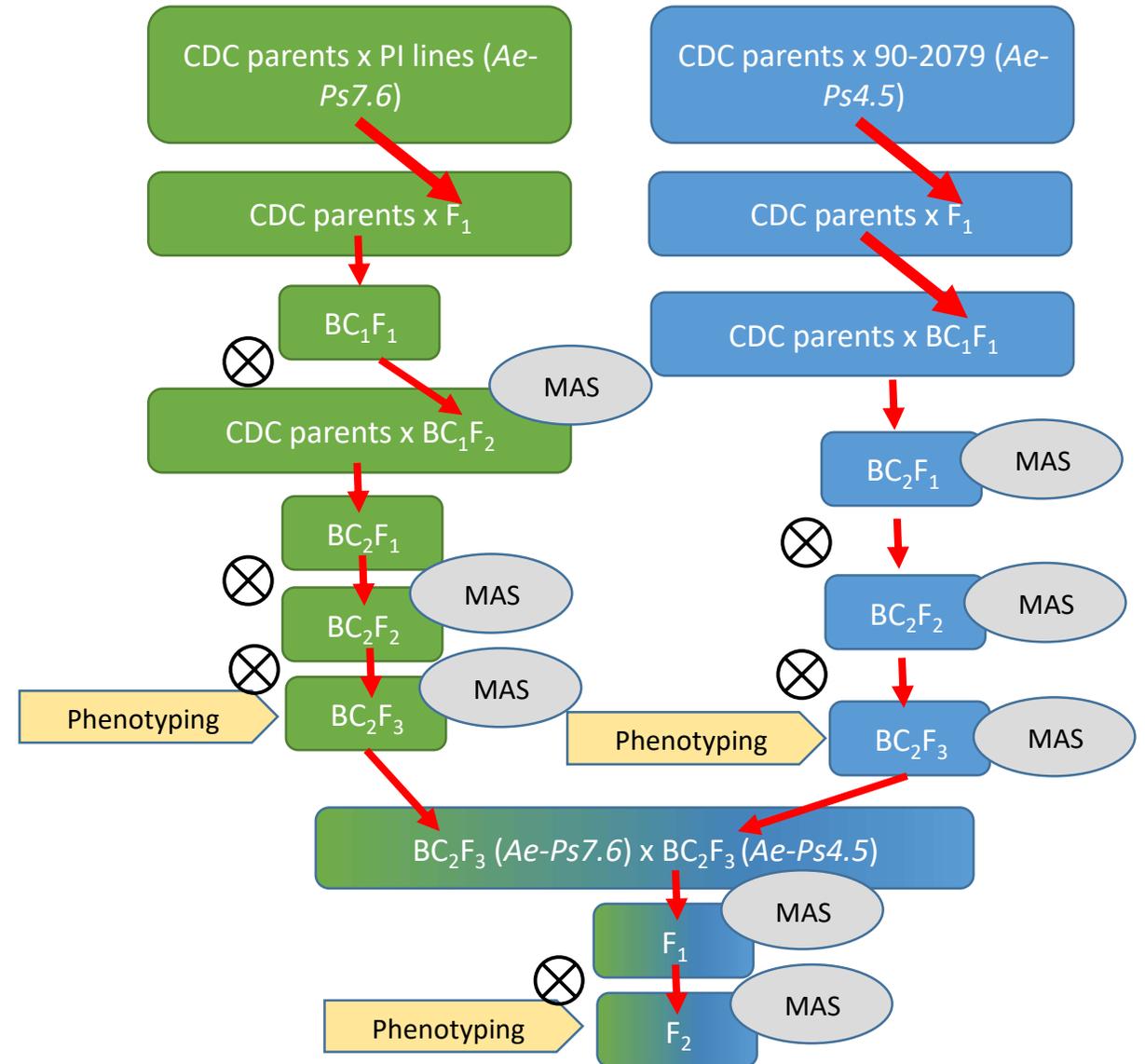
- Aphanomyces root rot in pea fields
- Aphanomyces root rot in lentil fields



Remember the
Western
Interior
Seaway?

