

# The origins of aster leafhoppers to Western Canadian prairies

My PhD project:

An early warning system to predict aster yellows outbreaks in  
Western Canada: origin and arrival of migrant leafhoppers.

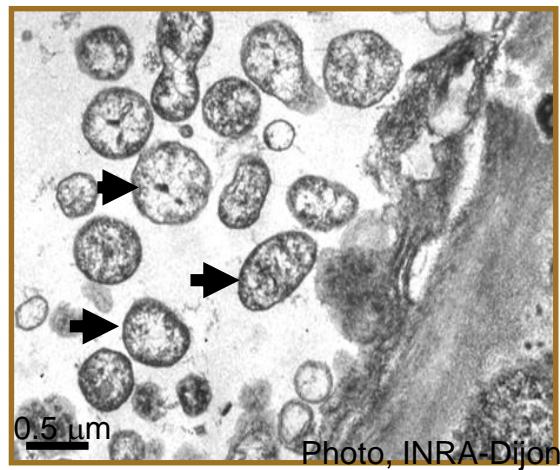


Karolina Pusz-Bochenska, Tyler Wist, Tim Dumonceaux, Erl Svendsen, Sean Prager, Chrystel Olivier, Keith Hobson  
Agriculture and Agri-Food Canada  
Saskatoon Research and Development Centre

# Aster yellows disease

- caused by the nonculturable, Gram-positive bacterium

*'Candidatus Phytoplasma'* spp.



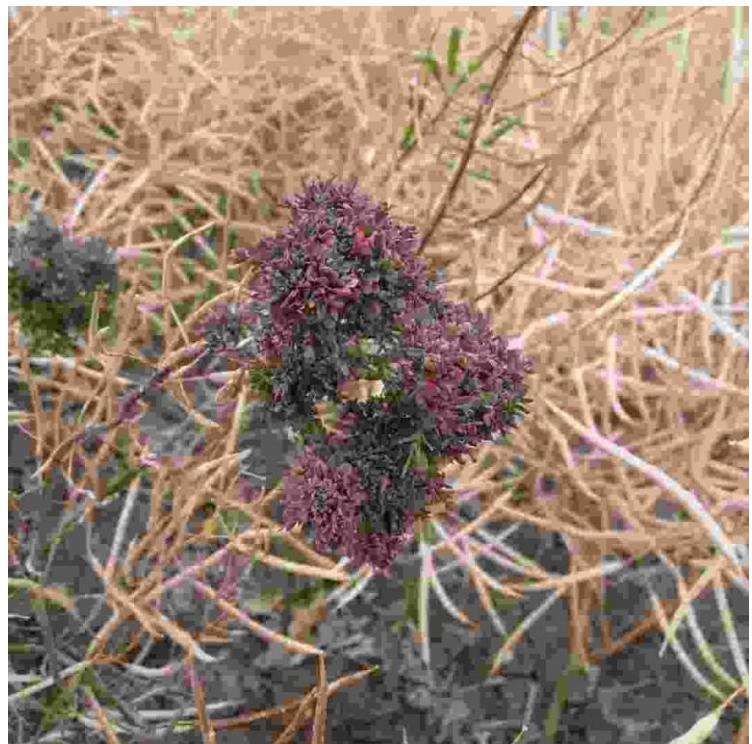
- The aster leafhopper *Macrosteles quadrilineatus* is the main vector of the AY phytoplasma.
- AY infects over 300 species of plants, including:  
**canola, barley, and the pulses.**
- **Symptoms:** Phyllody can lead to **very small** malformed seeds or **no seeds** at all.



(Tyler Wist)



In 2012 in SK loss was estimated at around **\$270 million**.

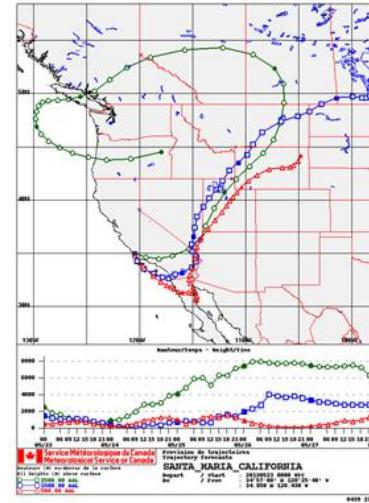


Tyler Wist

# Project objectives:

- Evaluate the origin of migrant aster leafhoppers to better understand AY risk each year.

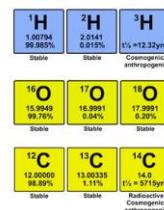
1. Wind trajectories monitoring.



2. Genetic profiling of leafhoppers sampled in SK and from Southern populations.

>KPB2\_LCO\_MP8014\_19 KPB2-SK3k6-Aberdeen\_CO1.0.2  
atgtgatcgttacatcccacgcatttataatttctttatagttataccattataa  
ttgggggtttgttaattgactacttcacttataattggagctcctgatatagcattc  
cacgactaaataatataagattctgacttcccccccttcattaacactattaatattaa  
gatcaatagtagaaaatggggtagggacgggtgaacagtataccccccctatctagaa  
atatgcacatgcggcccaagagtttatatatcaattttcttacatttagccgta

