

# Barley yellow dwarf virus (BYDV)

## Challenges & Opportunities for Effective Management

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13/01/2025



ROTHAMSTED  
RESEARCH







- Context
  - Collaborations are key
  - Barley yellow dwarf virus
- Challenges
  - Traditional control
  - Genetic resistance/tolerance?
  - Climate change (vectors? new strains?)
  - Rothamsted Insect Survey
  - What's out there?
- Opportunities
  - Diagnostics and high-throughput testing
  - Sampling

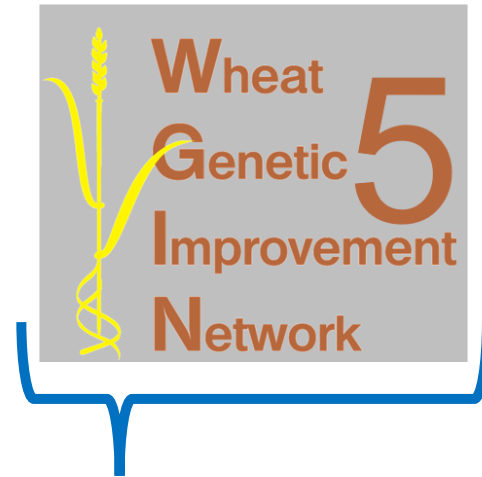


# Genetic Improvement Networks



Department for Environment Food & Rural Affairs

Defra supports the five GINs to improve major UK crop varieties:



## Wheat Genetic Improvement Network (WGIN):

- **BYDV** = previous UK viral target, now diversifying
- Diverse management team and stakeholders



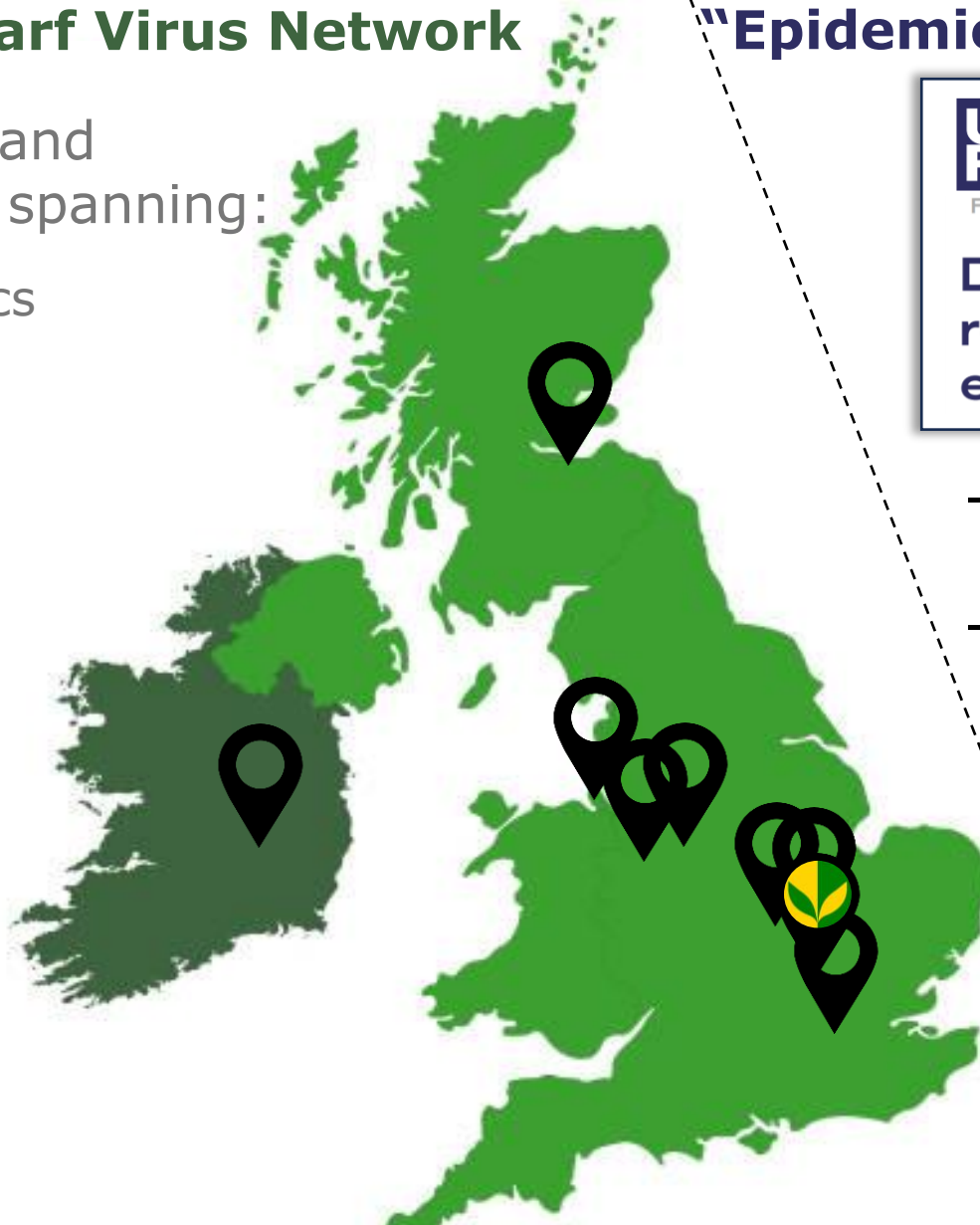




## UK & Ireland Yellow Dwarf Virus Network

Virologists, entomologists and modellers at key locations spanning:

- Biomathematics & Statistics Scotland
- University of Liverpool
- Keele University
- Harper Adams University
- Teagasc
- University of Cambridge
- AHDB
- ADAS
- Rothamsted Research
- University of Greenwich
- **More and growing...**



## "Epidemic Preparedness" Team



**Develop interdisciplinary research proposals to tackle epidemic threats**

- Workshop last Thu/Fri
- >20 experts tasked with evaluating the:

***"tools, technologies and policy devices required to tackle vector-borne viral and bacterial epidemics in agriculture"***



Subgroup within Wheat Pathogenomics Team  
(RRes, PI: Prof. Kim Hammond-Kosack)

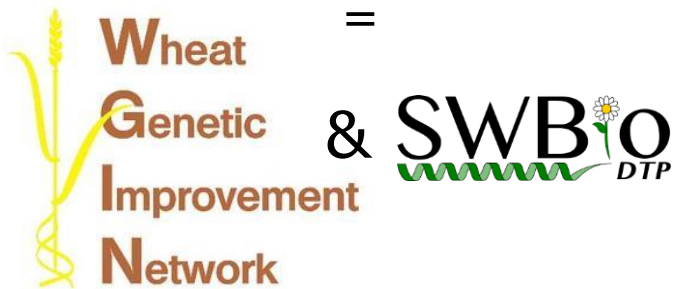


- Izayana Sandoval-Carvajal  
SWBio DTP PhD student (RAGT, Uni of Bristol)



- Postdoc / subgroup lead

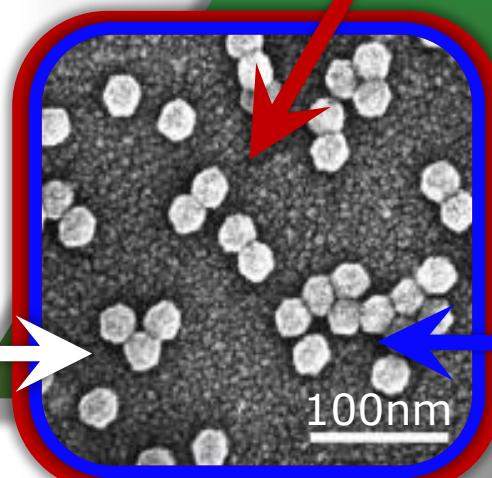
BYDV work =



Aphid resistance work =



Diagnostics



Viruses (BYDV)