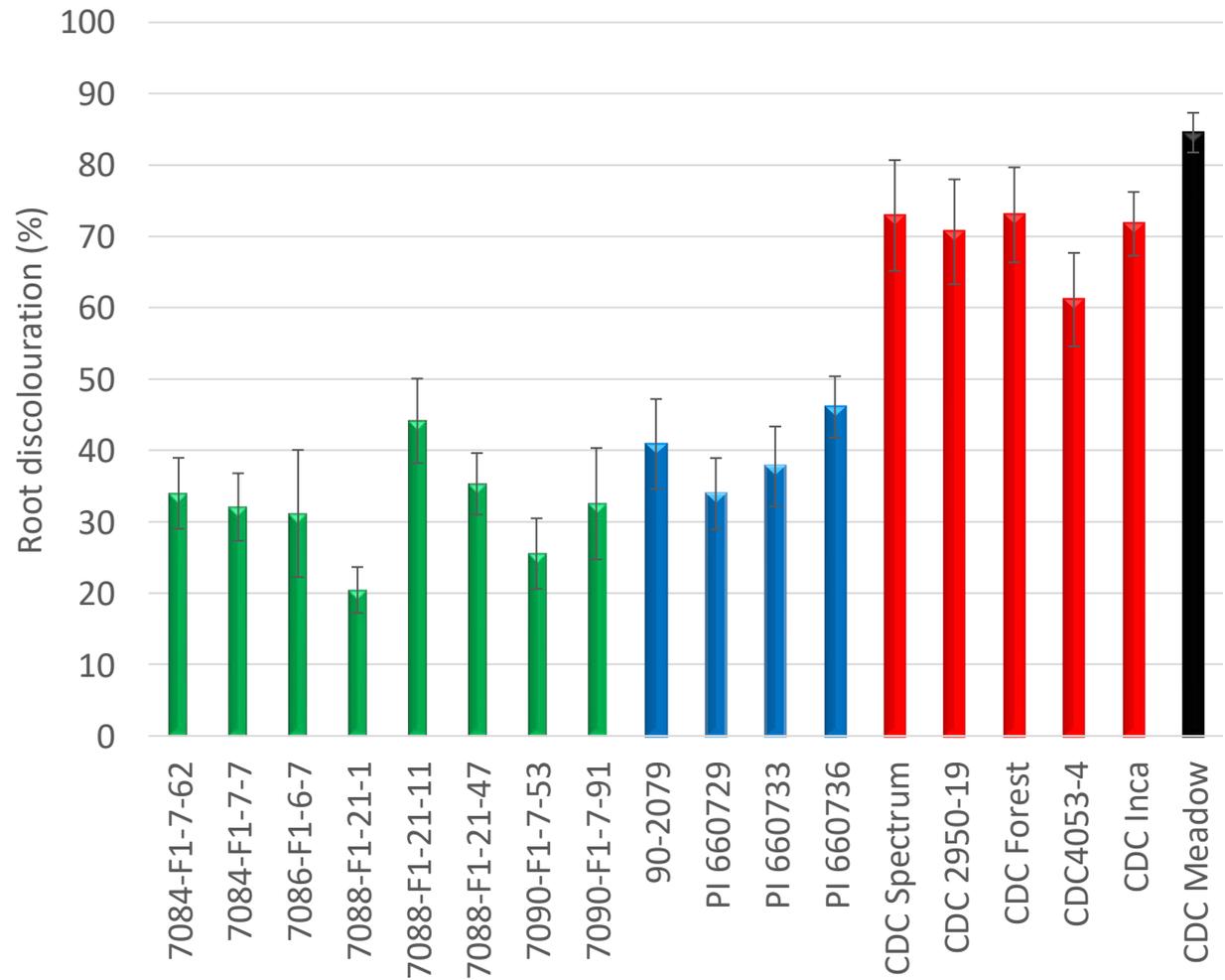
PEA: Backcrossing breeding for improved *Aphanomyces* resistance (Banniza & Warkentin)