3. Stable isotopes.





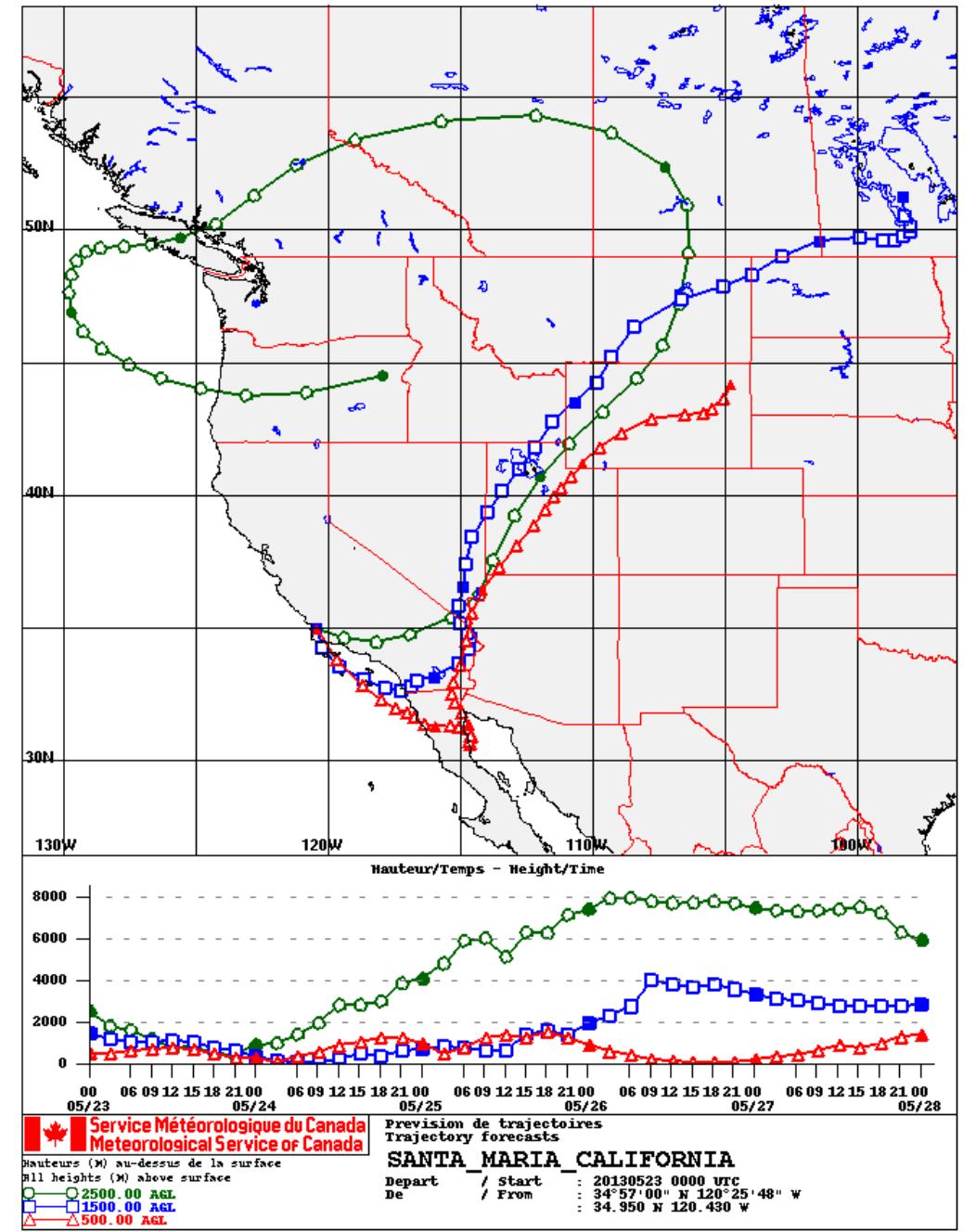
# Aster Yellows predictions

- South Wind: Reverse Trajectories
- Prairie Pest Monitoring Network (PPMN) monitors winds
- Environment Canada

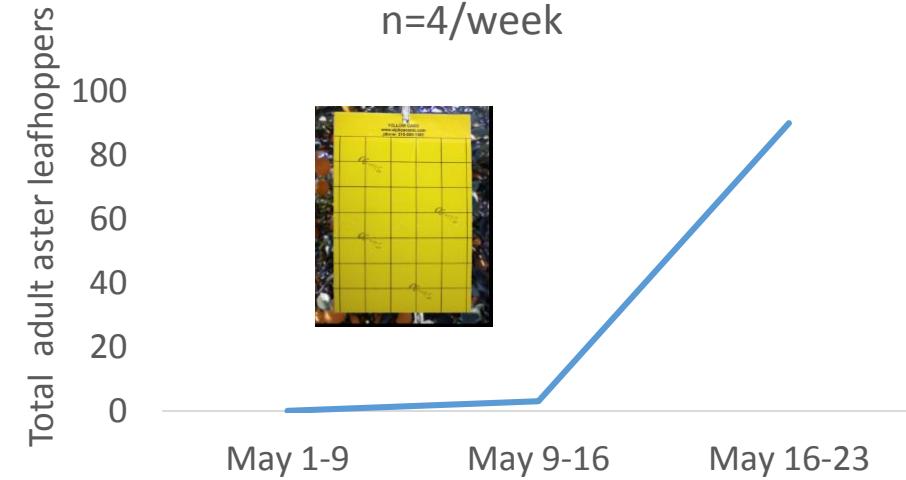
Owen Olfert<sup>1</sup>, Ross Weiss<sup>1</sup>, Meghan Vankosky<sup>1</sup> and Serge Trudel<sup>2</sup>

1 - AAFC

2 - ECCC



## MooseJaw SK 2018 Yellow Sticky Cards n=4/week



- 2018, May 9<sup>th</sup> - 16<sup>th</sup> **1<sup>st</sup>** arrival of aster leafhoppers (Southern Sask)
- 2018, May 23<sup>rd</sup> **2<sup>nd</sup>** arrival. Southern and Northern coverage

**RT wind from Washington**



RESEARCH ARTICLE

## Molecular data reveals California as the potential source of an invasive leafhopper species, *Macrosteles* sp. nr. *severini*, transmitting the aster yellows phytoplasma in Hawaii

J.J. Le Roux<sup>1,2</sup> & D. Rubinoff<sup>1</sup>

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**Keywords**

CO1; invasion history; *Macrosteles*; NADH1; phylogenetics; Wg.

**Correspondence**

J.J. Le Roux, DST-NRF Centre of Excellence for Invasion Biology, Department of Botany and

