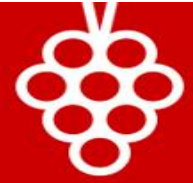


Advances in the diagnosis of grapevine virus diseases



Cool
Climate
Oenology &
Viticulture
Institute

Brock University



Outline

- History of plant virus detection
- Recent Advancements in virus detection
- Grapevine viruses – detection methods
- Applications of plant virus disease diagnostics

History

1892: Dmitry Ivanovsky: Tobacco mosaic disease could pass through a porcelain filter

1898: Martinus Beijerinck: *contagium vivum fluidum*

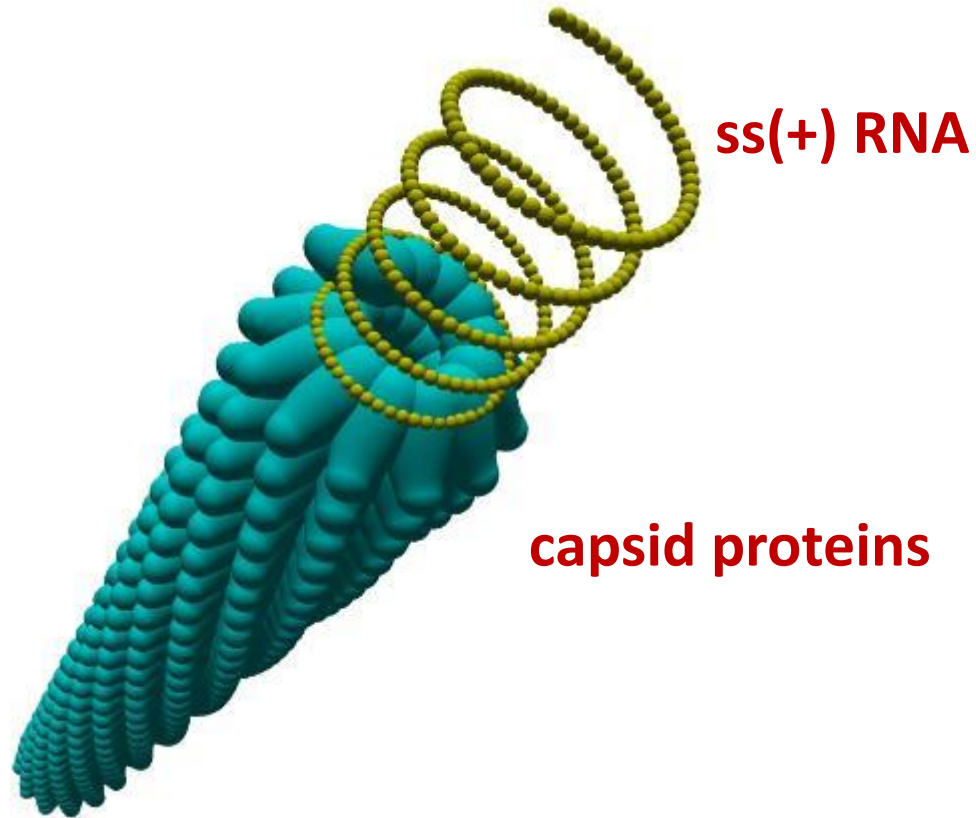
1931: Ernst Ruska: Electron microscope

“Plant viruses have obvious importance for food crops and ornamental plants, and a range of viruses are responsible for an estimated \$60 billion in crop losses worldwide each year”.

Lefeuvre et al. (2019). Evolution and ecology of plant viruses. *Nat Rev Microbiol* 17, 632–644

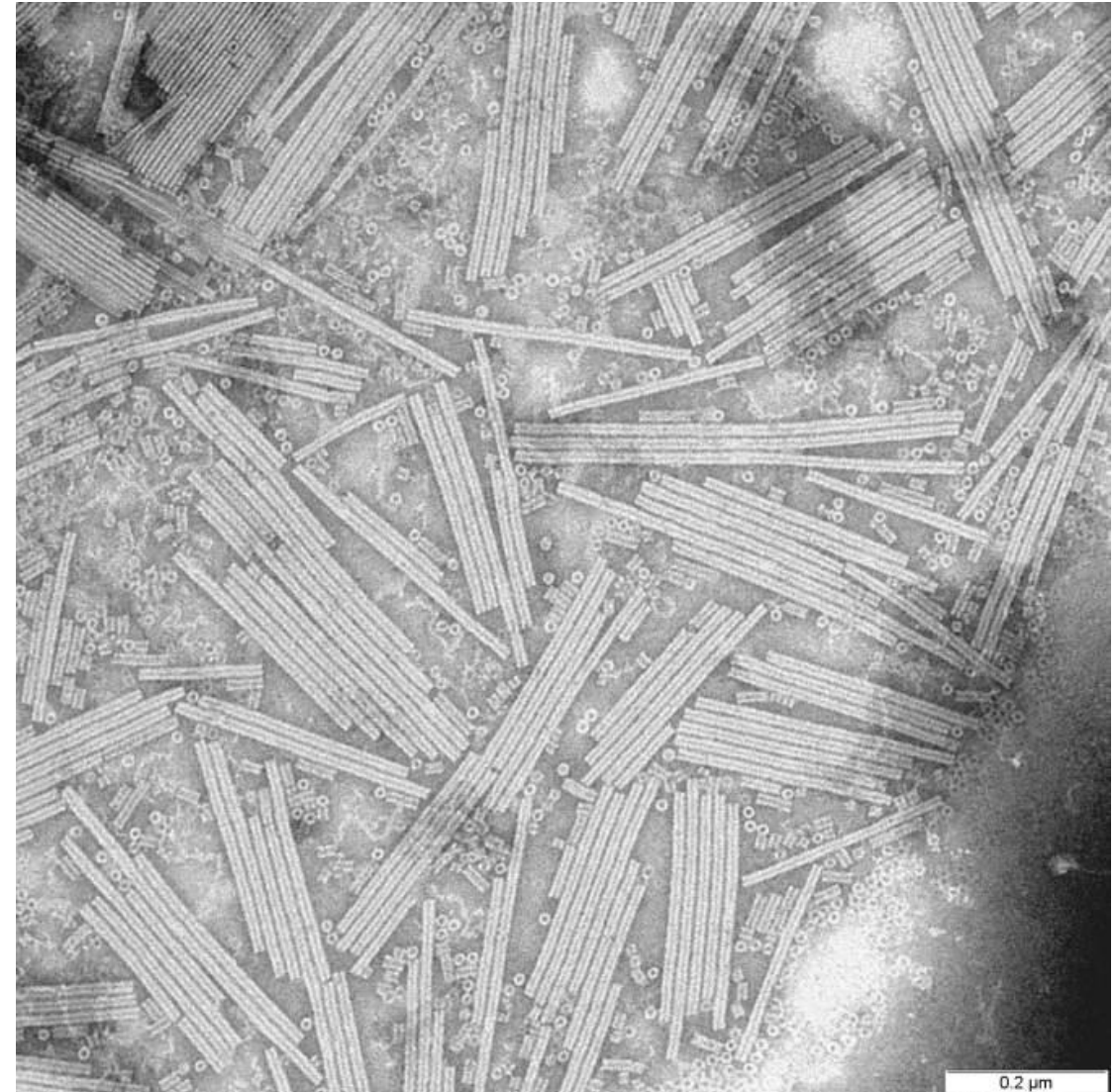
What is a virus

TMV (Tobacco Mosaic Virus)



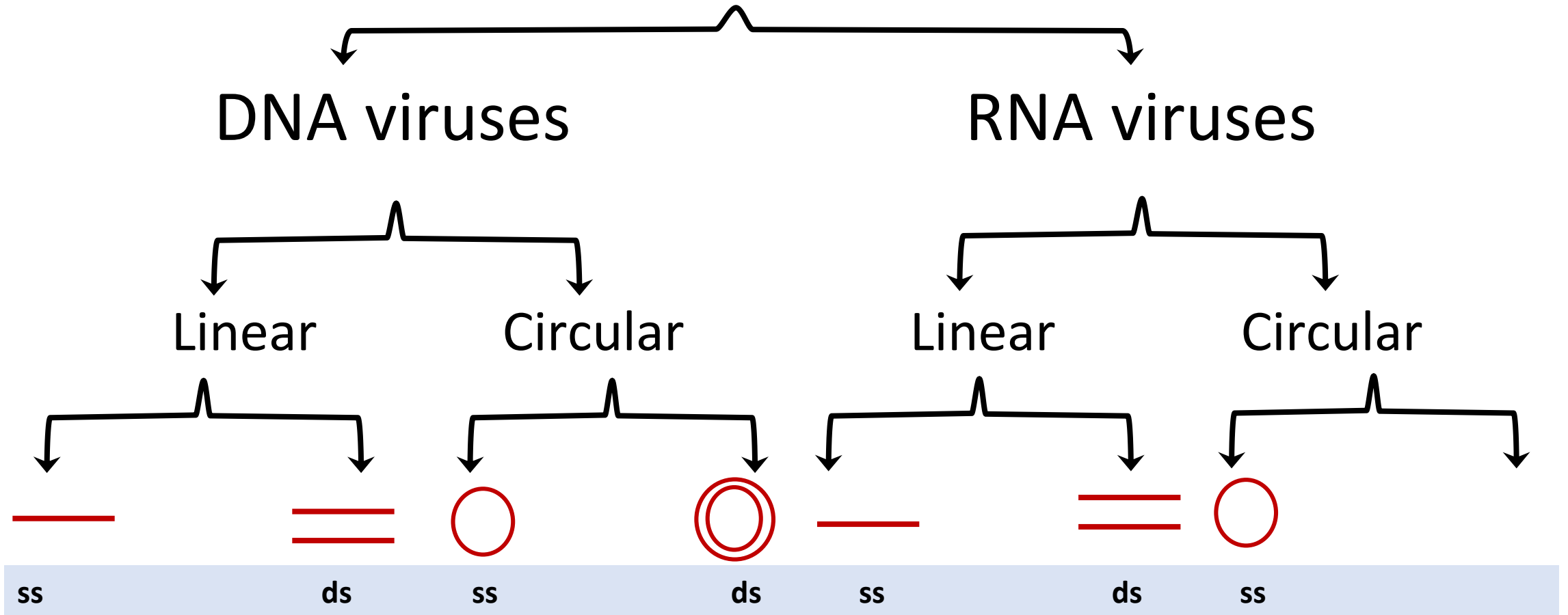
300 nm long by 18 nm in diameter

https://cronodon.com/BioTech/Virus_Tech.html



https://en.wikipedia.org/wiki/History_of_model_organisms





Virus Genome Diversity



Importance of Diagnostics

- Not all viruses show symptoms
- Cause of a specific pattern in symptoms
- Source? Where did the virus come from?
- How the virus transmitted?
- Spatial and temporal disease spread
- Plant Protection and Quarantine
- Impact on End Product – Mitigation
- Disease Management

Importance of Diagnostics in Grapevine Production

-  The sustainability of grapevine production system depends heavily on the health status of the propagating planting material being free of grapevine pathogens.
-  Grapevine virus diseases have been known to cause significant economic losses by negatively impacting the yield and quality of the wine.
-  The negative effects of grapevine virus diseases on rooting ability, graft in-take, vigour and berry quality have been demonstrated in commercial grapevine cultivars from distinct geo-climatic conditions.
-  Disease diagnostics has become extremely important for the effective management of the virus spread and mitigation practices.