- Multiple CDC pea varieties
- Crossed with 2 sources of resistance (PI lines & 90-2079) to transfer
 - 'Major' (*Ae-Ps4.5* & *Ae-Ps7.6*)
 - 'minor' QTLs

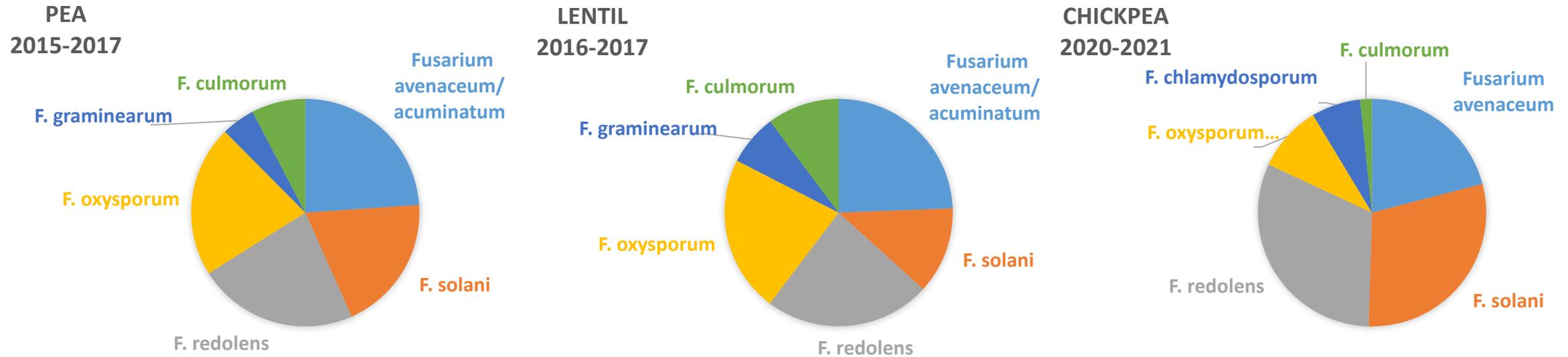


PEA: Some CDC lines carrying QTLs *Ae-Ps4.5* and *Ae-Ps7.6*

- Most promising lines arising were tested in
 - 2021 yield trials
 - 2021 field disease nursery
- 2022 pre-breeder seed development
- 2023 /2024 pea Co-op Test
- 2025 Registration?