**Abstract**

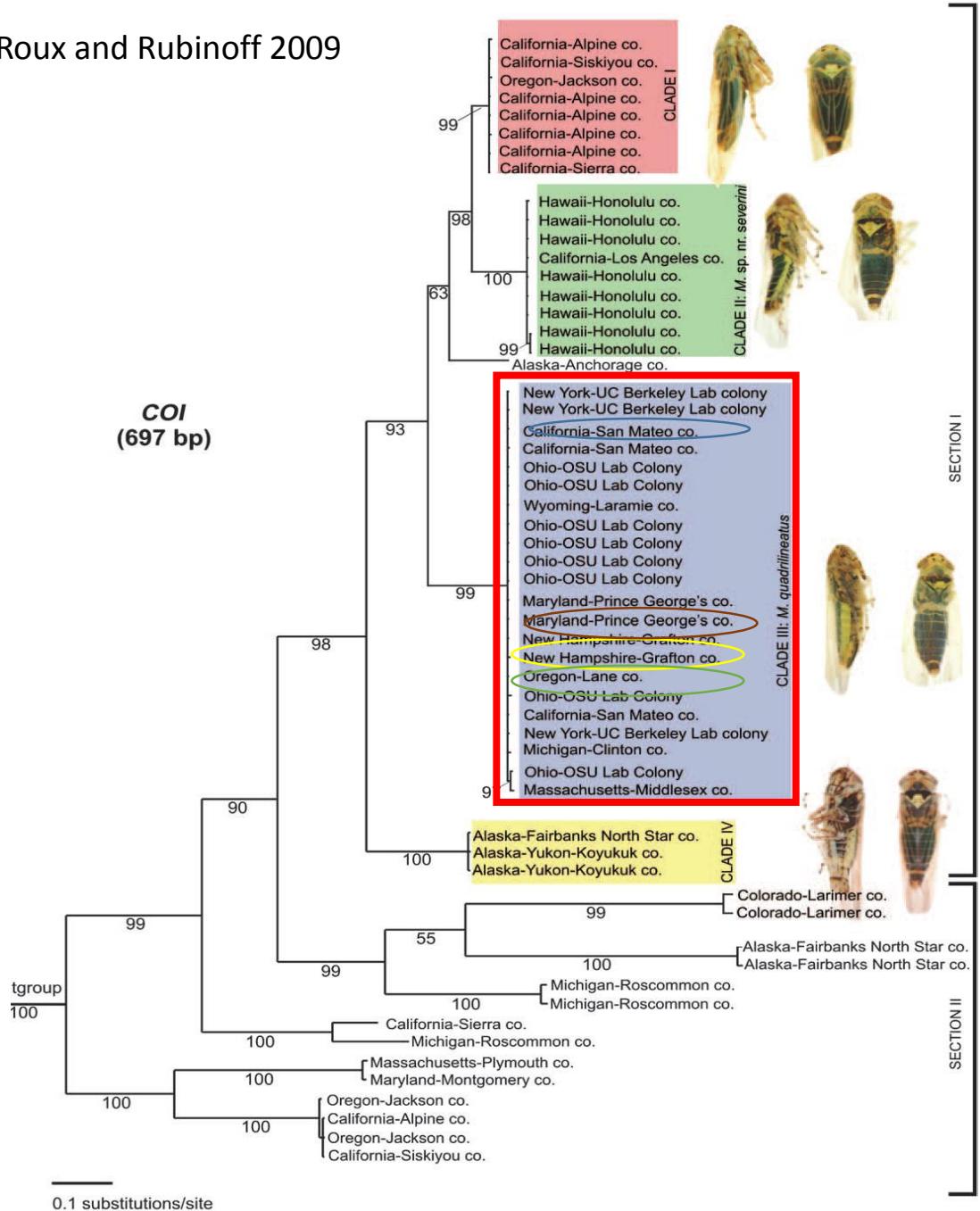
A species of aster leafhopper (*Macrosteles* sp.) became established in 2001 on Oahu, Hawaii, and through the transmission of the aster yellows phytoplasma, caused devastating losses to the island's watercress industry. DNA sequence data were analysed from two mitochondrial genes [cytochrome oxidase sub-

Genetic markers to ID  
popns of aster  
leafhoppers: CO1 and  
NADH1

Le Roux and Rubinoff  
2009 created  
GenBank records



Le Roux and Rubinoff 2009



## Sask leafhoppers from 2018

- SK7k68
    - SK8Mell
  - EU081848.1\_Mq\_Wyoming\_Laramie
  - SK7k102
  - SK7k56
  - SK7k74
  - SK7k83
  - SK7k13
  - SK7k41
    - SK7k104
  - SK7k60
  - EU081831.1\_Mq\_California\_San\_Mateo
  - SK7k66
  - SK7k54
  - SK7k88
  - SK7k24
  - SK7k88
  - SK7k15
  - SK7k36
  - SK7k65
    - SK7k48
  - EU081869.1\_Mq\_Oregon\_Lane
  - SK7k67
  - SK7k106
  - SK7k70
  - SK7k23
  - SK7k95
  - SK7k42
    - EU081859.1\_Mq\_New\_Hampshire\_Grafton
  - EU081884.1\_Mq\_California\_San\_Mateo
  - EU081854.1\_Mq\_Maryland\_Prince\_George's
  - SK3k6
  - SK7k53
  - SK4k1
  - SK7k21
  - SK7k92
  - SK7k18
  - SK7k35
    - SK7k85
  - SK7k8
  - EU081832.1\_Mq\_California\_San\_Mateo
  - SK7k1
  - SK7k51
  - SK7k71
  - SK7k25
  - SK7k10
  - SK7k19
  - SK7k62
    - SK37k
  - EU081858.1\_Mq\_New\_Hampshire\_Grafton
  - SK7k101
  - SK7k55



SK7k20	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK41k2	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k23	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k25	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k24	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k21	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK9k2	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k77	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k74	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k73	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG

- little sequence variation in regions tested
- Next: look for SNPs across the genome (Illumina)

SK7k51	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k54	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k53	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k50	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k100	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k102	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k101	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k67	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k1	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k66	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK3k6	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k37	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
SK7k39	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
EU981848.1_Mq_Wyoming_Laramie	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
EU981858.1_Mq_New_Hampshire_Grafton	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
EU981869.1_Mq_Oregon_Lane	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
EU981832.1_Mq_California_San_Mateo	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
EU981831.1_Mq_California_San_Mateo	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG
EU981854.1_Mq_Maryland_Prince_George's	TCATTAATTAA	ATAATGATCA	AGCATAACAAT	GTGATCGTTA	CATCCCACGC	ATTTATTATA	ATTTTCTTTA	TAGTTATACC	AATTATAATT	GG

# **Project objectives:**

- Create a fast, diagnostic tool to determine if leafhoppers are infected with AY.