Family *Closteroviridae*

Ampelovirus

Grapevine leafroll-associated virus 1

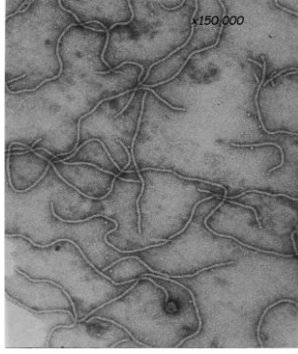
Grapevine leafroll-associated virus 3

Grapevine leafroll-associated virus 5

Little cherry virus 2

Pineapple mealybug wilt-associated virus 1

Pineapple mealybug wilt-associated virus 2



www.rothamsted.ac.uk

Closterovirus

Beet yellow stunt virus

Beet yellow virus

Burdock yellows virus

Carnation necrotic fleck virus

Carrot yellow leaf virus

Citrus tristeza virus

Grapevine leafroll-associated virus 2

Wheat yellow leaf virus



<http://cirs.ucr.edu>

Crinivirus

Abutilon yellows virus

Beet pseudoyellows virus

Cucurbit yellow stunting disorder virus

Lettuce infectious yellows virus

Sweet potato chlorotic stunt virus

Tomato chlorosis virus

Tomato infectious chlorosis virus

Potato yellow vein virus



www.bayercropscience.com

Karasev, 2000 *Annu. Rev. Phytopathol.* 38: 293–324

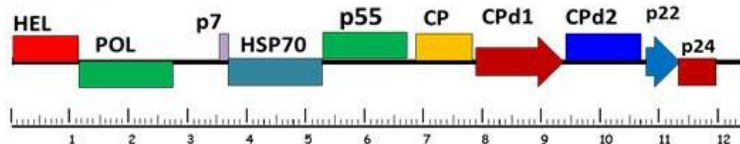
Martelli et al., 2002 *Arch. of Virol.* 147: 2029–2044

[www. ICTV.org](http://www.ICTV.org)

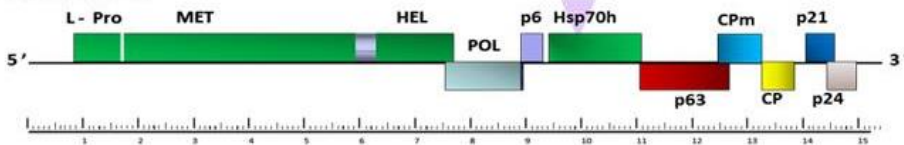
Family *Closteroviridae*

Genome Complexity

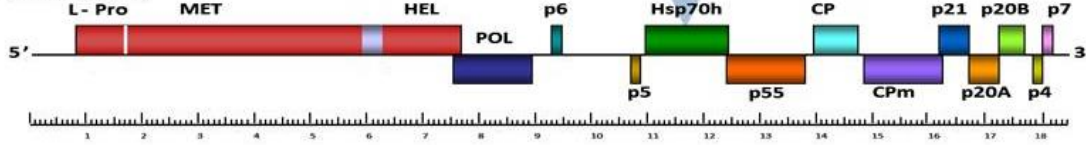
GLRaV-1



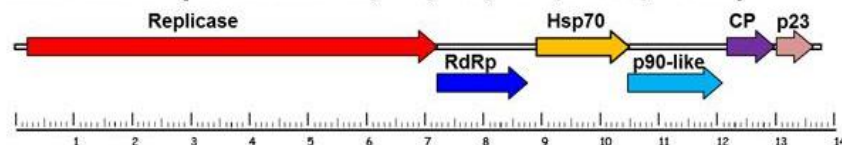
GLRaV-2



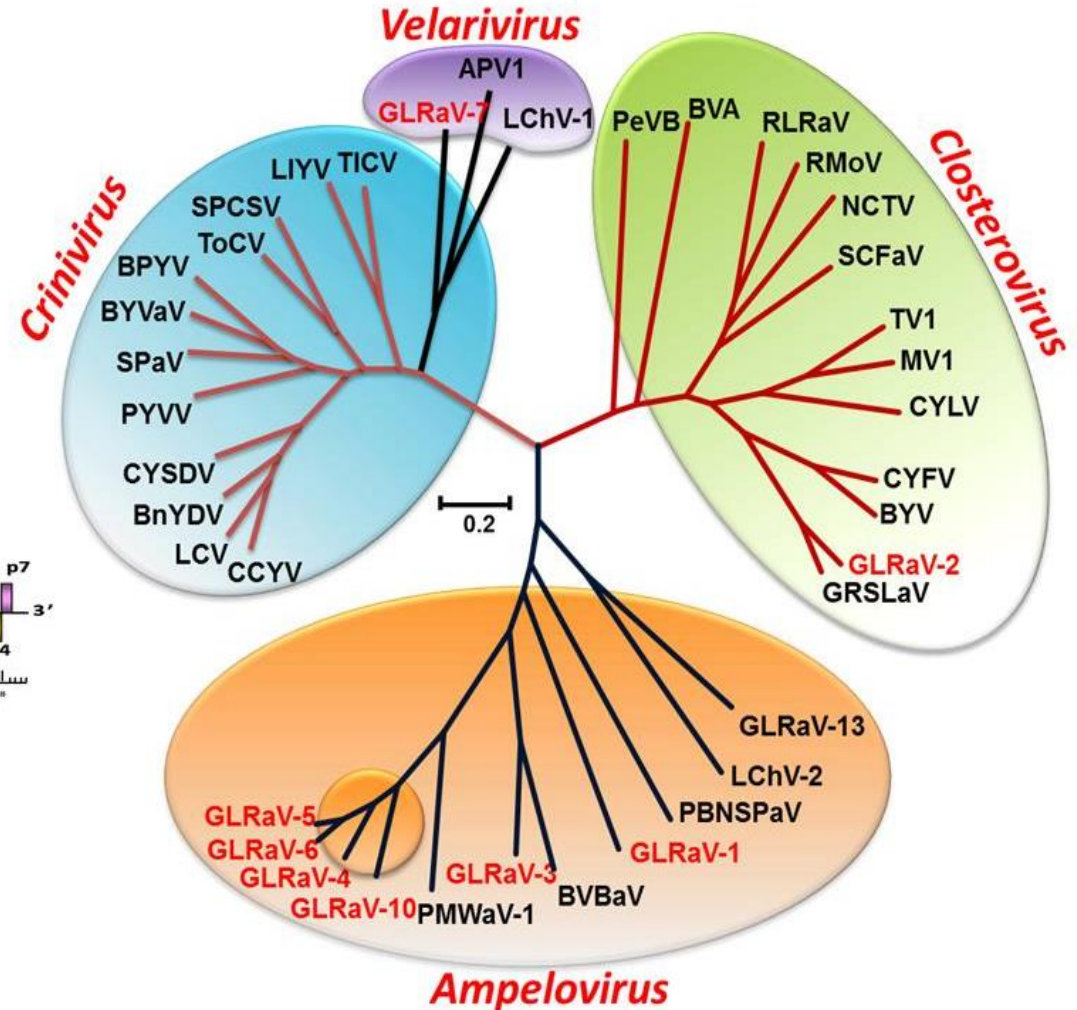
GLRaV-3



GLRaV-4 (strains: -5, -6, -9, -Pr, -De, -Car)