Aphids



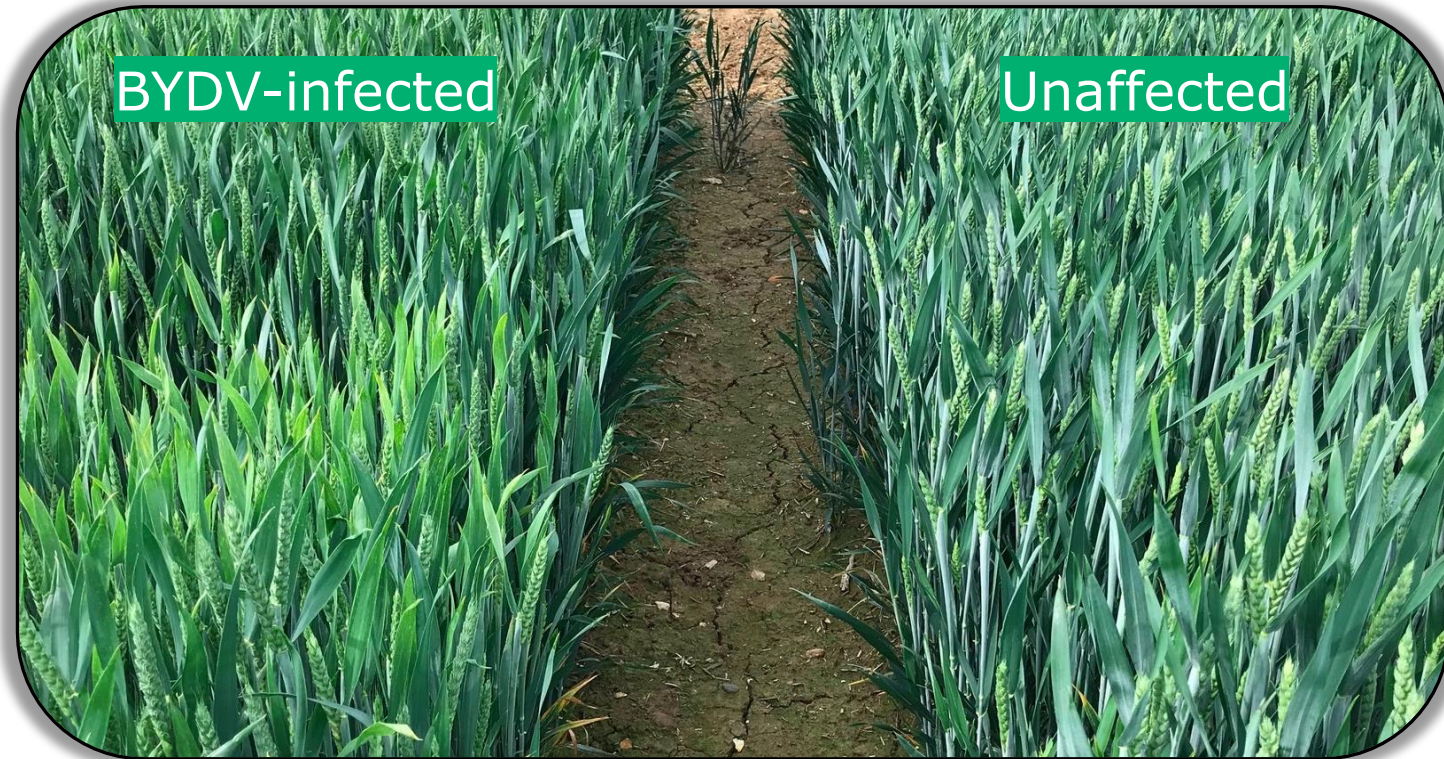
Wheat



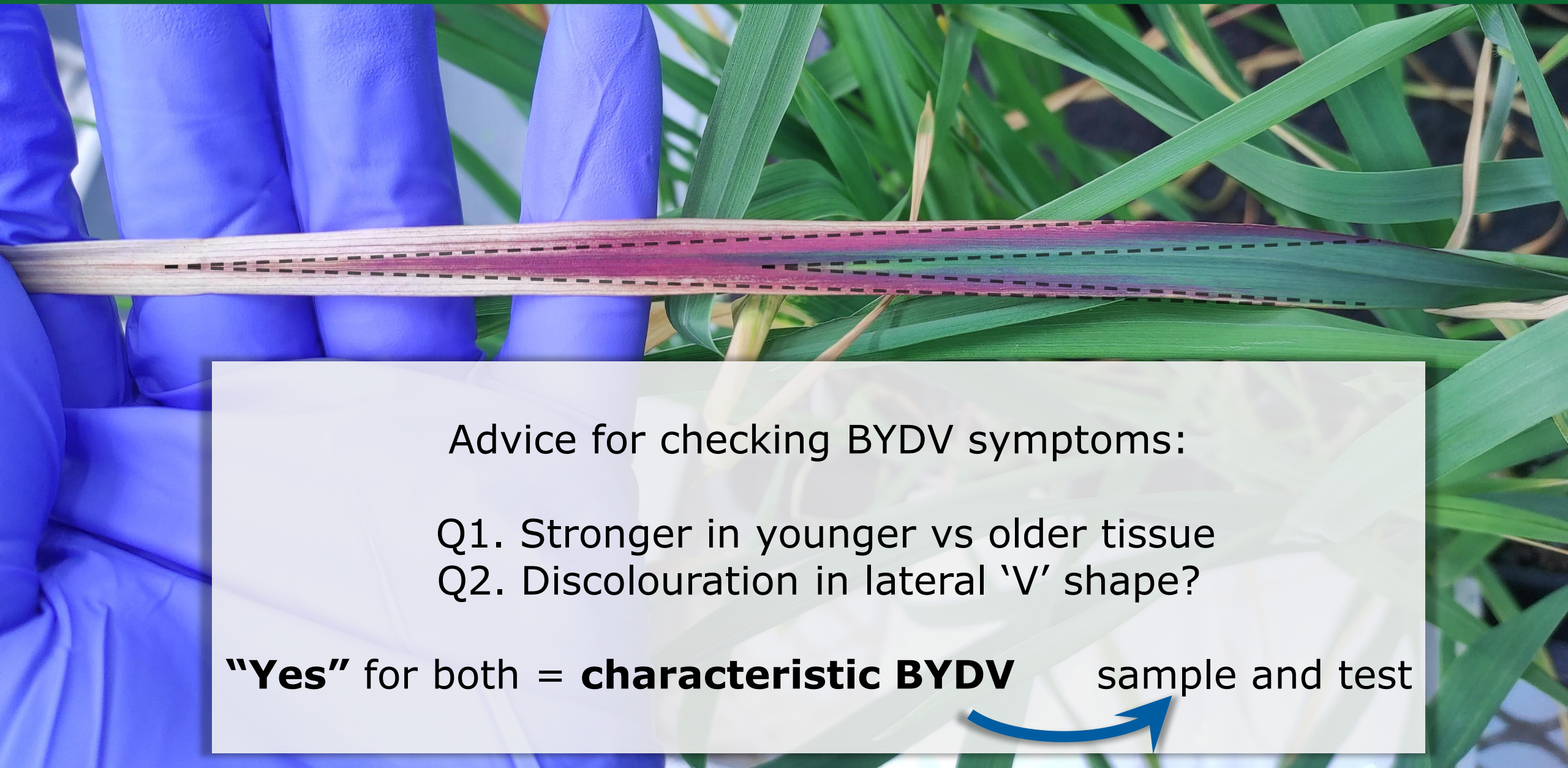




- Harvest yield loss in wheat due to natural BYDV infection = **5 - 80%**, average of **30%**\*
- Indistinct disease phenotype limits rapid diagnosis
- No practical virus-focussed chemical control
- Aphid vectors targeted, and "resistant" cultivars developed
- Interactions between aphid vector species and BYDV strains = complex







Advice for checking BYDV symptoms:

Q1. Stronger in younger vs older tissue

Q2. Discolouration in lateral 'V' shape?