# Aster yellows detection:

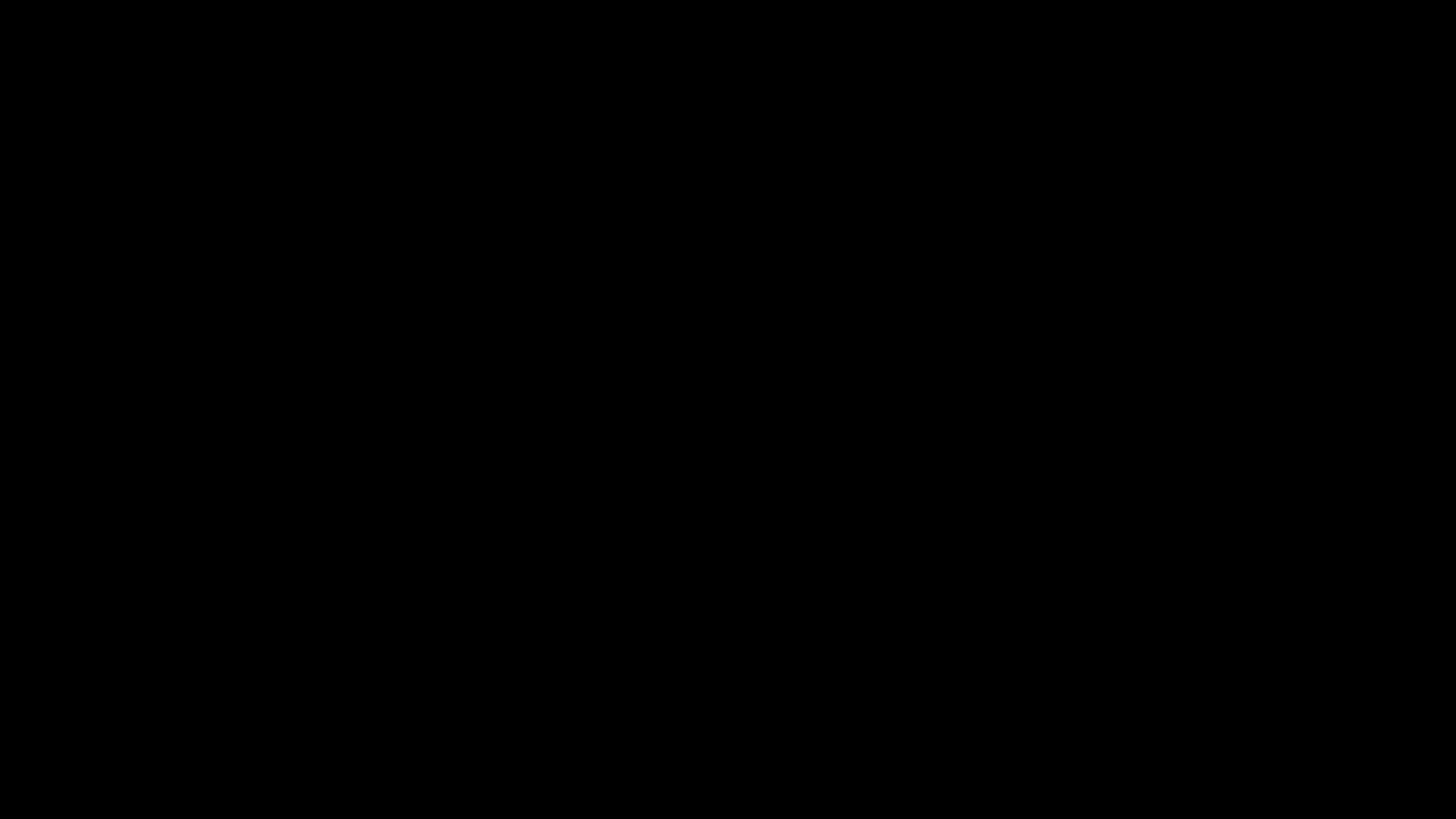
- Molecular diagnostics

1. DNA extraction



2. Molecular detection

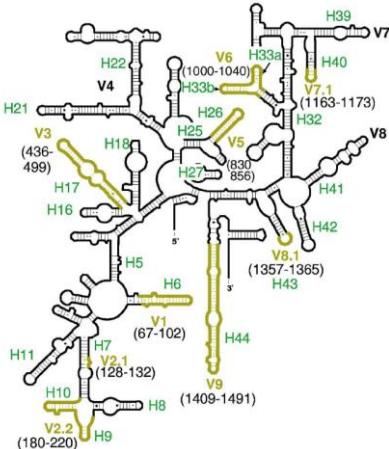
# Rapid AY diagnostics (RAYD)



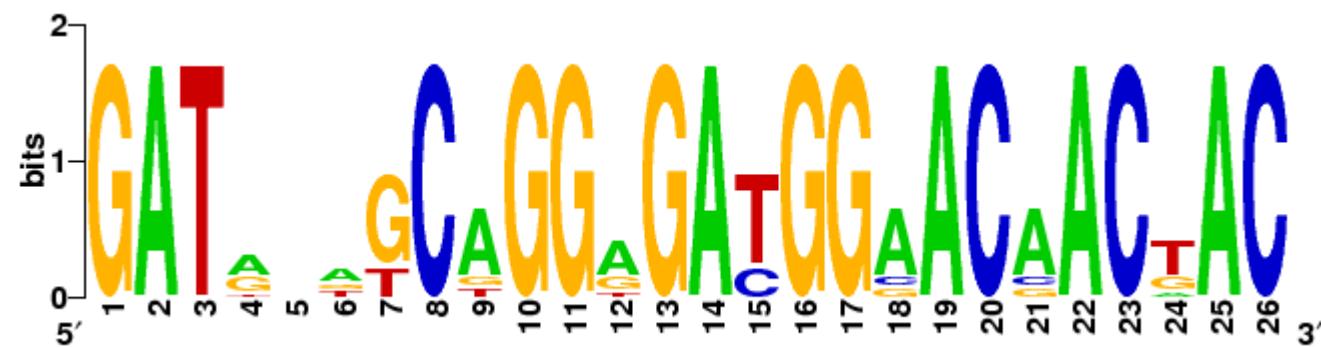
Starring Rebecca Green working out of Tim Dumonceaux's lab