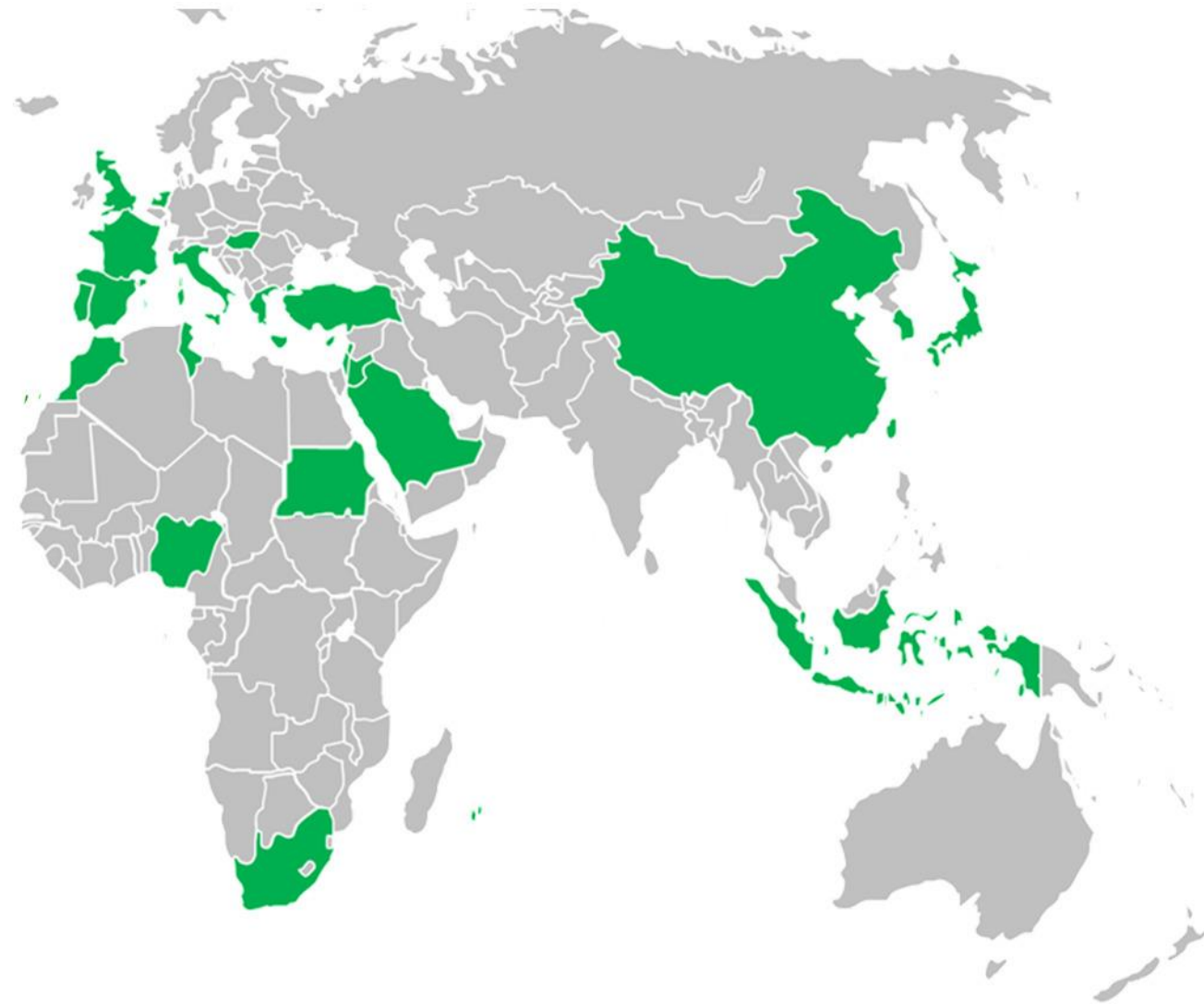
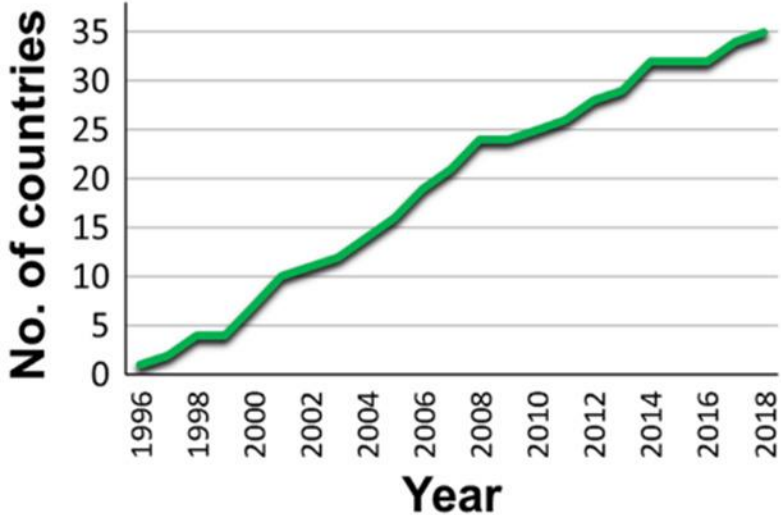
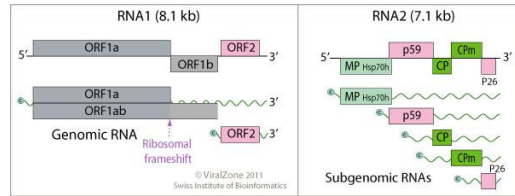