**"Yes"** for both = **characteristic BYDV** sample and test

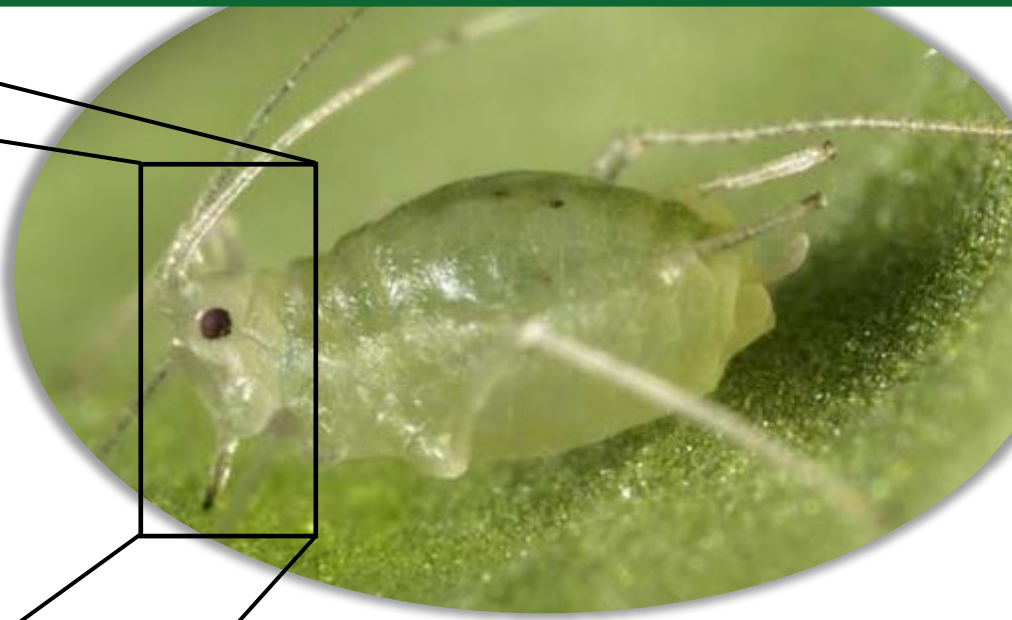
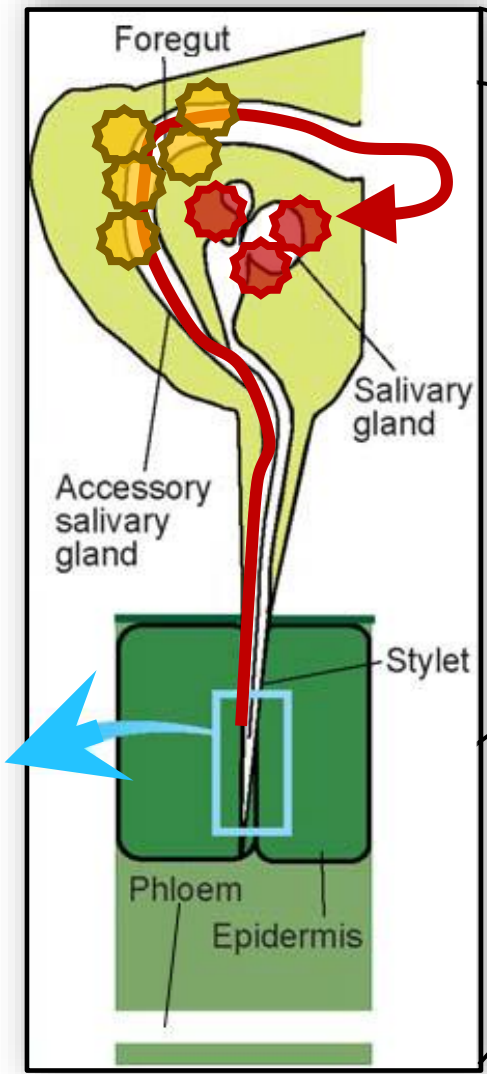
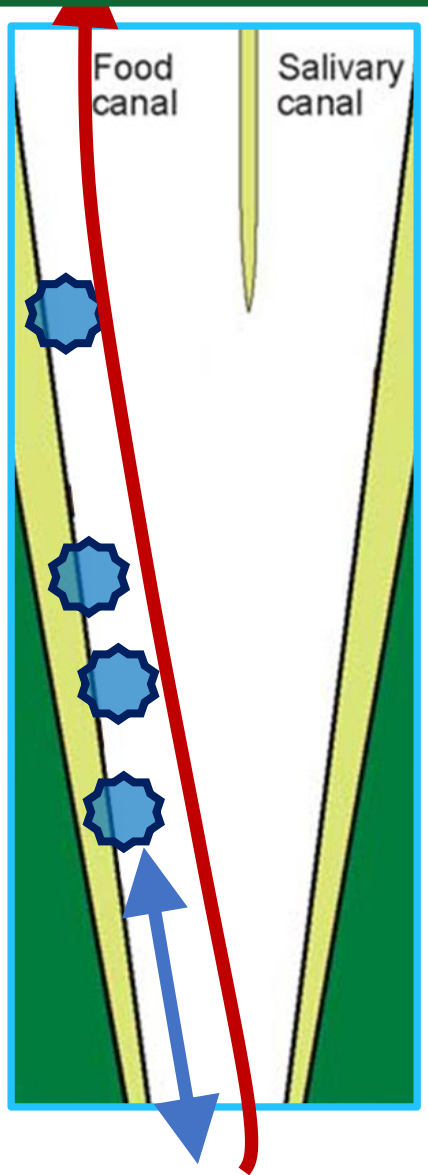






# Aphid viral transmission

-  = Non-persistent
-  = Persistent "circulative"
-  = Semi-persistent

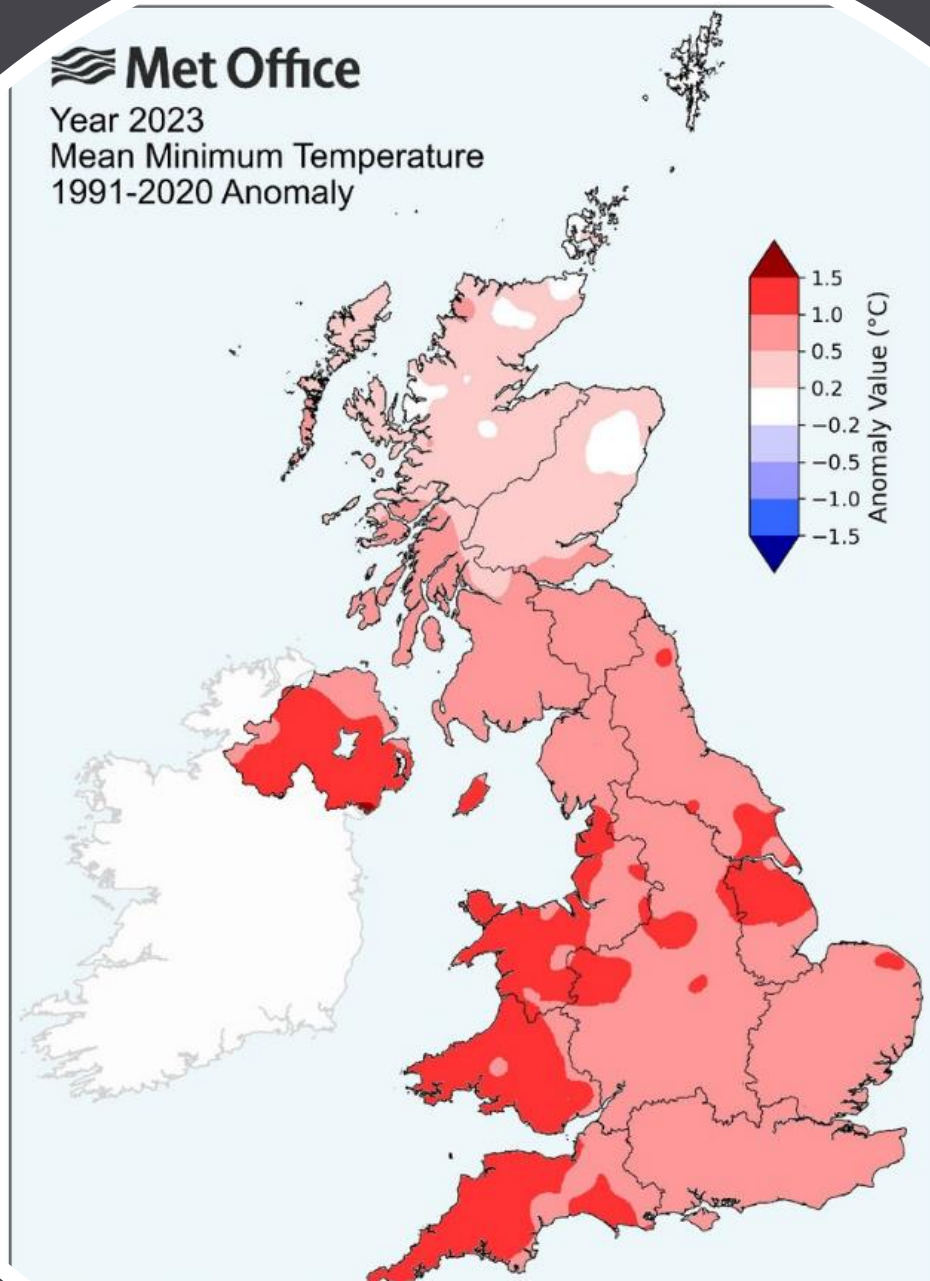


**BYDV = persistent**

- The ability and efficiency to vector BYDV strains varies between aphid species...and populations



# Challenges







# Yellow dwarf virus strains

Genus	Species	Strain	Known aphid vectors*
Luteovirus	BYDV	<b>PAV</b>	<i>R. padi</i> , <i>S. avenae</i> , <i>Metapolophium dirhodum</i> , <i>Schizaphis graminum</i> , <i>Sitobion fragariae</i>
		<b>MAV</b>	<i>S. avenae</i> , <i>M. dirhodum</i> , <i>R. padi</i> , <i>S. fragariae</i>
		<b>PAS</b>	<i>Rhopalosiphum maidis</i>
		<b>GAV</b>	<i>Scizaphis graminum</i> , <i>S. avenae</i>
Polerovirus	CYDV	<b>RPV</b>	<i>R. padi</i> , <i>S. graminum</i>