# Molecular diagnostic assays

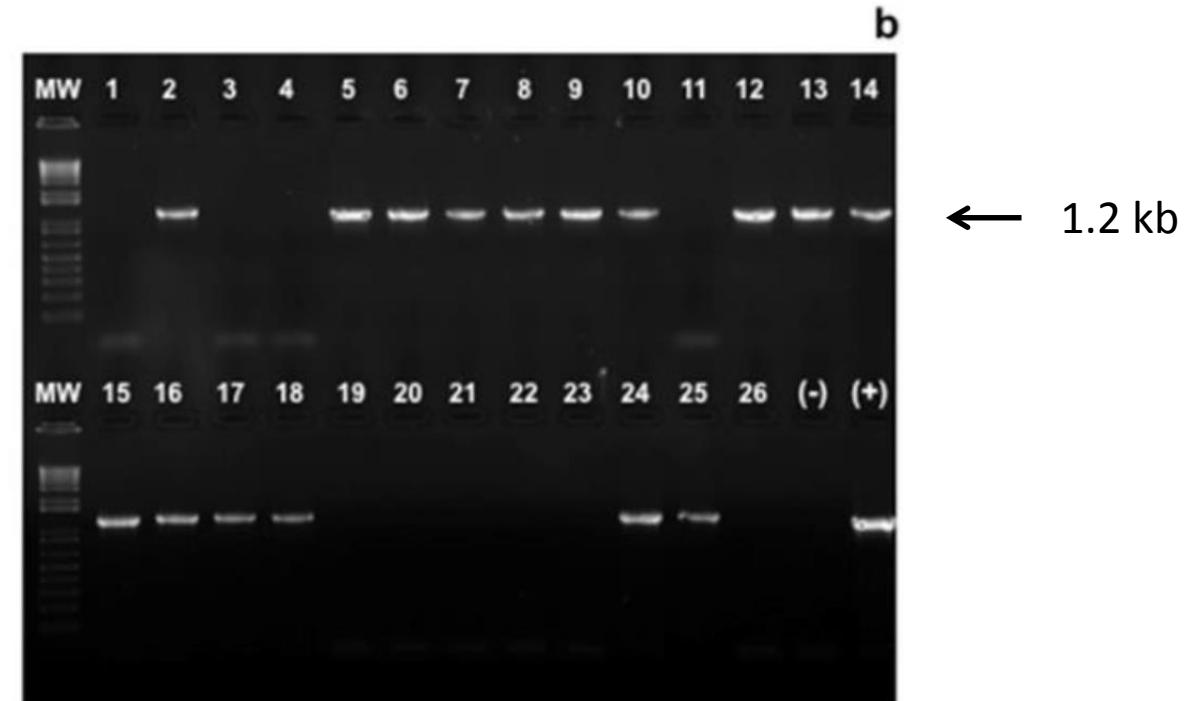
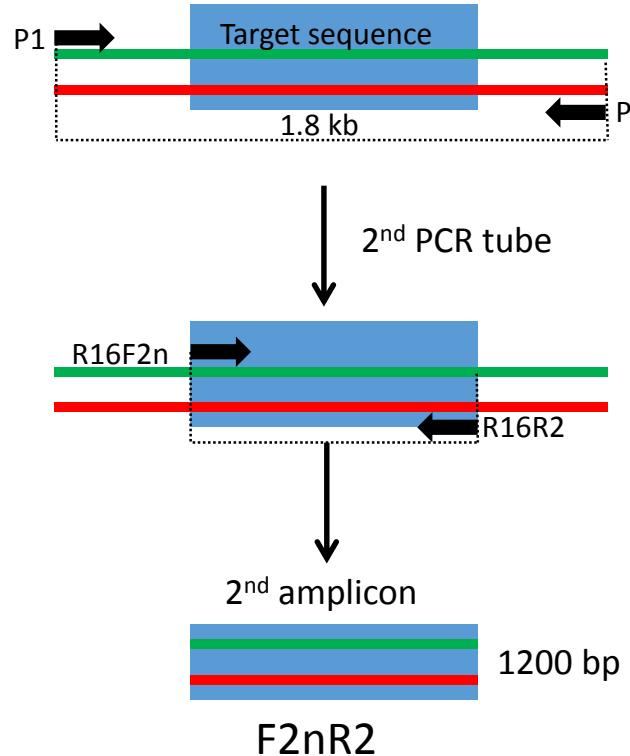
## Detection and quantification of phytoplasma