Fusarium species in pea, lentil and chickpea

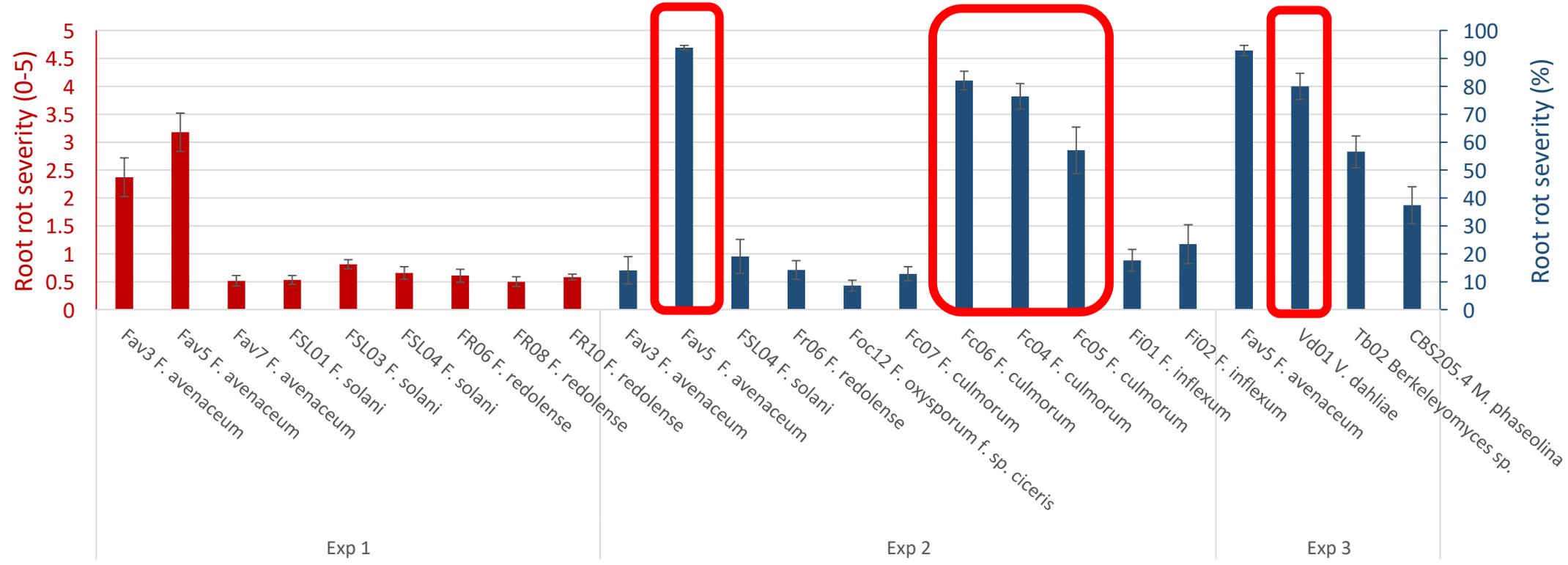
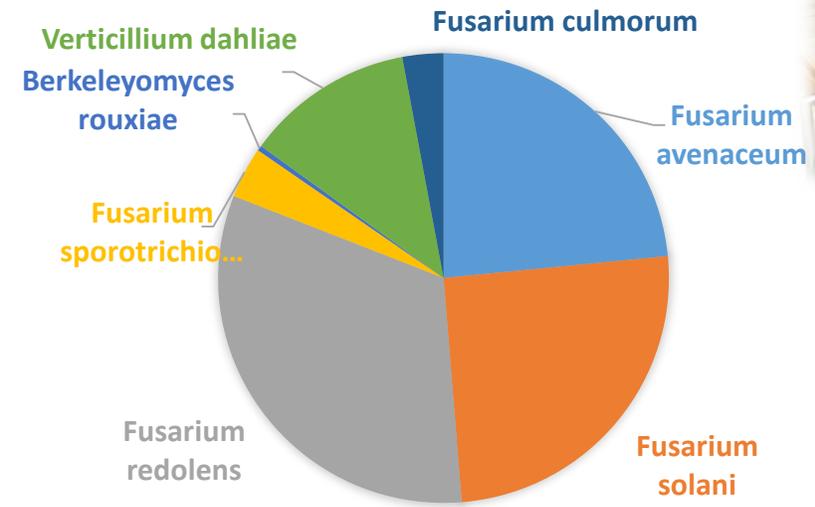


Chatterton et al. 2017. Can J Plant Pathol 41:98-114. <https://doi.org/10.1080/07060661.2018.1547792>