***Rhopalosiphum padi***  
bird cherry-oat aphid



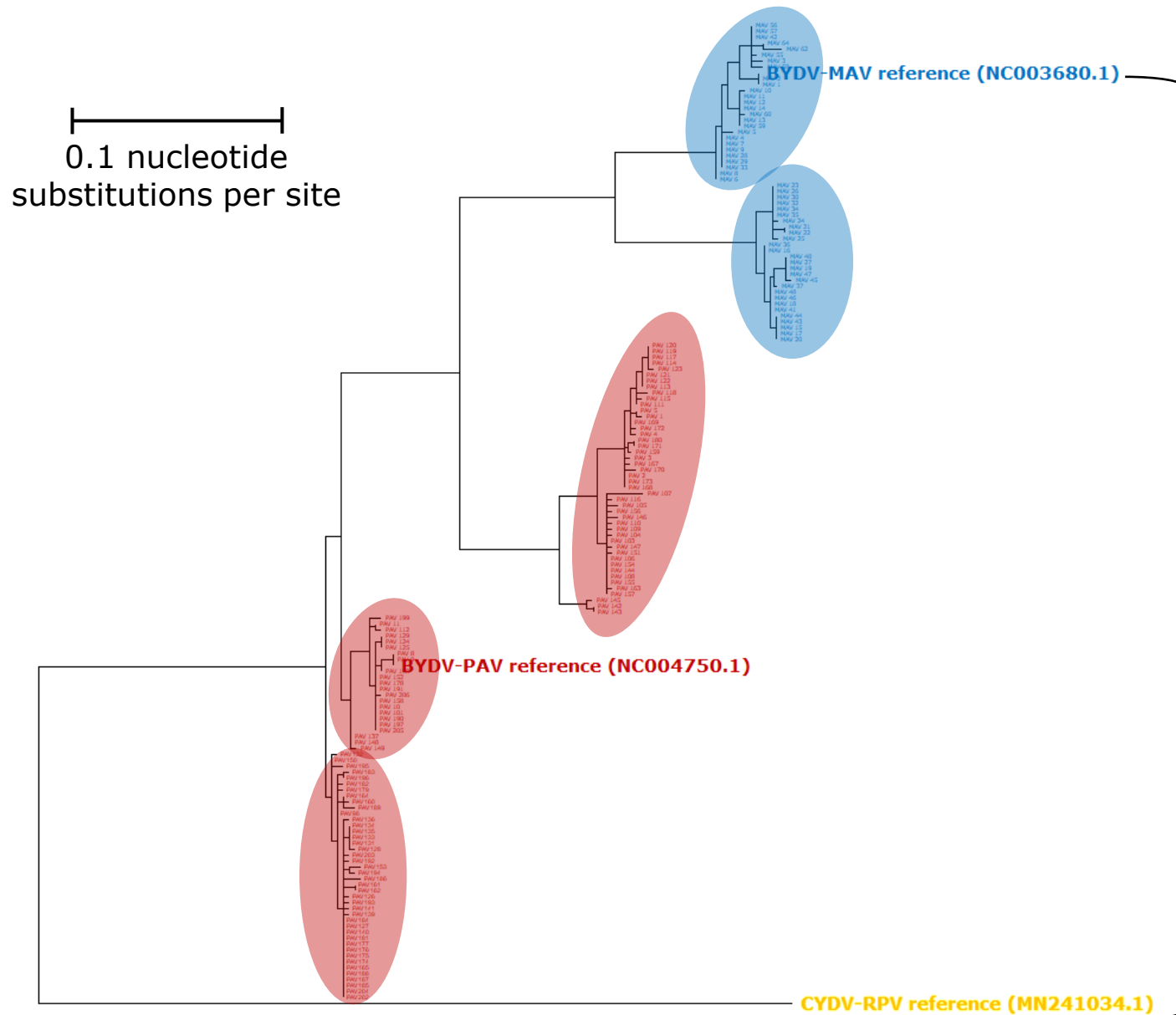
***Sitobion avenae***  
English grain aphid

\*Aradottir & Crespo-Herrera (2021). *Current Opinion in Insect Science.*, 45, 59-68





# BYDV strain variation



- Strains **BYDV-MAV** and **BYDV-PAV** considered most impactful for UK

- Global **MAV** and **PAV** diversity, based on subset of NCBI data

- Substrain variation suggested on global scale

- **Molecular BYDV diagnostics often based on limited data**

BYDV-MAV sequences from NCBI, n = 53

BYDV-PAV sequences from NCBI, n = 110

NCBI, National Centre for Biotechnology Information





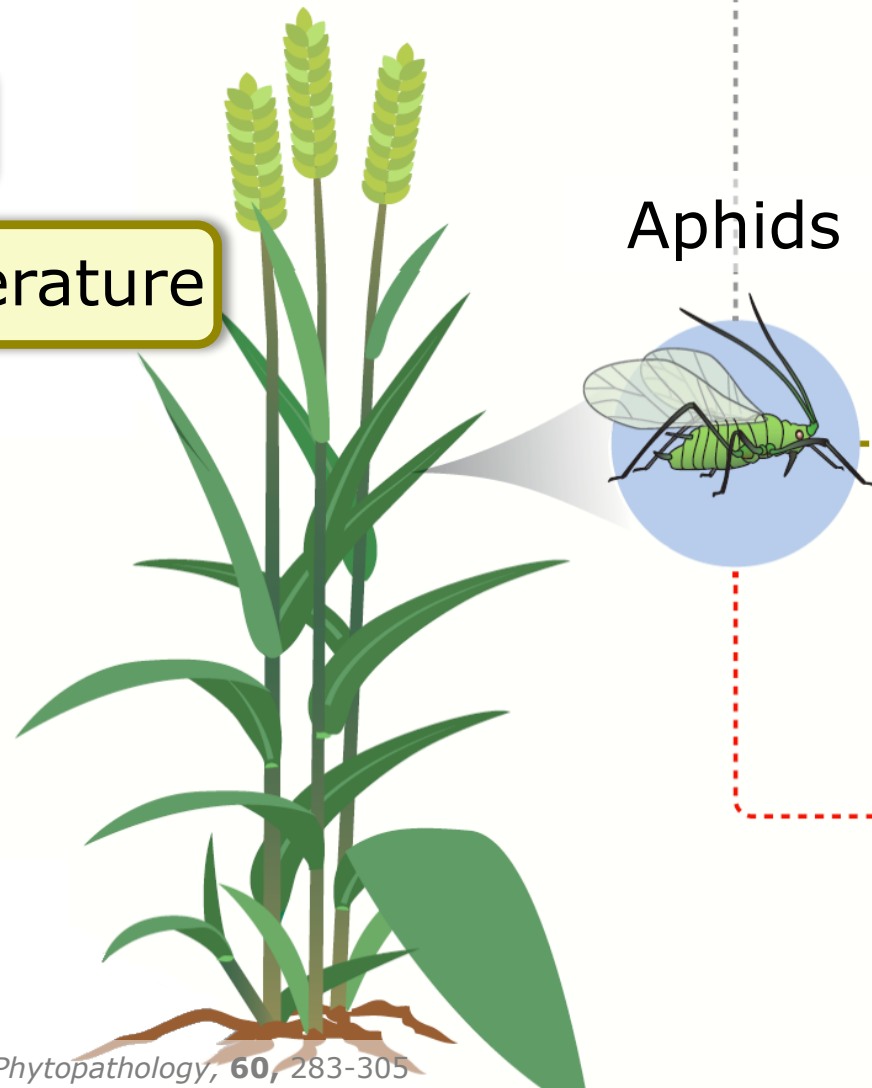
Future impacts upon aphid-wheat-virus pathosystems from:

- Elevated CO<sub>2</sub>

- Elevated temperature

- Drought

Climate change to exacerbate need for BYDV management



↓ (or no) Δ to population size  
(yellow dwarf viruses)  
eCO<sub>2</sub> × eTemperature × YDV  
show no change to aphid fitness

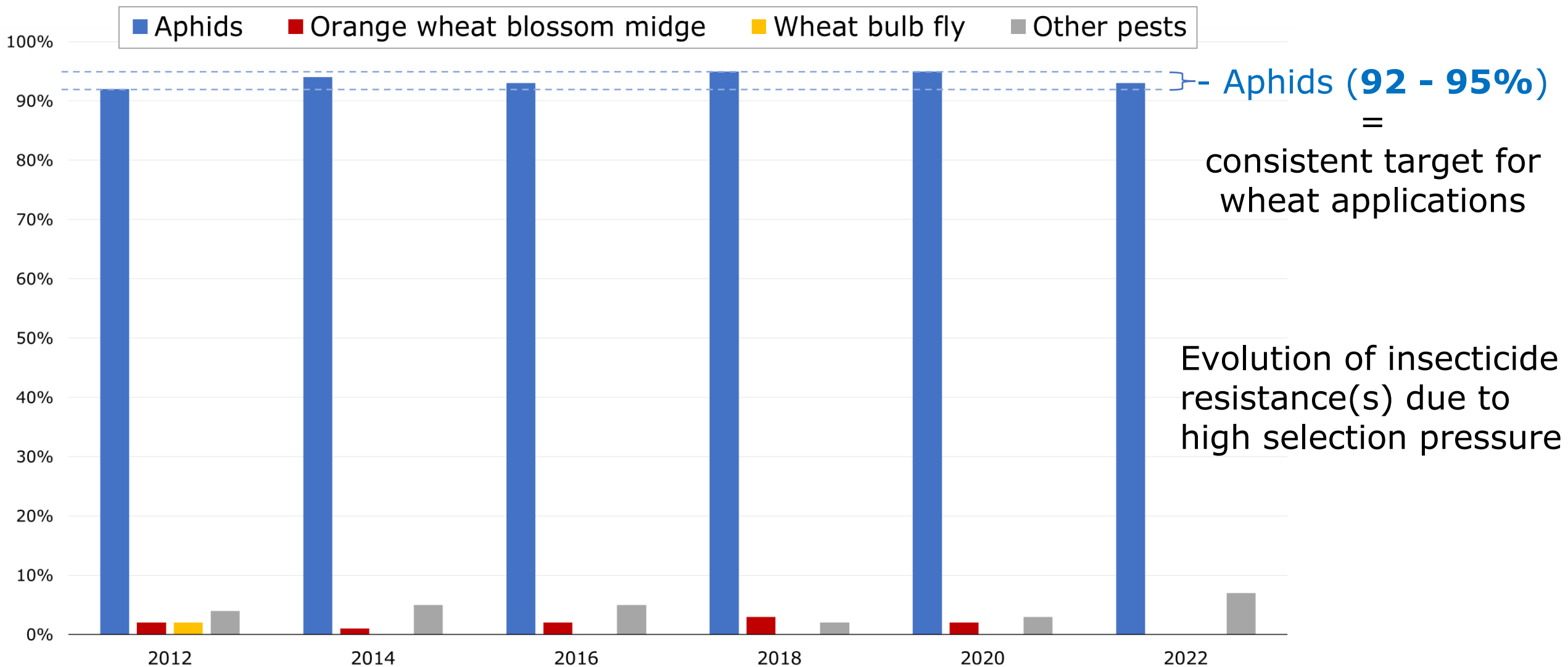
↑ Food consumption  
↑ Metabolic rate  
↑ Abundance  
↑ Alates  
↑ Flight activity/movement  
↑ Interplant movement by apterous aphids  
↑ Dropping rate from plants  
↓ Generation time