Phylogenetic Tree



Geographical distribution of Tomato Chlorosis Virus

Genome



Elvira Fiallo-Olivé and Jesús Navas-Castillo. 2019.
10.1111/mpp.12847

Family *Geminiviridae*

Symptoms

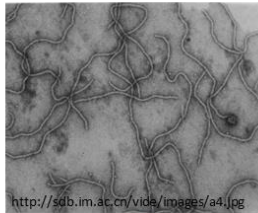
Grapevine leafroll virus



Grapevine red blotch virus

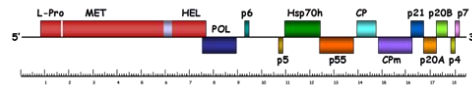


Virus particles

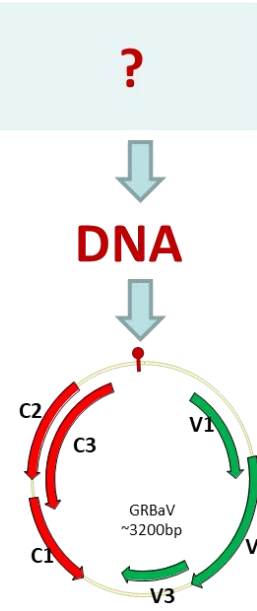


Genetic material

RNA



Genome Organization

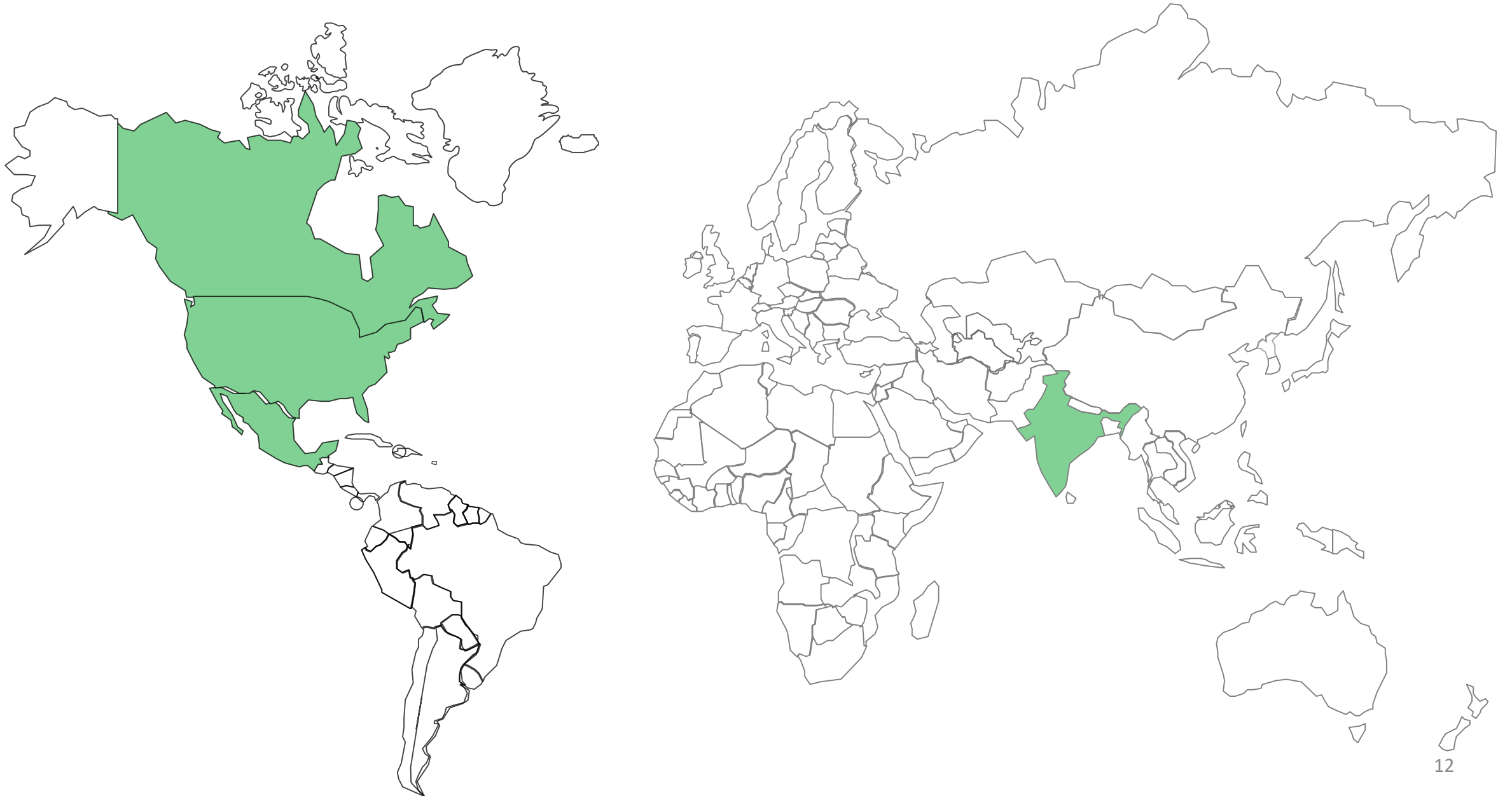


Geminiviridae

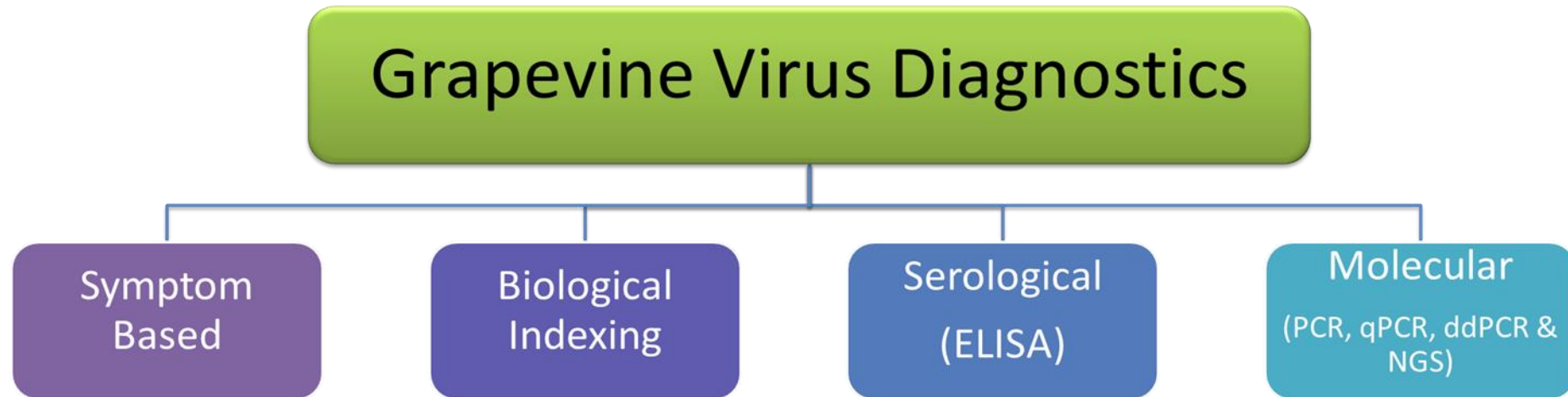


DOI: 10.1007/s00705-017-3268-6

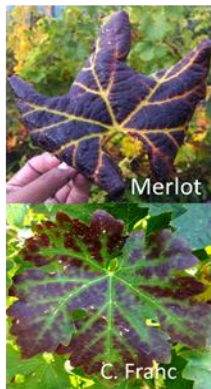
Geographical distribution of Grapevine Red Blotch Virus



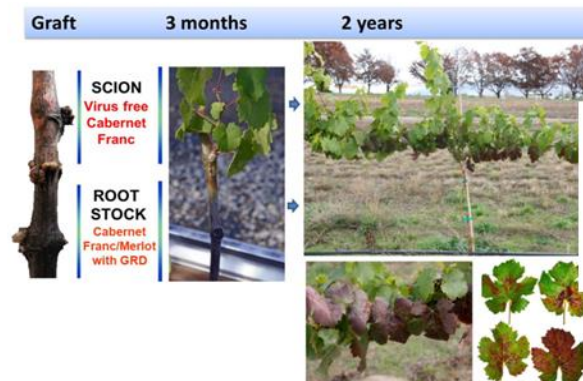
How to detect grapevine viruses?



Non-specific
Inaccurate

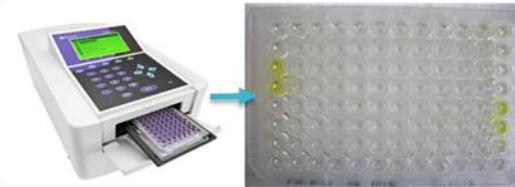


Symptom based
Labor intensive
Time consuming



Poojari et al. 2013. PLoS ONE 8(6): e64194

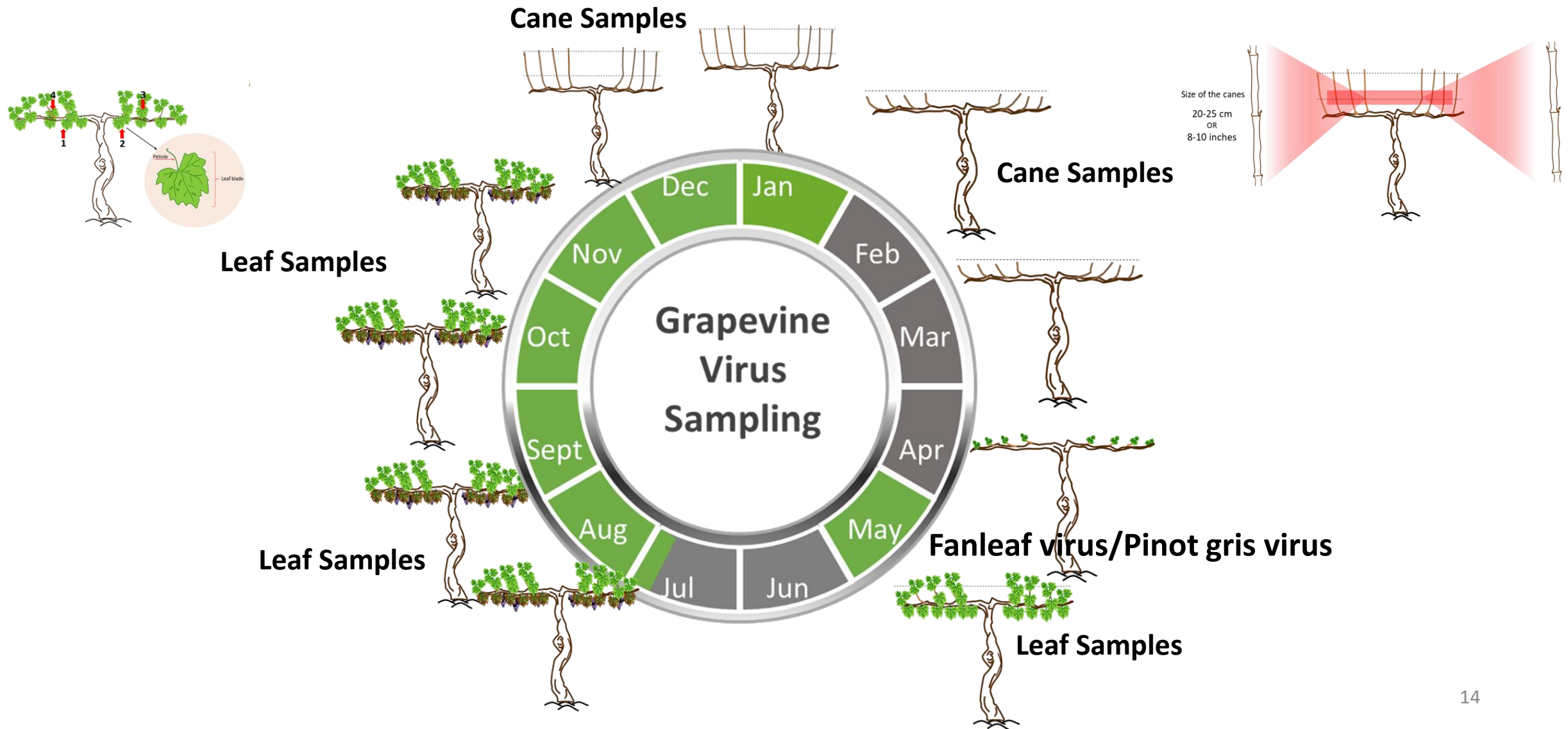
Specific
No Abs for all viruses
Less sensitive than PCR



Highly specific
Post-PCR process
Multiplex
End-Point
Quantitative

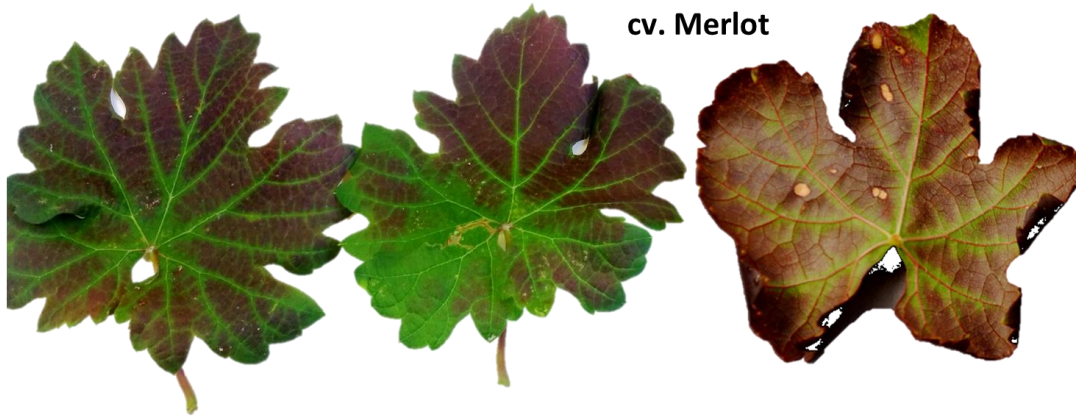


What is the preferred sampling time and tissue type for virus detection?



Symptom based detection

Grapevine leafroll-associated viruses



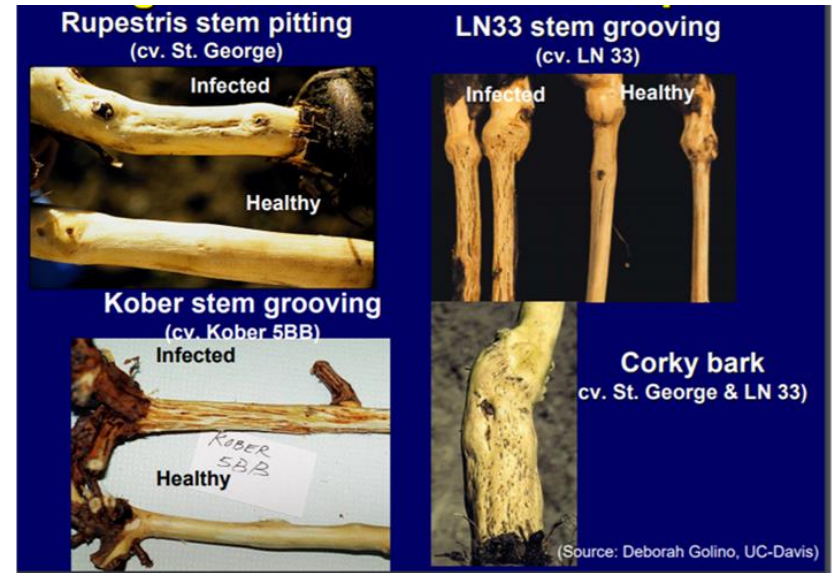
Grapevine red blotch virus



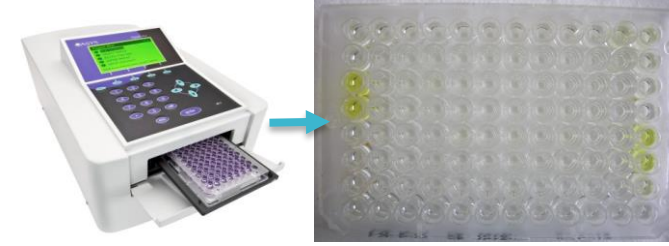
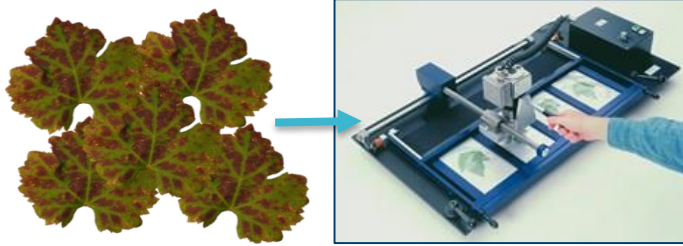
Grapevine fanleaf virus