Armstrong-Cho et al., in press

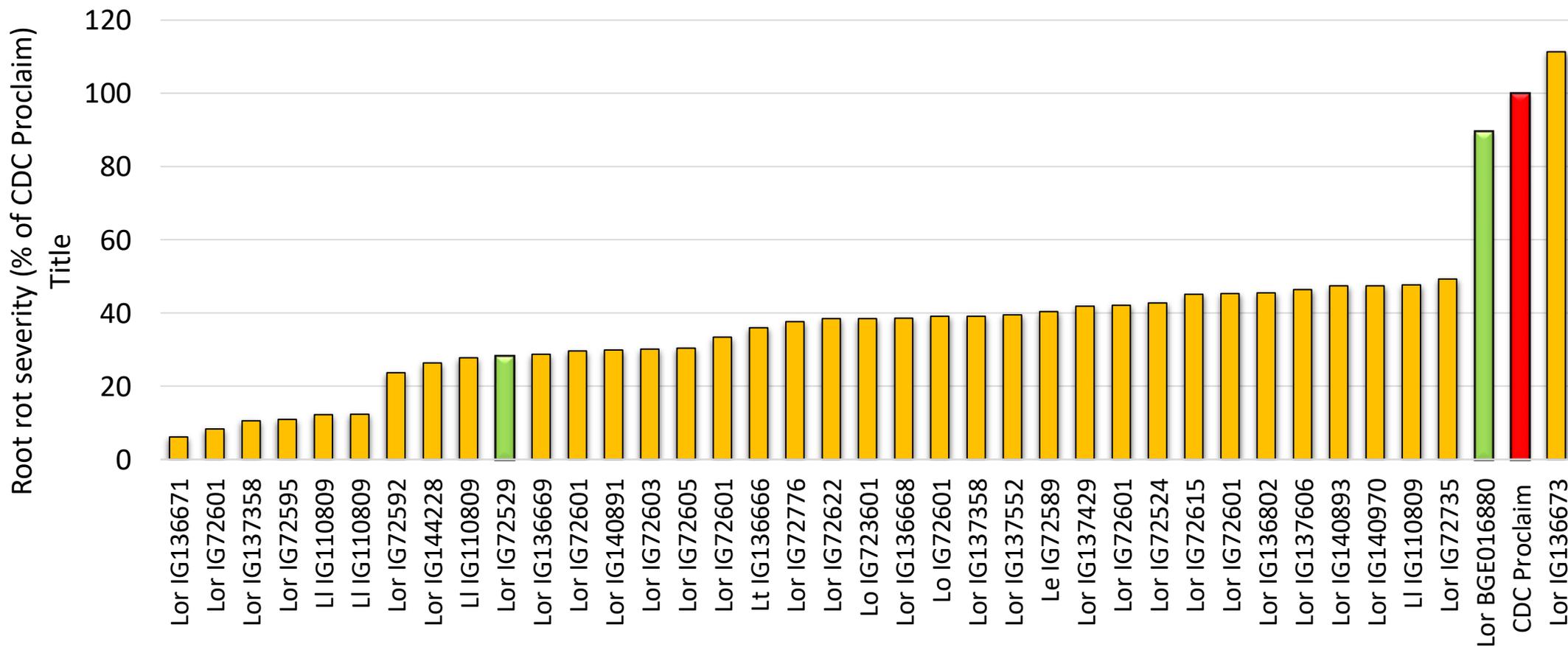


New pathogen in chickpea in 2021





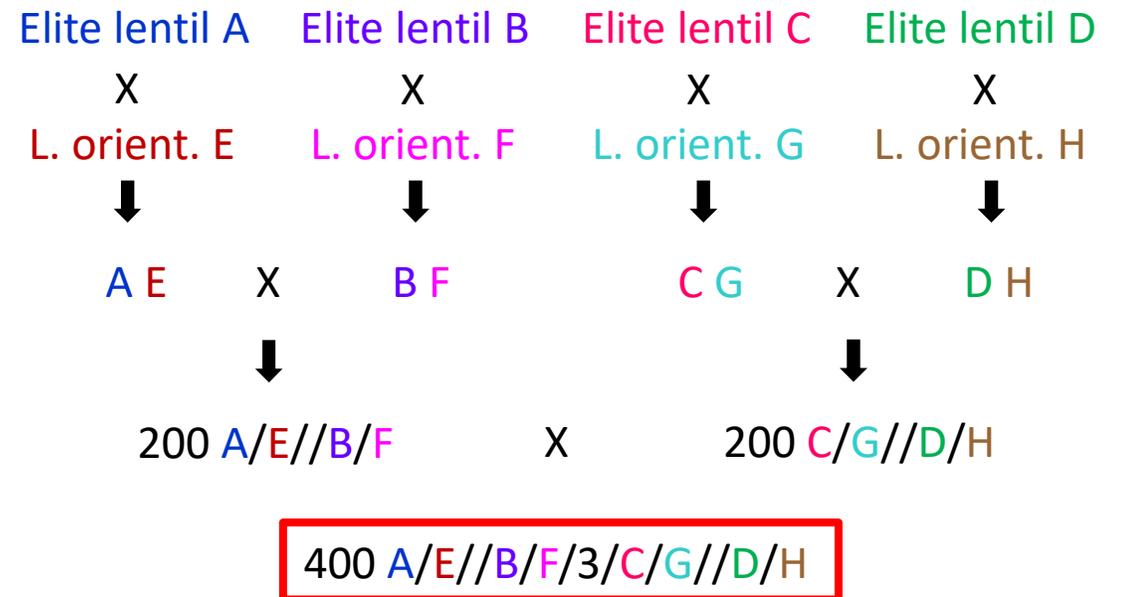
Aphanomyces root rot resistance in lentil