↑ Movement  
↑ YDV transmission  
↑ Alates  
Drought × eTemperature  
show increased aphid landing on plants



# Aphids = key insecticide target

Insect targets, where stated, for foliar insecticide application to wheat in the UK\*:



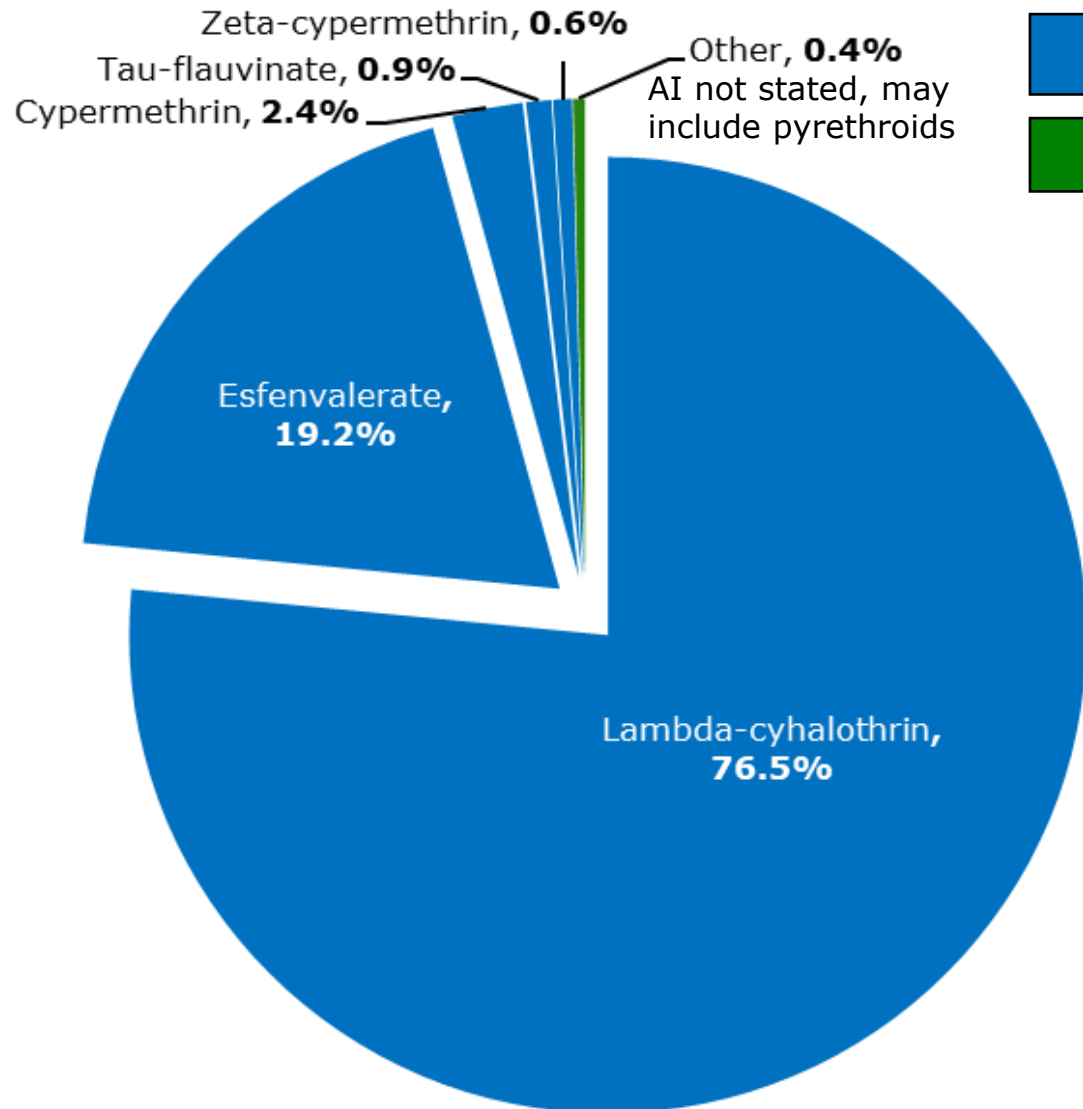
\*UK Pesticide Usage Survey Reports: 250, 263, 271, 284, 295 & 309





# Aphids = key insecticide target

Main foliar insecticide AI formulations (most recent data available, 2022 survey<sup>1</sup>):



- - Pyrethroid-based active ingredient (AI)
- - Other



***Rhopalosiphum padi*** (bird cherry-oat aphid) & ***Sitobion avenae*** (English grain aphid)

= impactful UK cereal aphids/BYDV vectors

- Moderate pyrethroid resistance widespread across UK *S.avenae*
- Evidence of pyrethroid tolerance in *R.padi* in Ireland<sup>2</sup>

<sup>1</sup>Ridley et al., (2023) UK Pesticide Usage Survey Report 309  
<sup>2</sup>Walsh et al., (2020). *Outlooks on Pest Management*, 31, 5-9

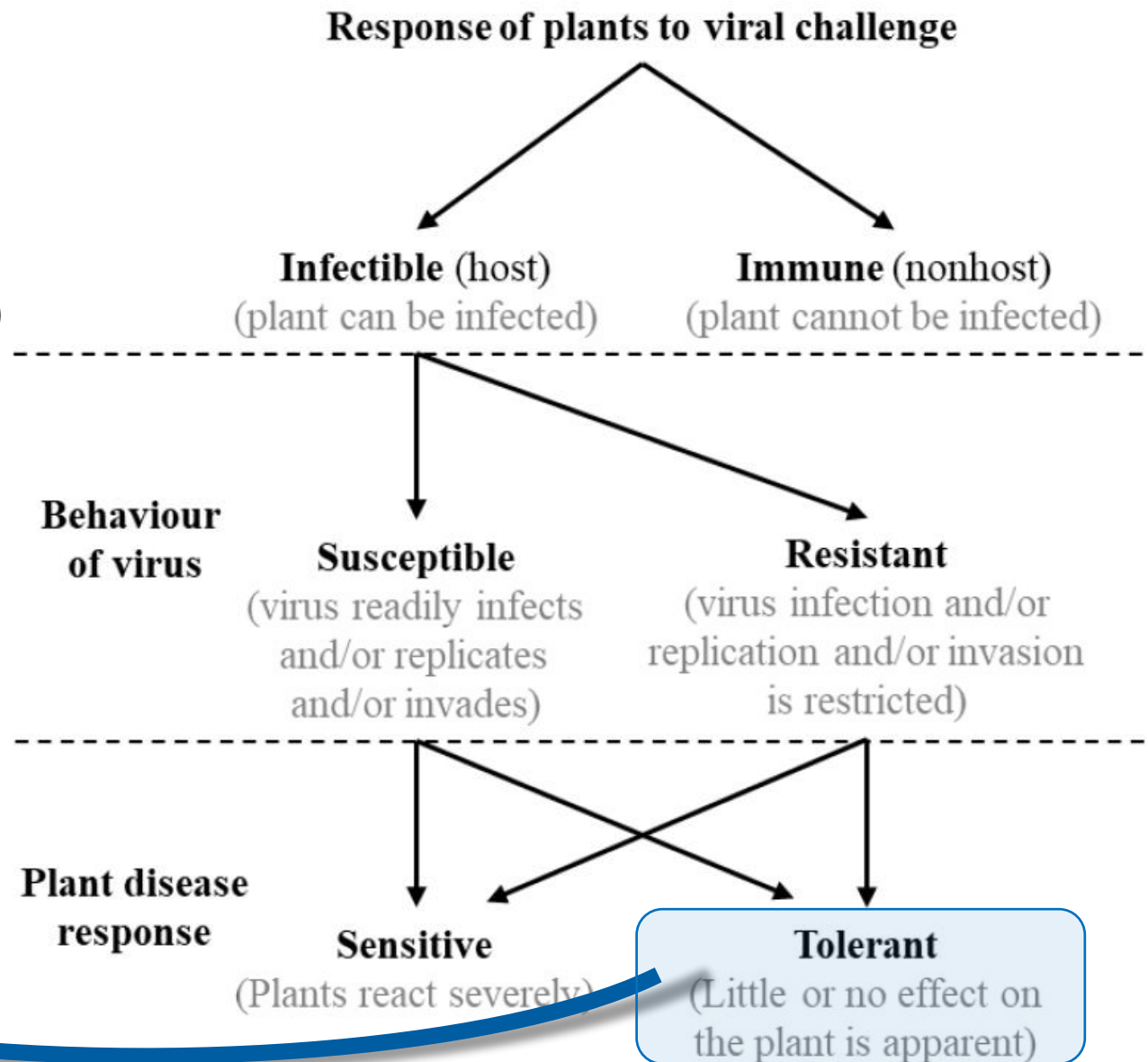


# BYDV resistance?

Effective deployment of genetic resistances depends upon knowledge of:

- Inoculum (BYDV strains, distribution etc.)
- Resistance mode-of-action
- Resistance spectra (all strains vs one?)
- Resistance or tolerance...