Grapevine Pinot Gris Virus



ELISA

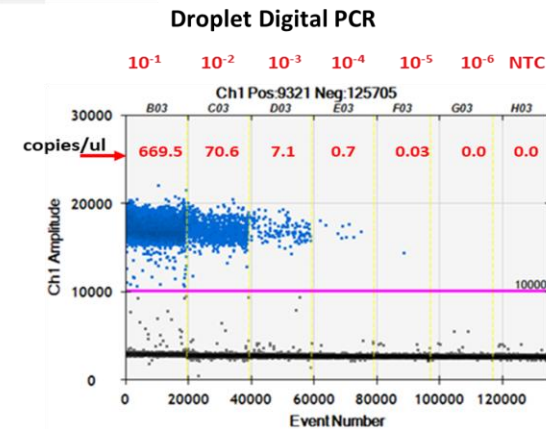
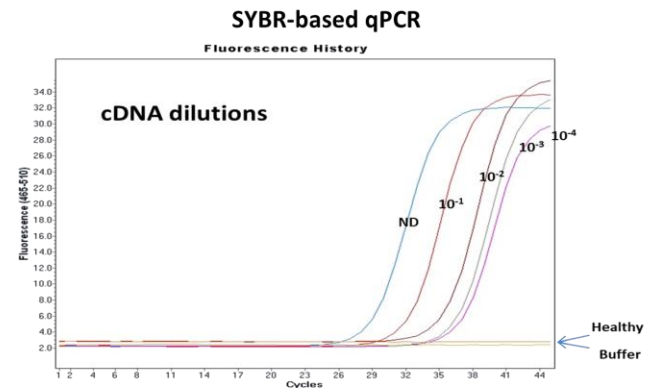
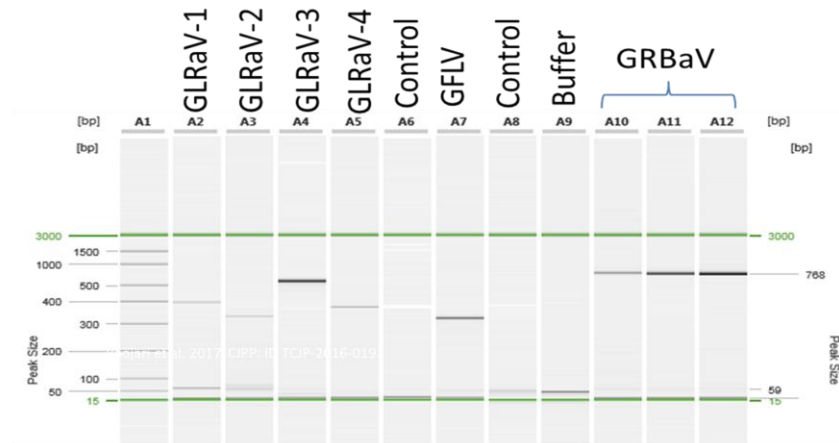
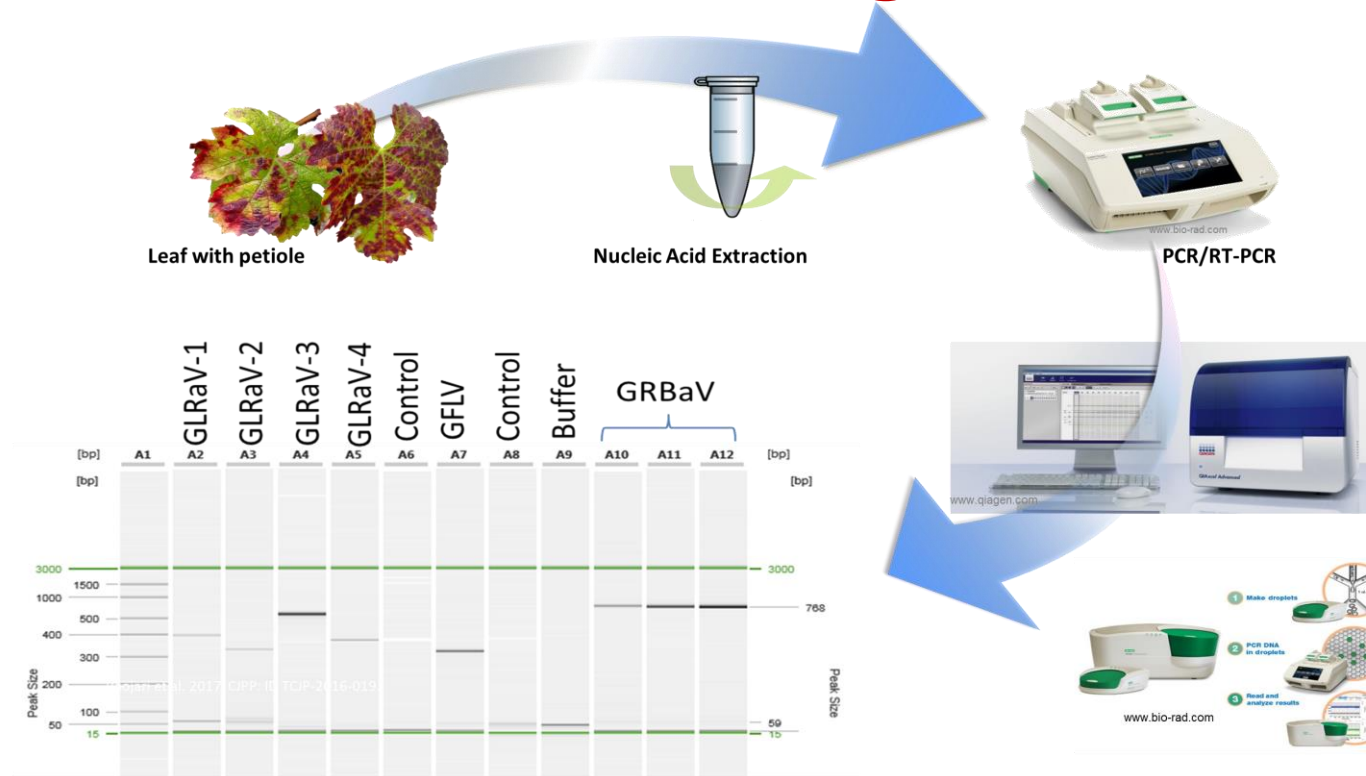


Advantages	Limitations
Less technical knowledge Large scale detection	Extraction of large number of samples in short time Time consuming Antibody availability is limited Nonspecific reactions Quality of antibodies Detection limitations with virus variants/strains