Lor: *Lens orientalis*, Ll: *L. lamottei*, Lt: *L. tomentosus*, Lo: *L. odemensis*, Le *L. ervoides*

New: Combining *Aphanomyces* and *F. avenaceum* resistance in lentil (Banniza & Bett)

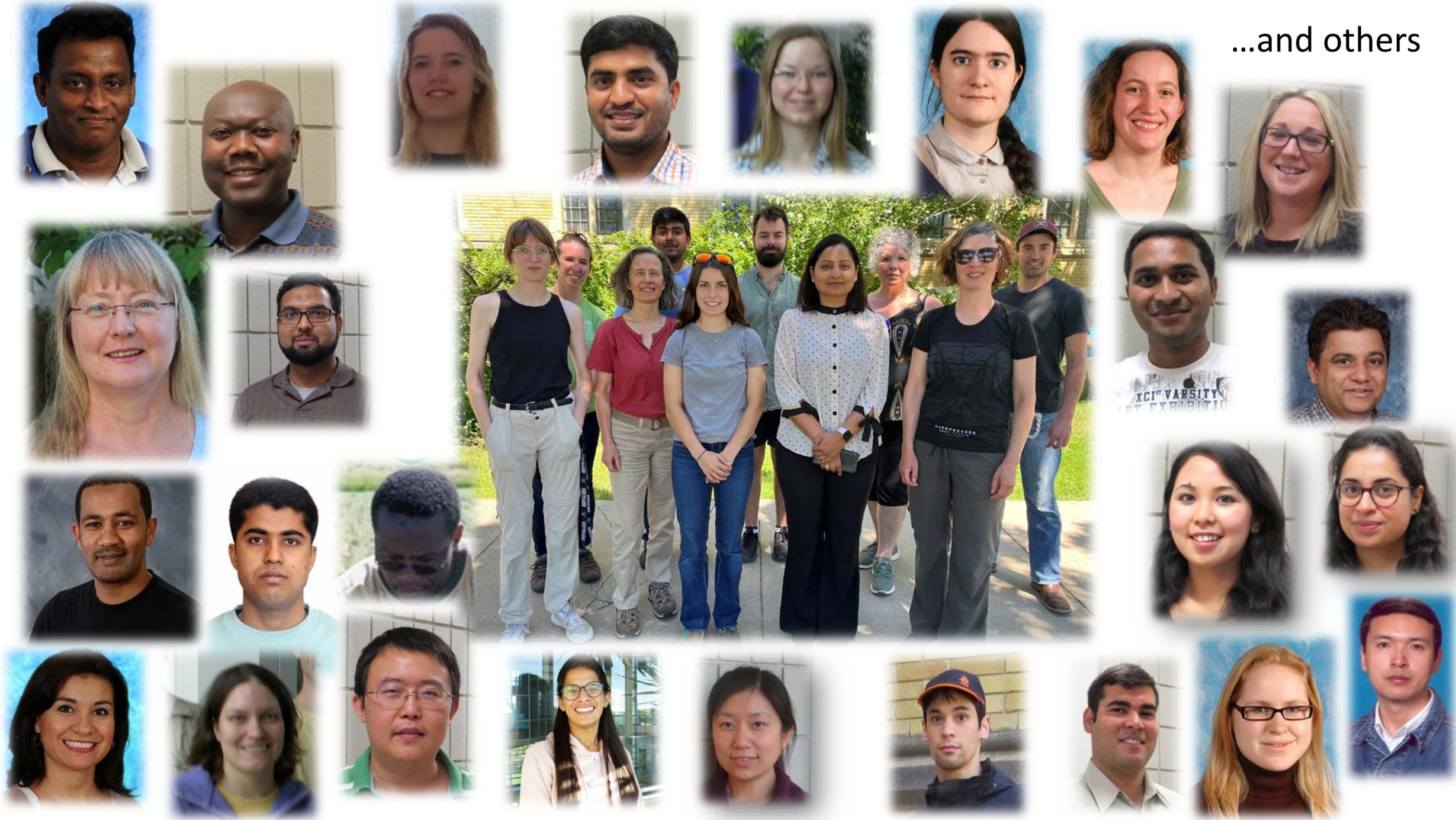
- Development of a multi-parent advanced generation inter-cross (MAGIC) population
- Identify lines with improved *Aphanomyces* and *F. avenaceum* resistance
- Determine effect of seed coat colour
- Develop markers for breeding