“Can tolerant varieties harbour BYDV and/or influence BYDV prevalence?”







Genetic sources of resistance are relatively scarce...

## Barley:

- Four genes linked to tolerance/resistance ("*Ryd1 - Ryd4Hb*", & 3 QTLs)
- *Ryd2* = most heavily deployed<sup>1</sup>
- Reduces viral titre in young plants, some doubt for older plants
- Virus can still spread systemically

## Wheat:

- Similarly, four BYDV-resistance linked genes ("*Bdv1 - Bdv4*")<sup>2</sup>
- *Bdv2* originated from *Thinopyrum intermedium*
- Winter wheat developed by:



<sup>1</sup>Jarošová, et al., (2016). *Field Crops Research*, 198, 200-214.

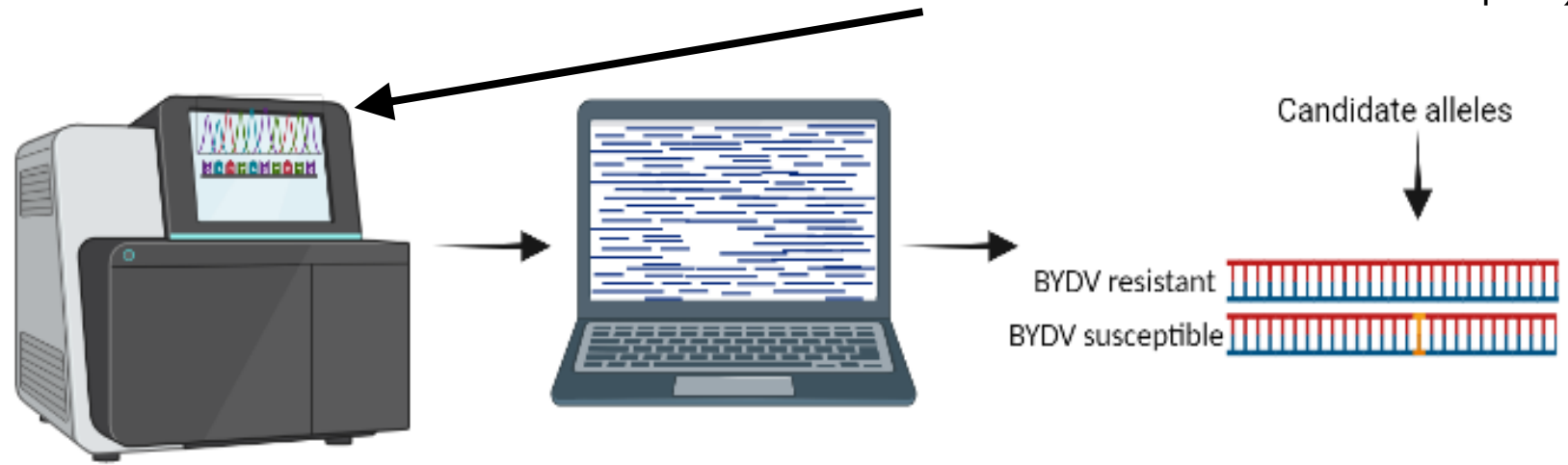
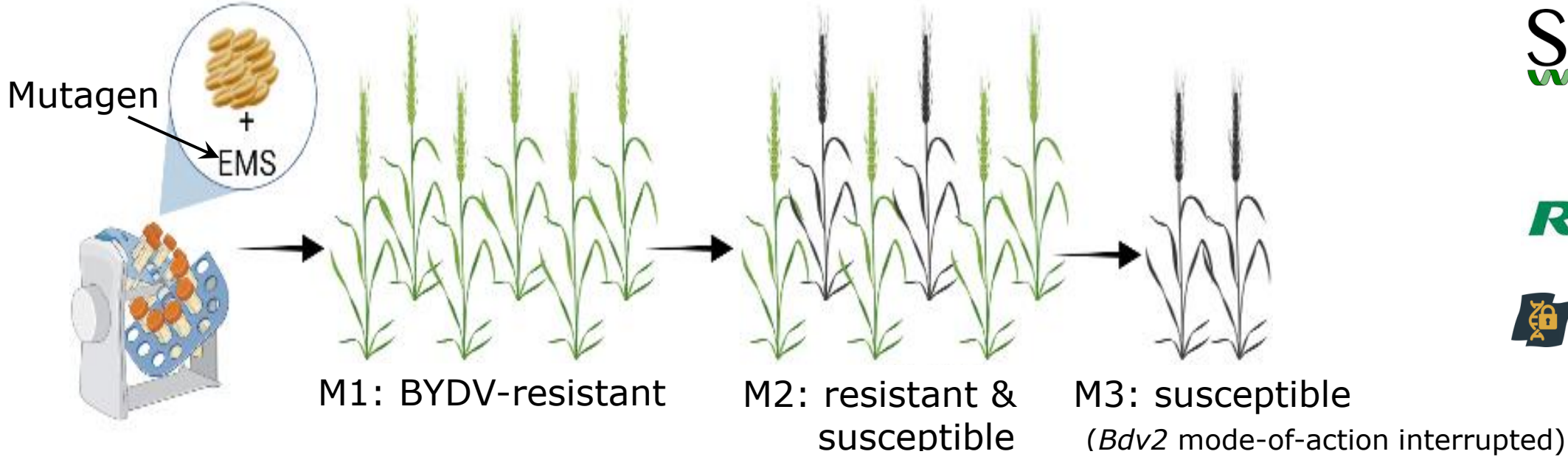
<sup>2</sup>Aradottir & Crespo-Herrera (2016). *Current Opinion in Insect Science*, 45, 59-68.