- “Nested” PCR
- PCR (conventional)
- qPCR
- ddPCR
- LAMP



# “nested” PCR based on 16S rRNA-encoding locus (F2nR2)



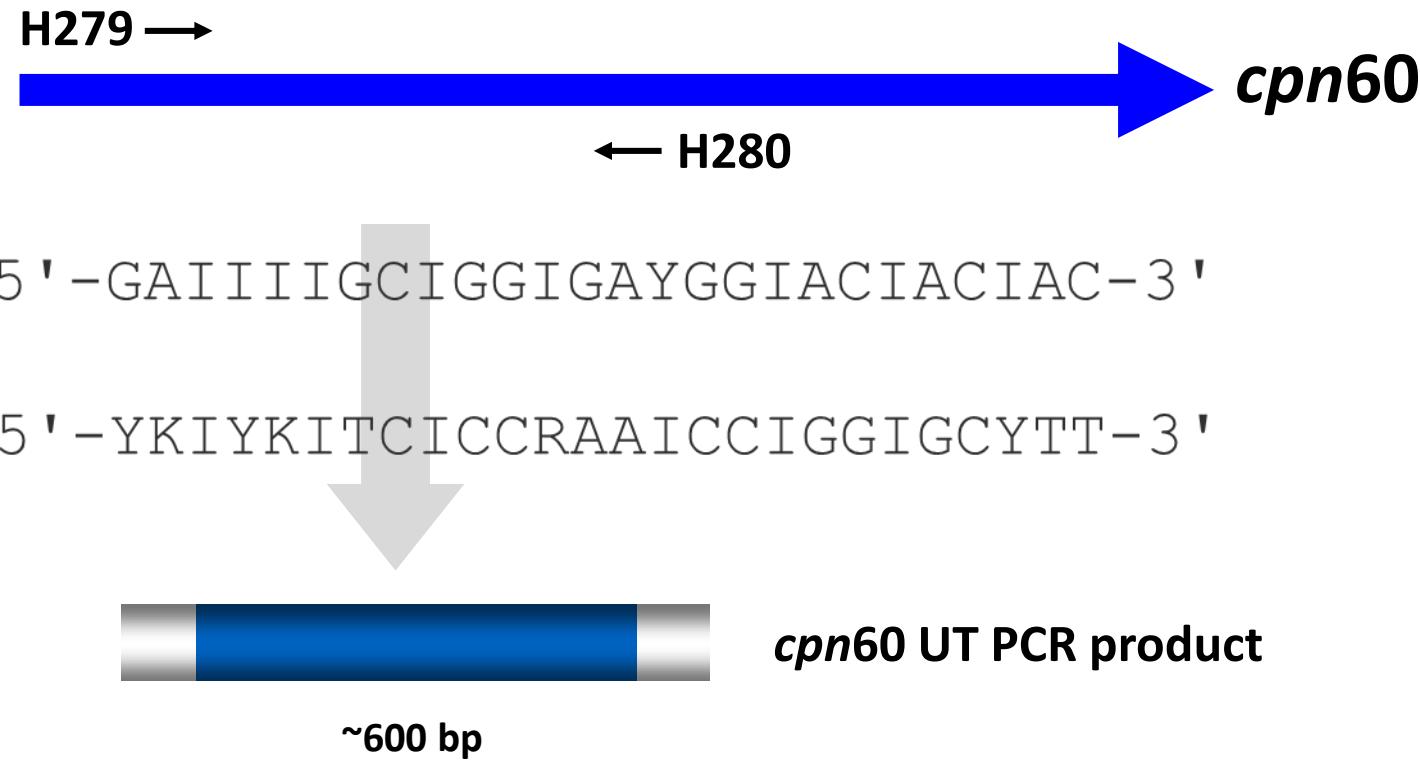
~5hr

~45 min

Gundersen, D. E., & Lee, I. M. (1996). Ultrasensitive detection of phytoplasmas by nested-PCR assays using two universal primer pairs. *Phytopathologia Mediterranea*, 35(3), 144-151. doi:10.2307/42685262

- second “nested” PCR based on different encoding gene- rp ribosomal protein

# “Universal” PCR amplification tools targeting *cpn60*

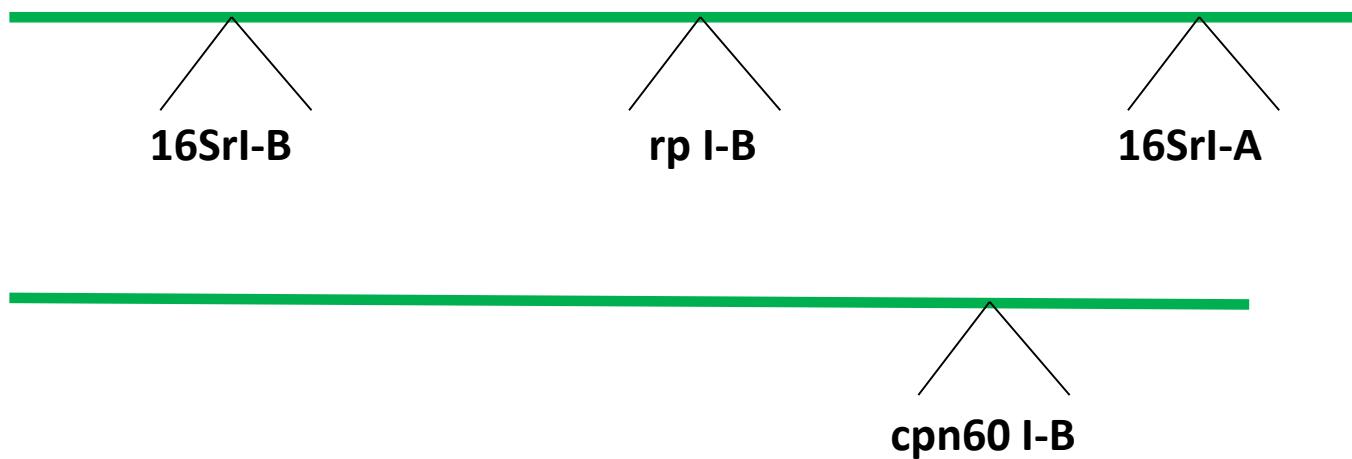


549-567 bp of template-derived sequence

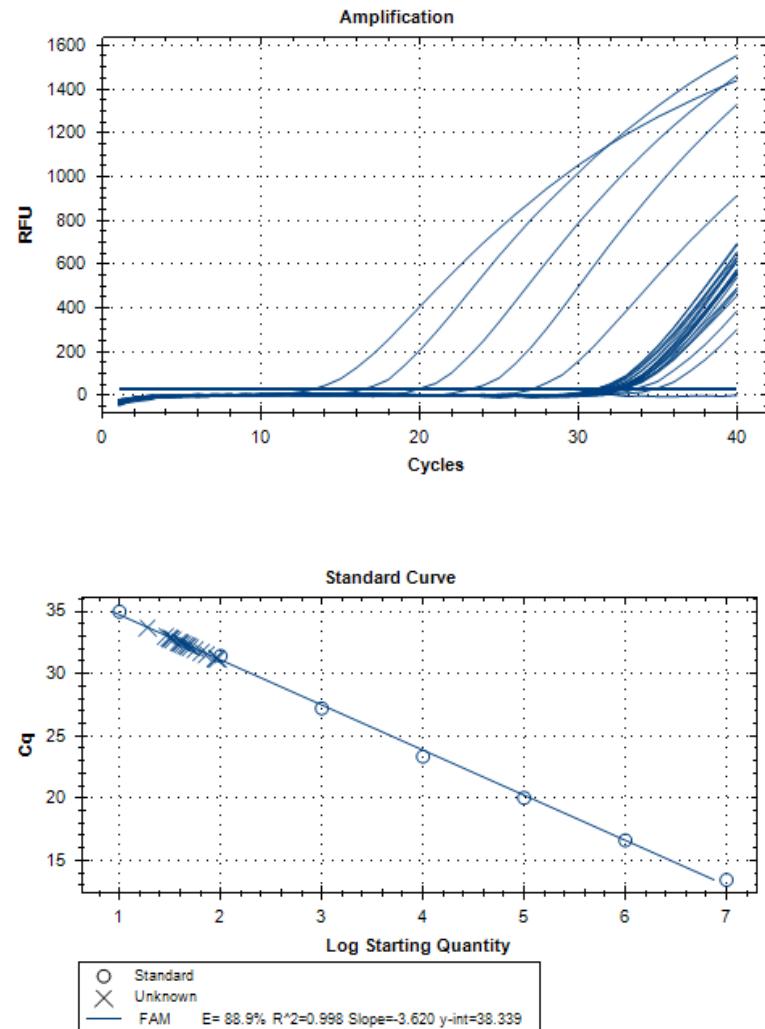
# Phytoplasma genome

AY-canola (Saskatoon, SK)