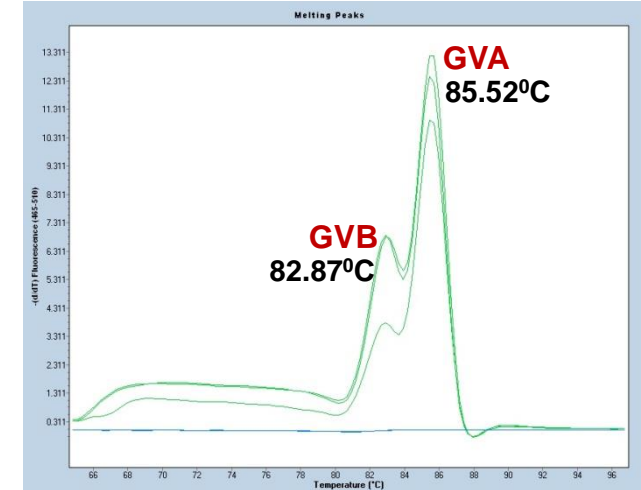
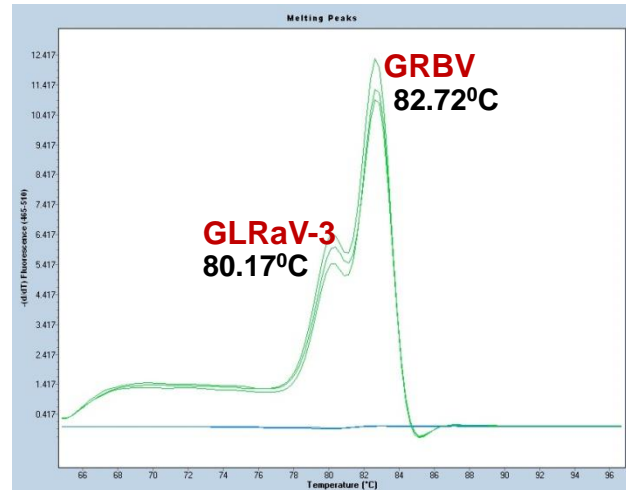
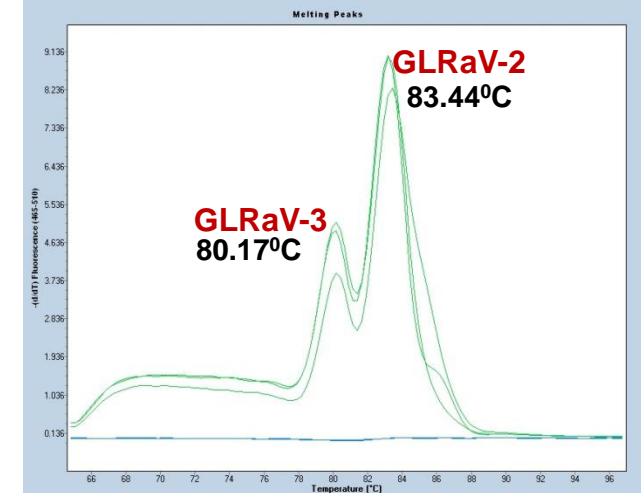
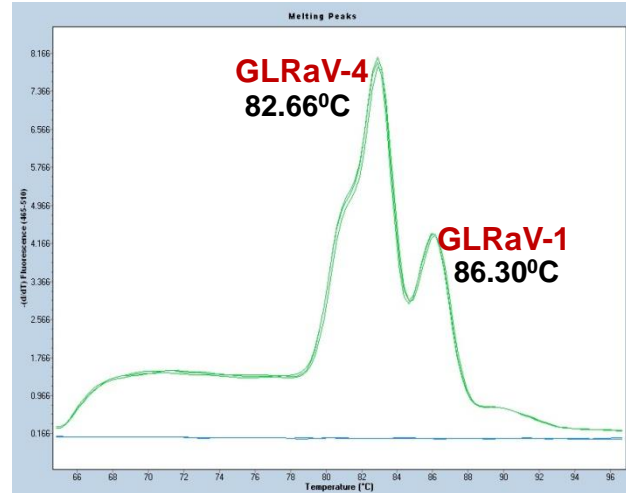
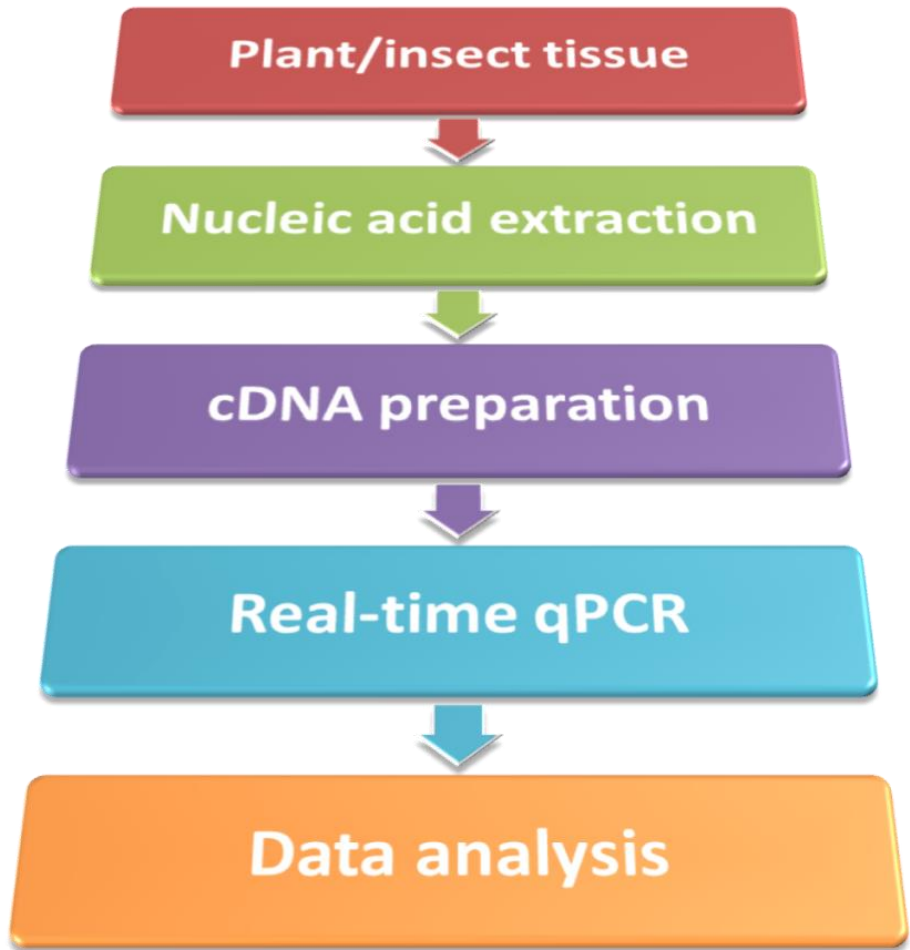
Arabis mosaic virus
Grapevine fanleaf virus
Raspberry ringspot virus-ch
Raspberry ringspot virus-g
Tomato black ring virus
Tobacco ringspot virus
Tomato ringspot virus
Tomato ringspot virus-Ch
Strawberry latent ringspot virus

Grapevine leafroll-assoc. virus 1
Grapevine leafroll-assoc. virus 2
Grapevine leafroll-assoc. virus 3
Grapevine leafroll-assoc. virus 4 and strain 6
Grapevine virus A
Grapevine fleck virus
Grapevine pinot gris virus

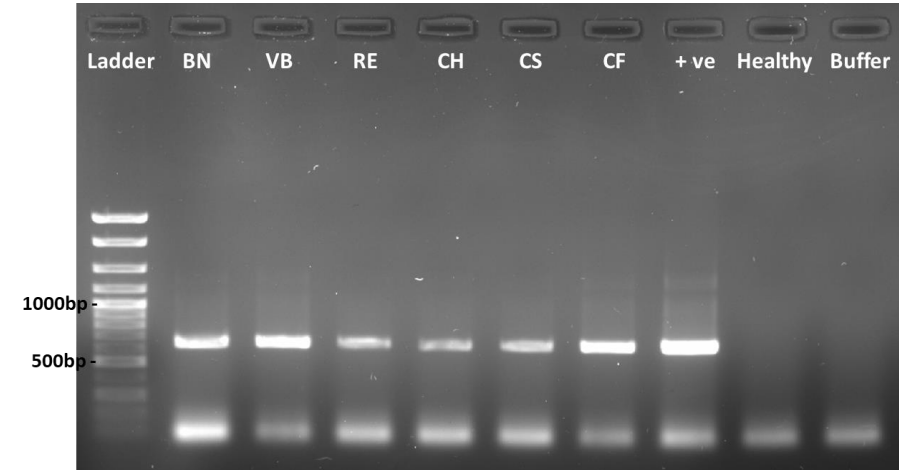
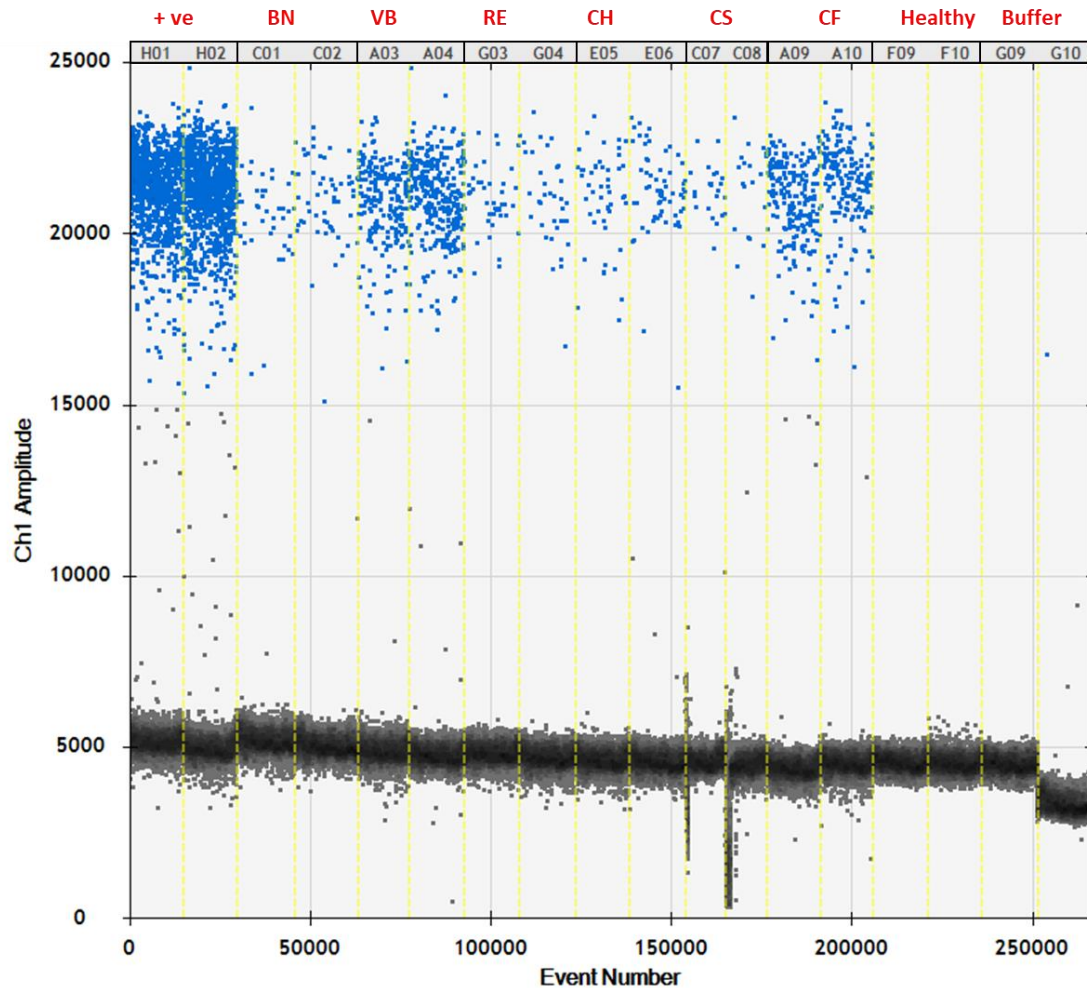
PCR-based diagnostics