# BYDV resistance?



Exploring RGT cv. Wolverine BYDV resistance/tolerance



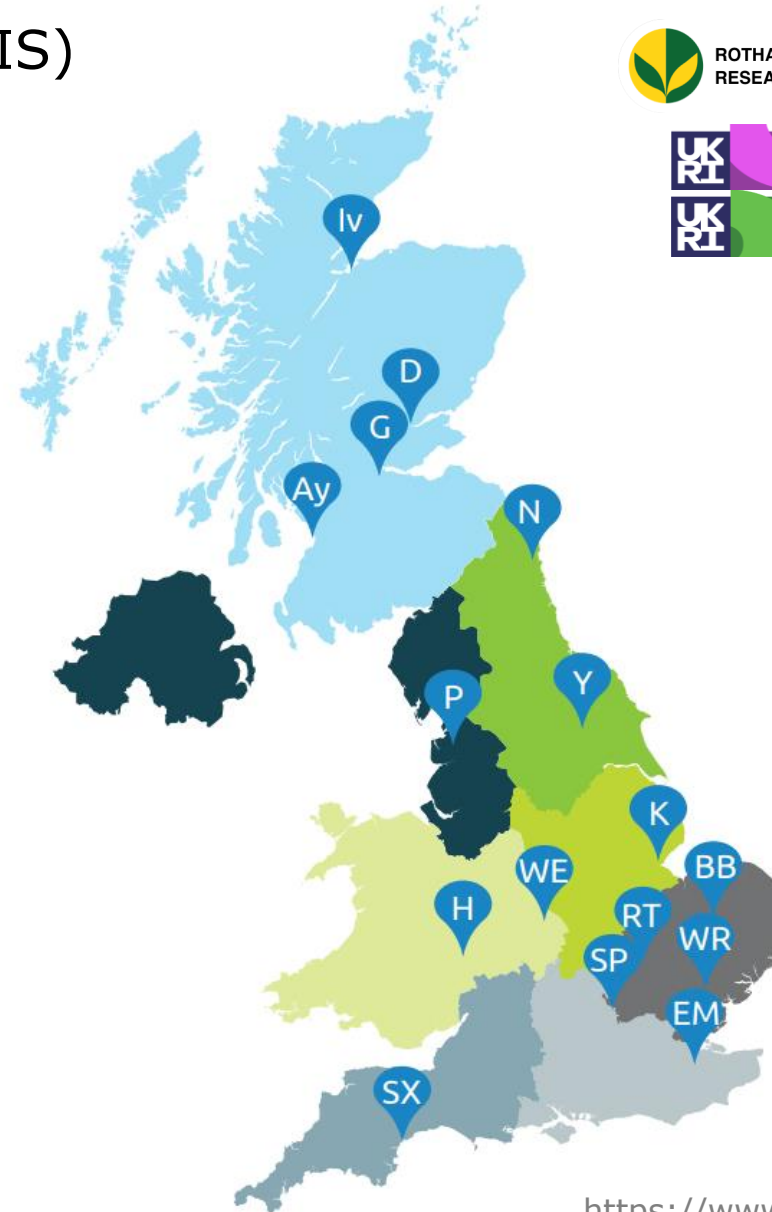




- Nationwide Rothamsted insect survey (RIS)
- Started in 1964

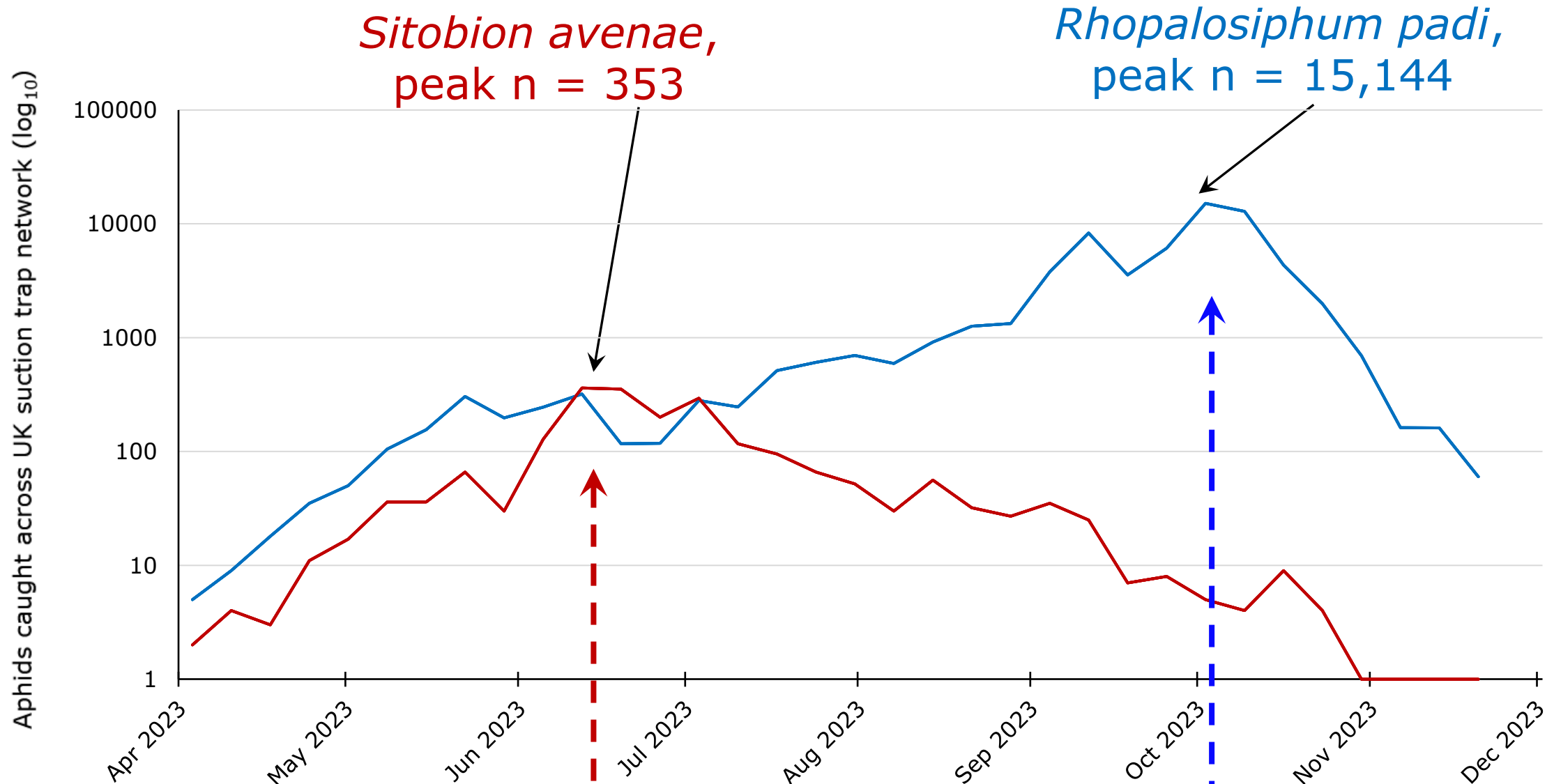


Suction traps (12.2m)





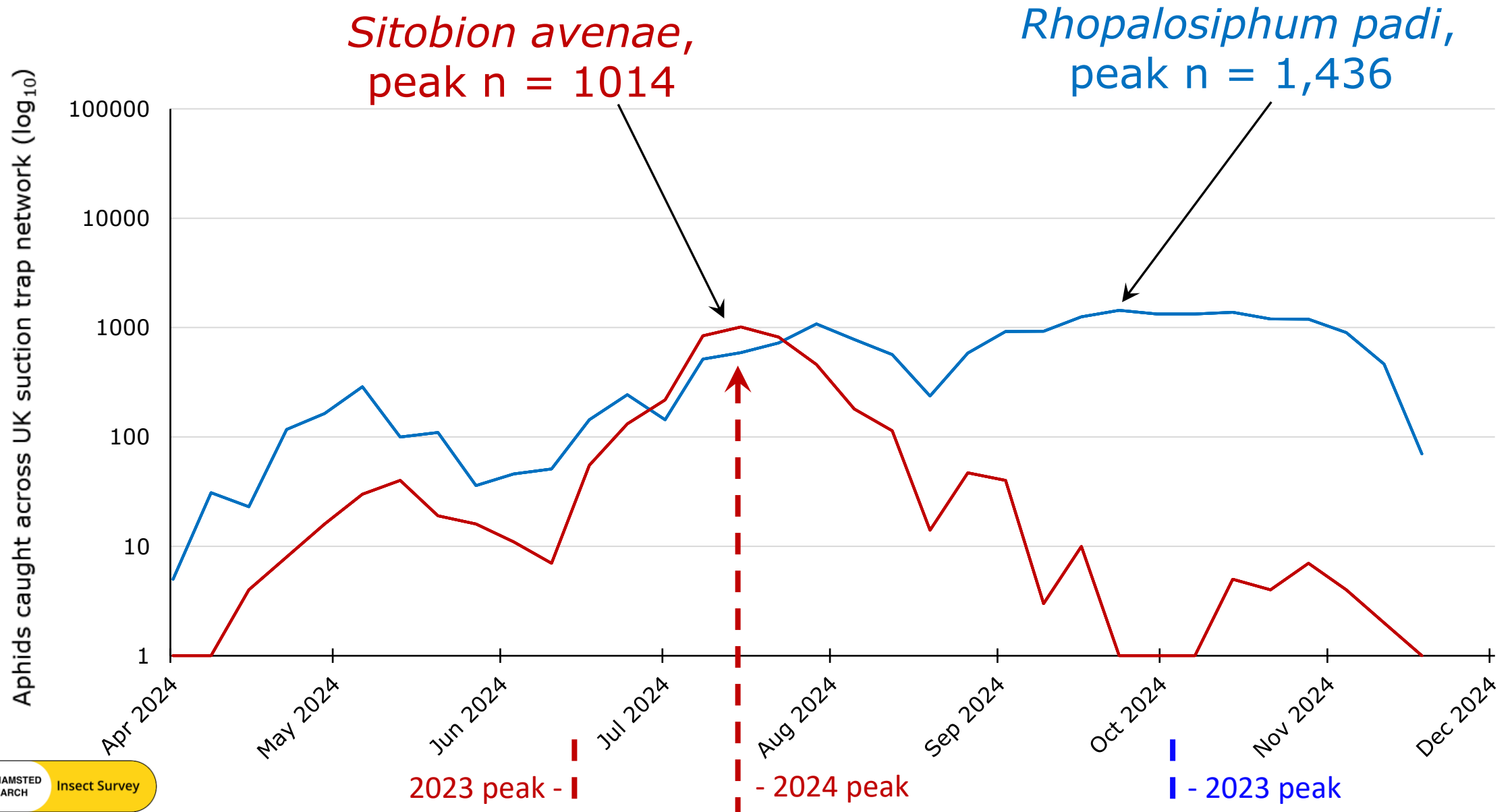
# UK aphid pressure (2023)







# UK aphid pressure (2024)





# Routine BYDV detection



\*

Year	<i>R. padi</i> carrying BYDV/total tested (%)	<i>S. avenae</i> carrying BYDV/total tested (%)
2020	329/1509 ( <b>22%</b> )	77/252 ( <b>31%</b> )
2021	231/1342 ( <b>17%</b> )	34/224 ( <b>15%</b> )
2022	128/980 ( <b>13%</b> )	20/108 ( <b>19%</b> )
2023	147/920 ( <b>16%</b> )	29/88 ( <b>33%</b> )
2024	153/990 ( <b>15%</b> )	23/88 ( <b>26%</b> )

- BYDV consistently detected UK-wide
- Highest incidence = 46% *S. avenae*, late July 2020
- lowest incidence = 5% *S. avenae*, early Aug 2021