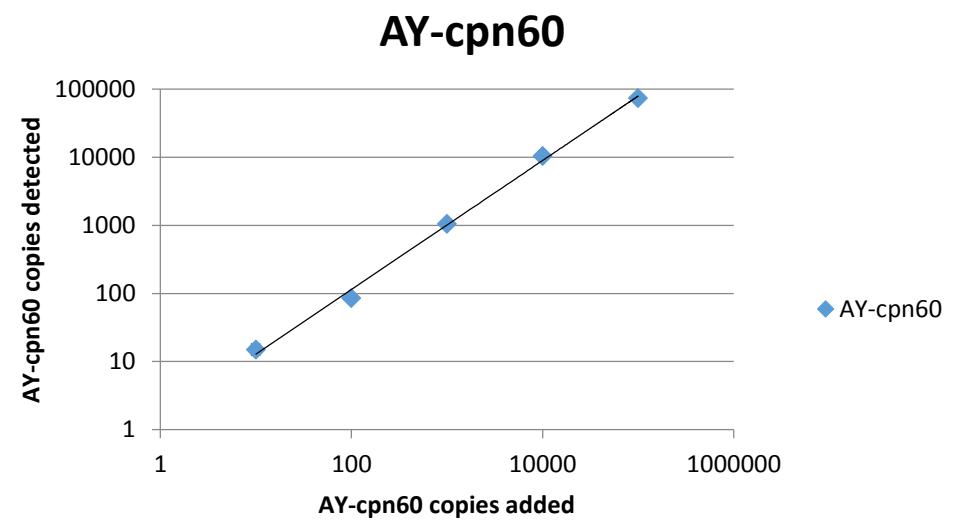
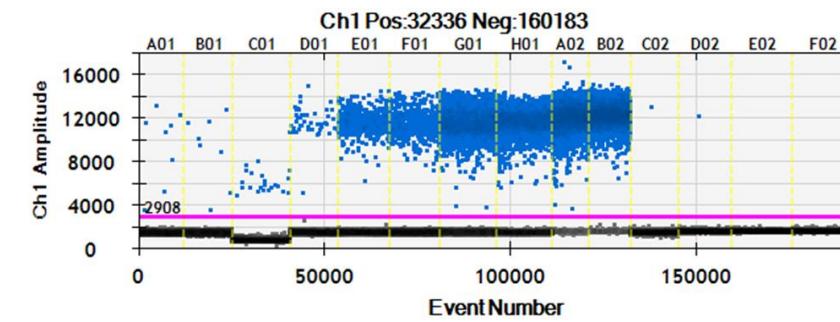
## Sequencing results



# qPCR targeting cpn60



# ddPCR



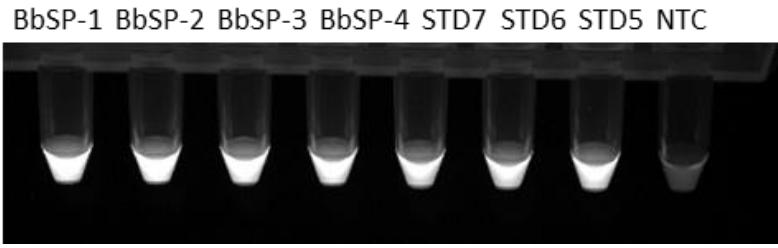
# LAMP- rapid molecular test of Aster yellows detection in a single leafhopper.