Going forward

- Root rot resistance in pea, lentil and chickpea
- *Colletotrichum lentis* race 0 (anthracnose) resistance in lentil
- Explore foliar fungal pathogens of faba bean
- Resistance screening to bacterial blights and anthracnose in bean
- Support provincial disease surveys through training

...and others



Incomplete list of formal and informal research collaborators

Bob Conner (AAFC Morden)
Randy Kutcher (AAFC Melfort)
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Yangdou Wei (Dept. of Biology, UofS)
Bruce Gossen (AAFC Saskatoon)
Sue Boyetchko (AAFC Saskatoon)
Clarice Coyne (USDA, USA)
Rebecca McGee (USDA, USA)
Marie-Laure Pilet-Nayel (INRA, FRA)
Anne Moussart (INRA, FRA)
Bernard Tivoli (INRA, FRA)
Christophe Le May (Agrocampus Ouest, FRA)
Mary Burrows (Montana State University, USA)
Michael Harding (Government of Alberta)
Robyne Davidson (Lakeland College)
Syama Chatterton (AAFC Lethbridge)
Deborah McLaren (AAFC Brandon)
Hossein Borhan (AAFC Saskatoon)
Judith Lichtenzveig (University of Western Australia, AUS)

Rebecca Ford (University of Melbourne, AUS)
Jenny Davidson (SARDI, AUS)
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Julie Pasche (North Dakota State University, USA)
Axel Diederichsen (AAFC Saskatoon)
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Tobin Peever (Washington State University, USA)
Weidong Chen (Washington State University, USA)
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Tom Wolf (AAFC Saskatoon)
Terry Hogg (ICDC)
Keith Seifert (AAFC Ottawa)
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Art Davis (Dpt. of Biology, UofS)
Brienne McInnes (NARF Melfort)
Jessica Pratchler (NARF Melfort)
Diego Rubiales (CSIC, Cordoba, ES)
Roman Labuda (University of Veterinary Medicine Vienna, AT)
Many colleagues in the Dpt. Of PLSC, UoS



Agriculture and
Agri-Food Canada

Agriculture et
Agroalimentaire Canada



Acknowledgments



We acknowledge we are on Treaty 6 Territory and the Homeland of the Métis.

We pay our respect to the First Nations and Métis ancestors of this place and reaffirm our relationship with one another.