Validation of Duplex RT-qPCR



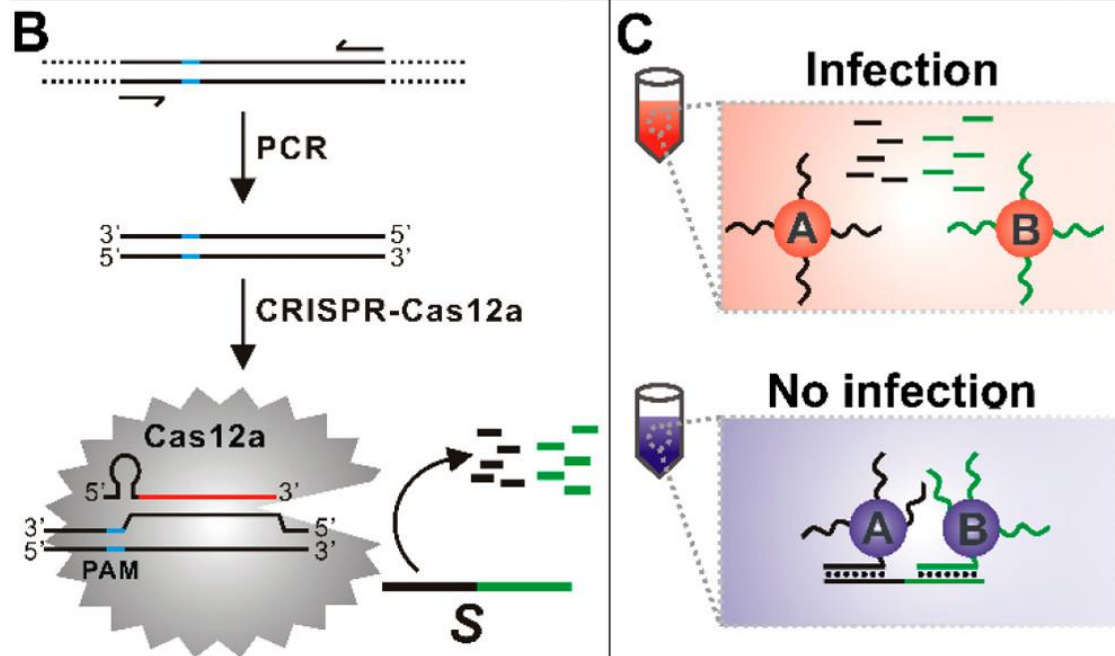
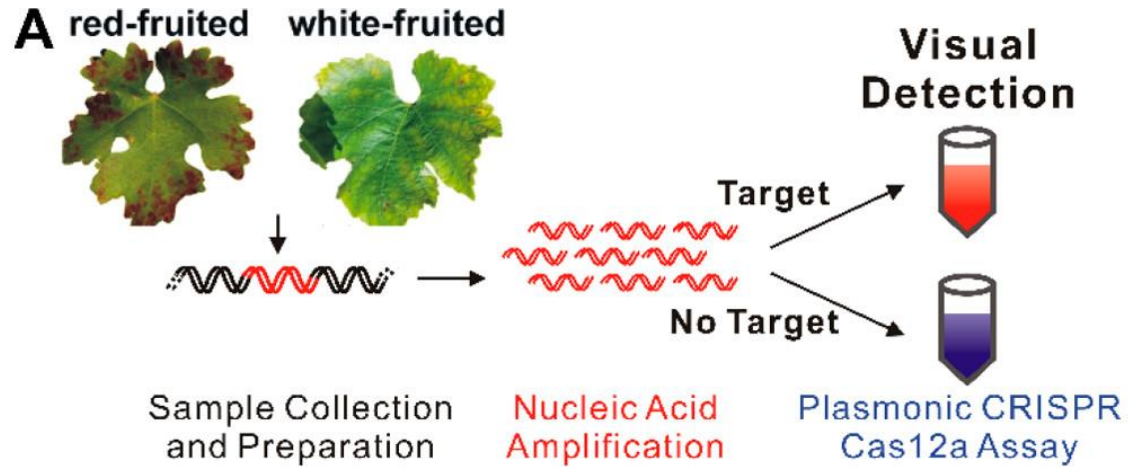
Detection of GRBV by Droplet Digital™ PCR (ddPCR™)



End-point PCR

Fluorescence amplitude plot representing the absolute quantification of grapevine red blotch virus (GRBV) DNA from 6 different isolates. BN-BacoNaoir; VB-Vidal Blanc; RE-Riesling; CH-Chardonnay; CS-Cabernet Sauvignon; CF-Cabernet Franc

Plasmonic CRISPR Cas12a assay for GRBV



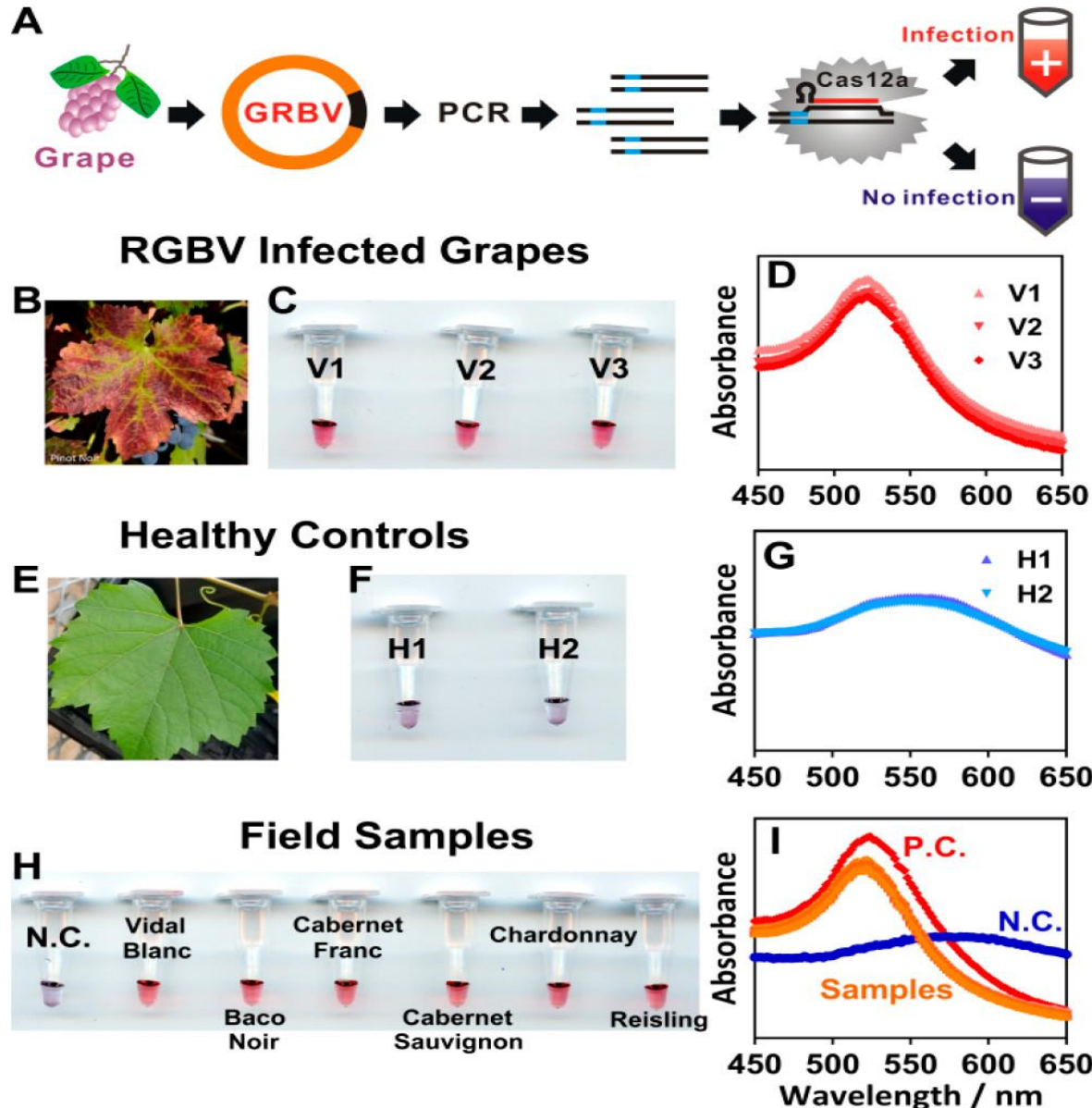
- (A) Workflow for visual detection of GRBV infections in red-fruited and white-fruited grapevine samples.
- (B) Target recognition using CRISPR-Cas12a and the activation of incriminate ssDNase activity.
- (C) Color development using plasmonic DNA functionalized AuNPs.

Yongya Li et al., Naked-Eye Detection of Grapevine Red-Blotch Viral Infection Using a Plasmonic CRISPR Cas12a Assay
Analytical Chemistry. 2019. 91. 18. 11510-11513

Cool Climate Oenology and Viticulture Institute, Brock University

Department of Chemistry, Centre for Biotechnology, Brock University

Plasmonic CRISPR Cas12a assay for GRBV



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Yongya Li et al., Naked-Eye Detection of Grapevine Red-Blotch Viral Infection Using a Plasmonic CRISPR Cas12a Assay
***Analytical Chemistry*. 2019. 91. 18. 11510-11513**

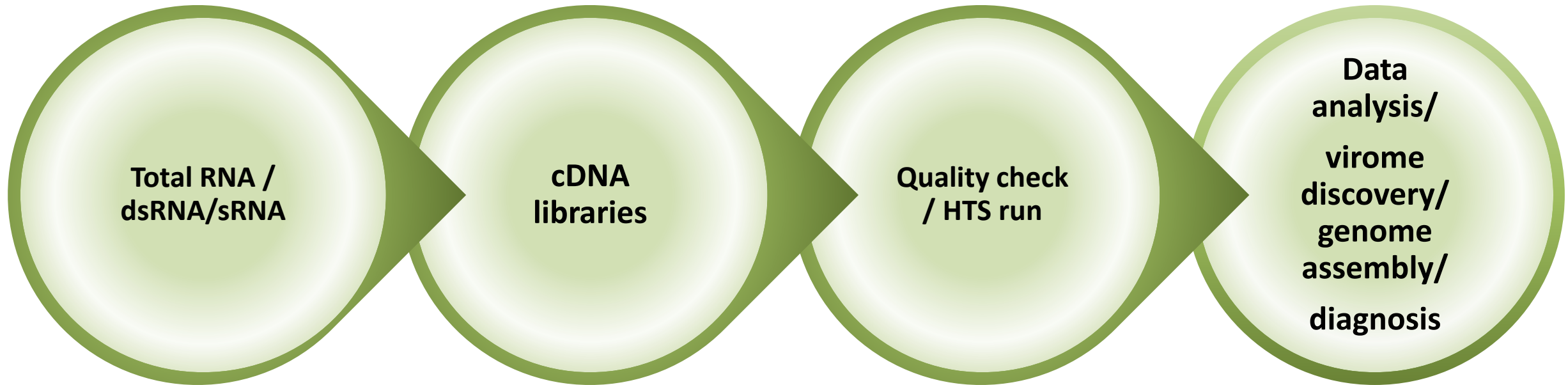
Cool Climate Oenology and Viticulture Institute, Brock University