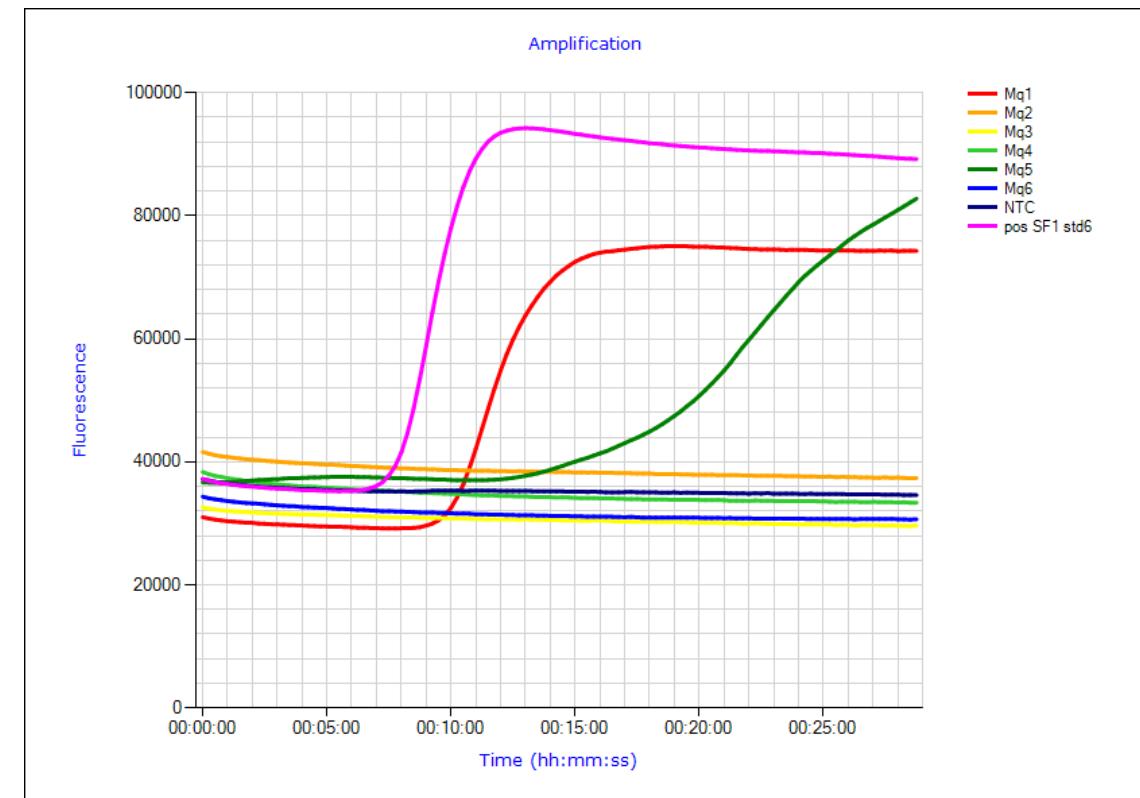
# Rapid, non-PCR amplification

## Loop-mediated isothermal DNA amplification (**LAMP**)

*cpn60*-targeted LAMP assay



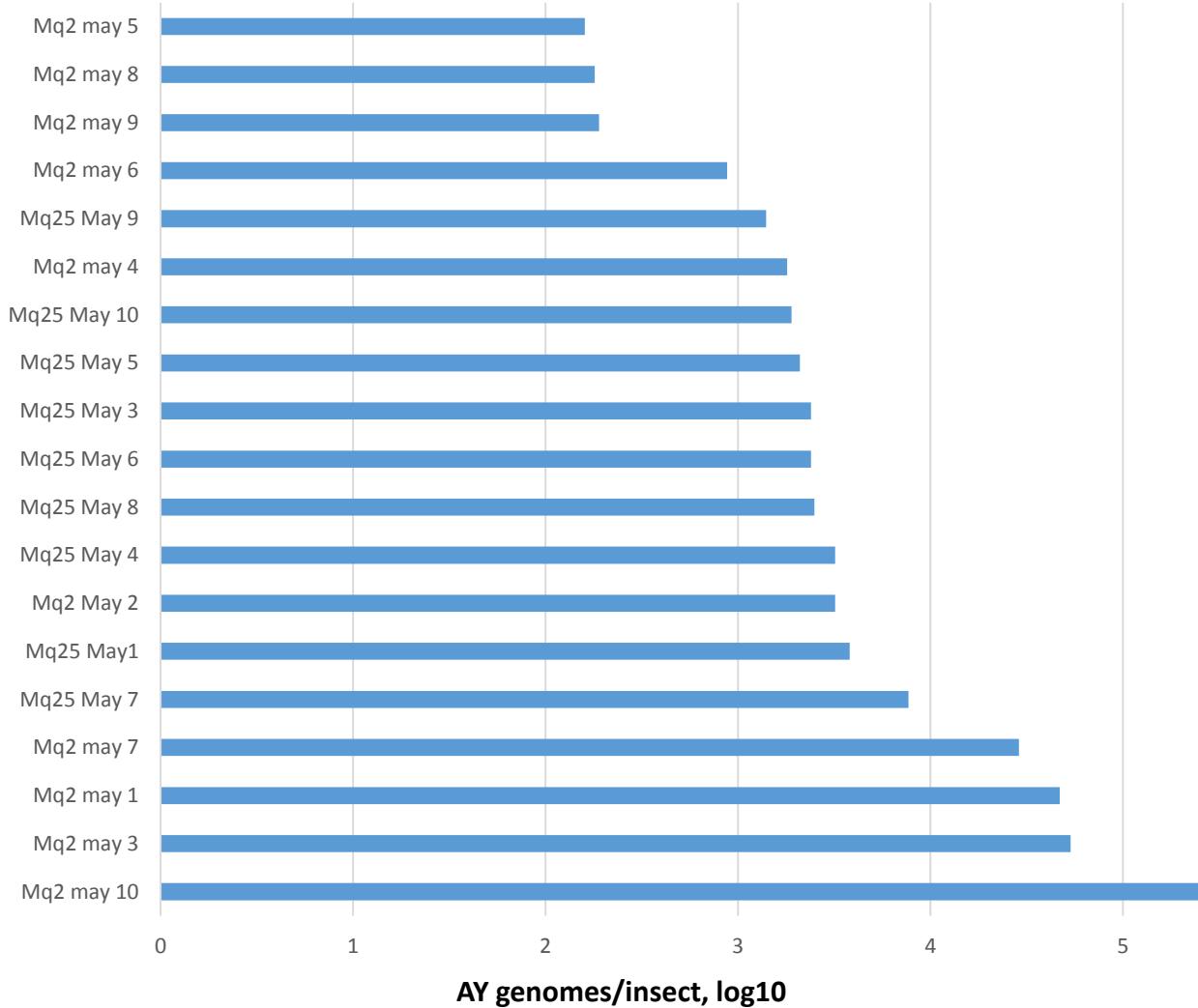
Binomial (pos/neg)



Time to positive (Tp)



## AY- ddPCR copies per insect



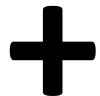
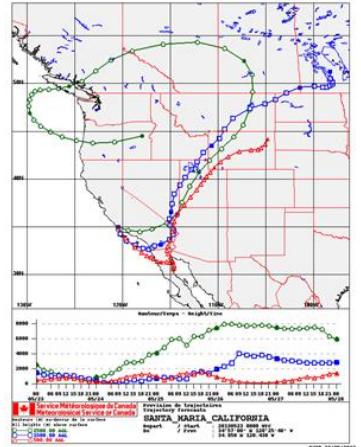
LAMP= ~ 1-2 hr total time

	Nested PCR- 16S	Nested PCR- rp	qPCR	LAMP
-	+	-	-	-
-	-	-	-	-
-	-	-	-	-
-	-	-	-	-
-	+	+	+	+
-	+	-	-	-
-	-	+	+	+
-	+	+	+	+
-	+	+	+	+
-	+	+	+	+
-	+	+	+	+
-	+	+	+	+
-	-	+	+	+
-	-	+	+	+
+	+	+	+	+
+	+	+	+	+
+	-	+	+	+
+	-	+	+	+
+	+	+	+	+

Nested PCR= ~ 1 week total time



# Take-home message



>KPB2\_LCO\_MP8014\_19 KPB2-SK3k6-Aberdeen\_CO1.0.2  
atgtgatcgttacatcccacgcatttattataatttccttatagtataccattataa  
ttgggggtttgtaattgactacttcacttataattggagctcctgatatacgattcc  
cacgactaaataatataagattctgacttctcccccttcattaacactattaatattaa  
gatcaatagtagaaatggggtagggacgggtgaacagtataccccccctatctagaa  
atatcgacatgcggcccaagagttgatatacaattttcttacatttagccgta



<sup>1</sup> H	<sup>2</sup> H	<sup>3</sup> H
1.00794 99.985%	2.0141 0.015%	3.0149 0.0001%
Stable	Stable	Cosmogenic anthropogenic
<sup>16</sup> O	<sup>17</sup> O	<sup>18</sup> O
16.00449 99.75%	16.00981 0.04%	16.00991 0.20%
Stable	Stable	Stable
<sup>12</sup> C	<sup>13</sup> C	<sup>14</sup> C
12.00069 98.95%	13.00335 1.11%	14.0 0.04%
Stable	Stable	Cosmogenic anthropogenic

= Origin of leafhoppers  
to Western Canadian  
Prairies

- quick, cheap, and reliable procedures to detect infected canola



# Acknowledgments

- Funding from Western Grains Research Foundation
- Christine Hammond, Edel Lopez, Dana Leedahl
- Owen Olfert<sup>1</sup>, Ross Weiss<sup>1</sup>, Meghan Vankosky<sup>1</sup> and Serge Trudel<sup>2</sup>

1 - AAFC

2 - ECCC





# Thank you

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