Department of Chemistry, Centre for Biotechnology, Brock University

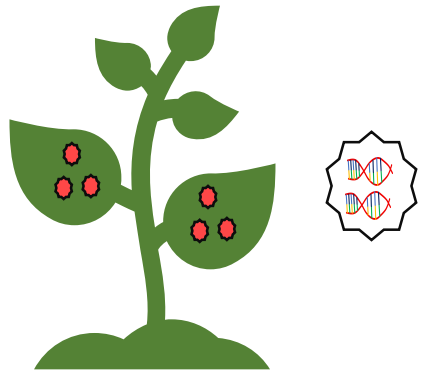
PCR-based diagnostics

	Conventional PCR	Real-time PCR	Digital PCR
Detection	Yes	Yes	Yes
Quantitative	Semi	Relative	Absolute
Advantages	Sophisticated Rapid (in hours) Downstream applications <ul style="list-style-type: none"> - Sequencing - Evolutionary analysis 	Increased sensitivity Records amplification as it happens No post-PCR processing Rapid (in minutes) Require less template melting curve analysis	Not dependent on house keeping genes / standards Highly accurate and useful to differentiate minor fold changes No post-PCR processing Require less template
Limitations	Size based results Post-PCR processing	Depend on reference genes Non-specific binding? Extensive optimization? Long amplicons? Multiplexing needs skills and knowledge	Large scale? Long amplicons? Multiplexing needs skills and knowledge

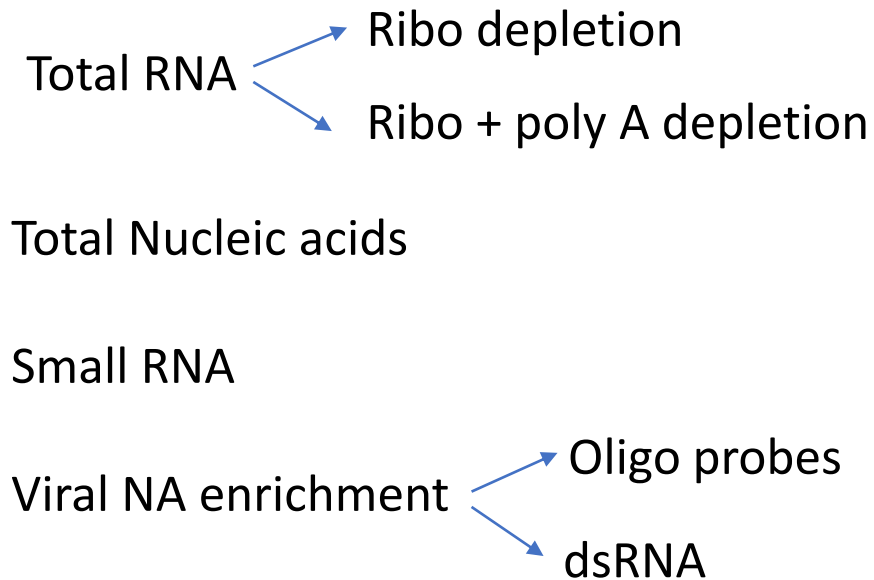
High Throughput Sequencing (HTS) based virus detection



High Throughput Sequencing (HTS) for virus detection



Virus
suspected
plant
sample



Quality Check

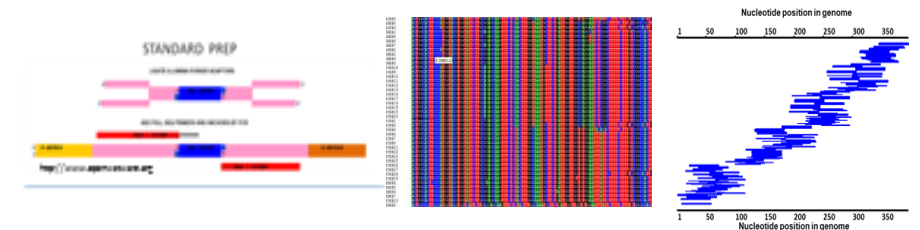
NGS sequence reads

Quality Check \ Adaptor sequence filter

De novo assembly of sequence reads

Blastn: host genome \ plant virus database


















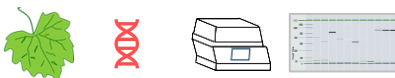



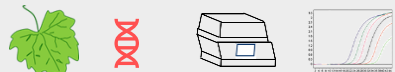






Virome discovery \ diagnosis



Virtool

Viral infection diagnostics using next-generation sequencing.

Virus Detection Methods and Applications

METHOD	HQP	EQP+DATA	TIME	APPLICATION
VISUAL				
BIOLOGICAL				
SEROLOGICAL				
-ELISA				
-LFIA				
PCR				
-End-point				
-qPCR/ddPCR				
HTS				

Acknowledgments

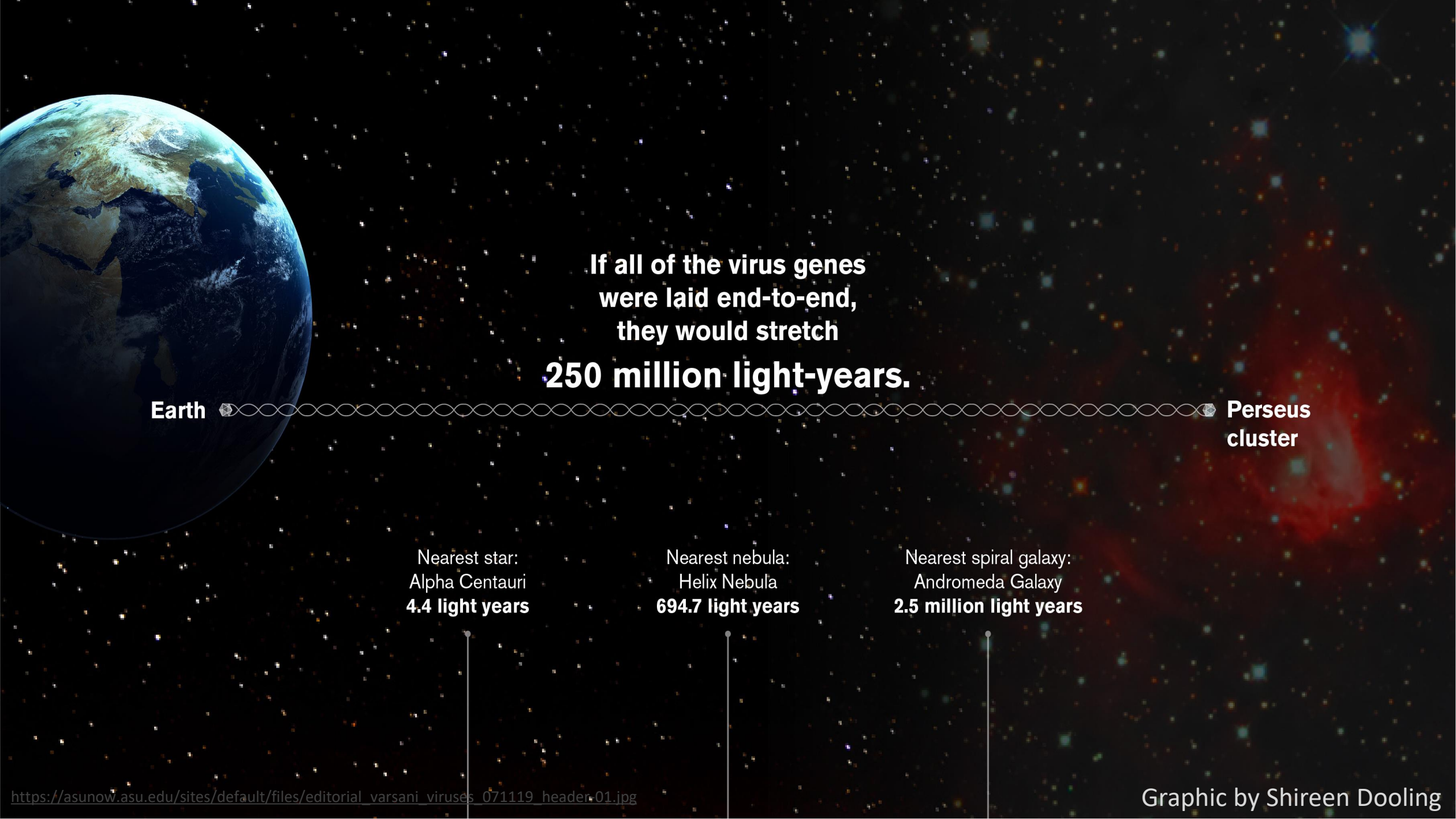


Educational Resources

- [Improvements in Grapevine virus diagnostics: Sudarsana Poojari](#)
- [Epidemiology and management of grapevine virus diseases: Tom Lowery](#)
- [Demystifying the status of grapevine viruses in British Columbia: José Ramón Úrbez Torres](#)
- [OMAFRA Grapevine leafroll virus](#)
- [OMAFRA: Grapevine red blotch](#)

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Tel: 905 688 5550 ext. 4227
Email: spoojari@brocku.ca
<https://brocku.ca/ccovi/>



If all of the virus genes
were laid end-to-end,
they would stretch
250 million light-years.

Earth

Perseus
cluster

Nearest star:
Alpha Centauri
4.4 light years

Nearest nebula:
Helix Nebula
694.7 light years

Nearest spiral galaxy:
Andromeda Galaxy
2.5 million light years