**Subset of aphids taken forward for informative CP sequencing**





# Strain variation across the UK



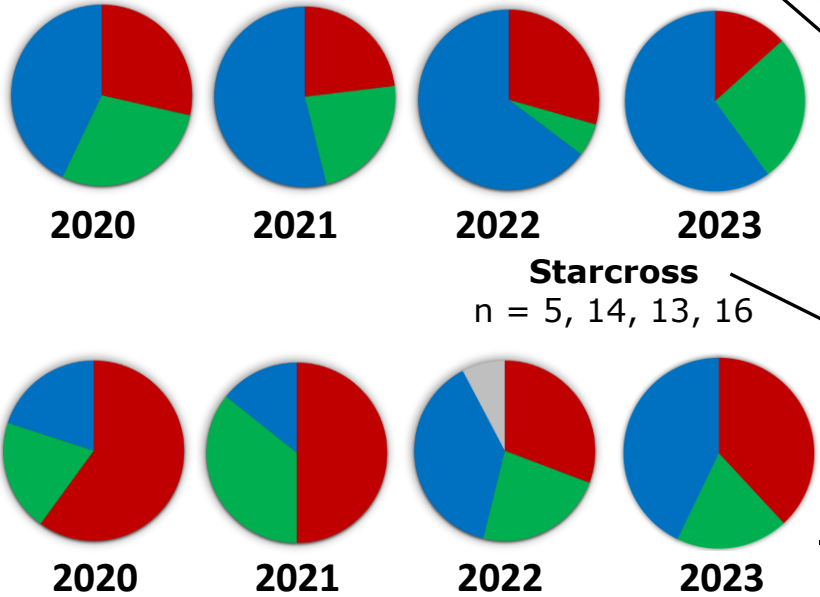
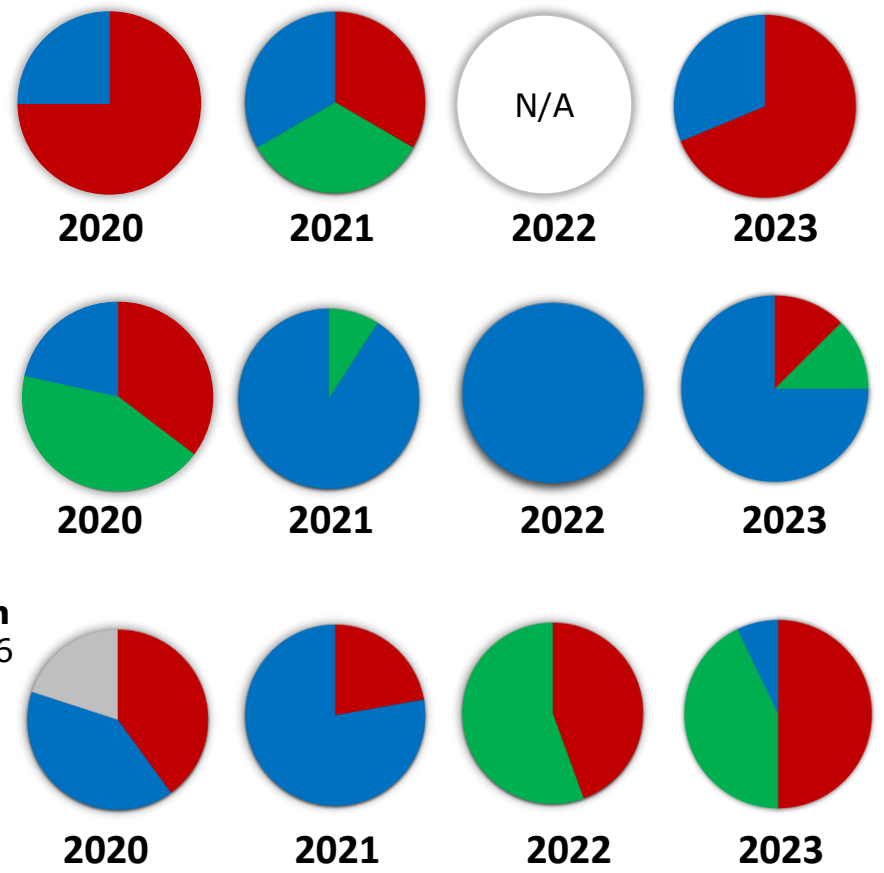
**Hereford**  
n = 7, 13, 21, 15

**Edinburgh**  
n = 4, 3, 0, 16

**York**  
n = 14, 11, 12, 16

**Brooms Barn**  
n = 5, 9, 9, 16

**Starcross**  
n = 5, 14, 13, 16



- BYDV strains not strictly conserved geographically
- Implications for resistance screening/germplasm deployment



# Yellow dwarf virus strains

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***Rhopalosiphum padi***  
bird cherry-oat aphid



***Sitobion avenae***  
English grain aphid

**Previously unreported** and confirmed via RIS sampling/testing

\*Aradottir & Crespo-Herrera (2021). *Current Opinion in Insect Science.*, 45, 59-68

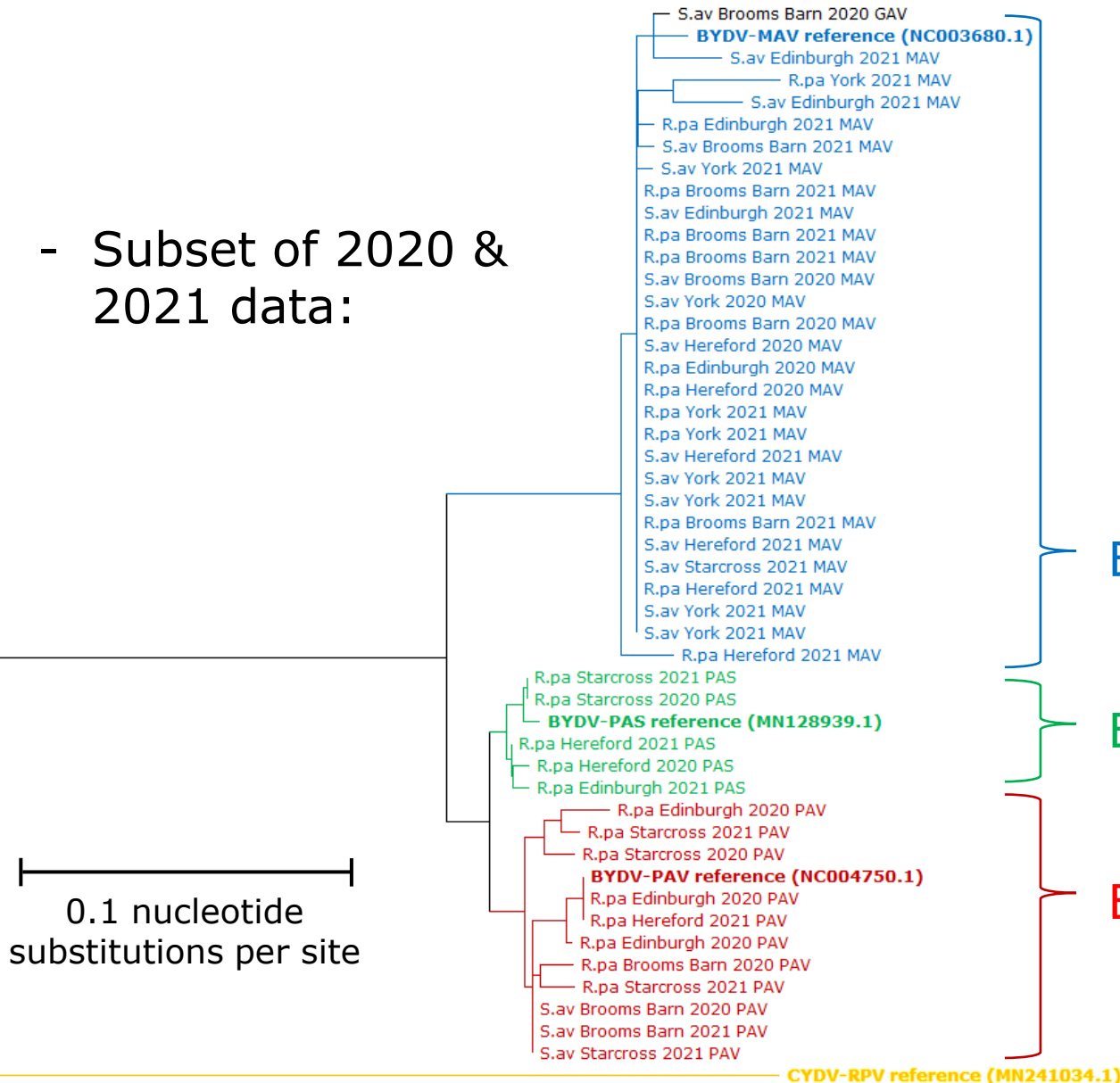




# Better diagnostics

- Subset of 2020 & 2021 data:

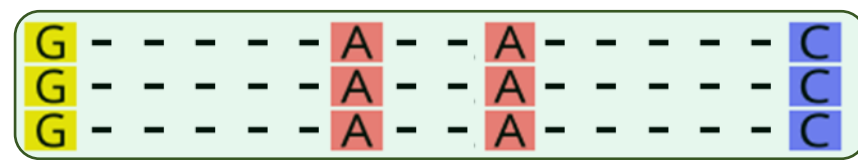
- Sequencing information used to identify conserved, UK strain-associated polymorphisms



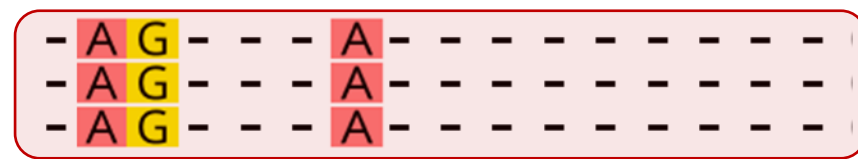
BYDV-MAV



BYDV-PAS



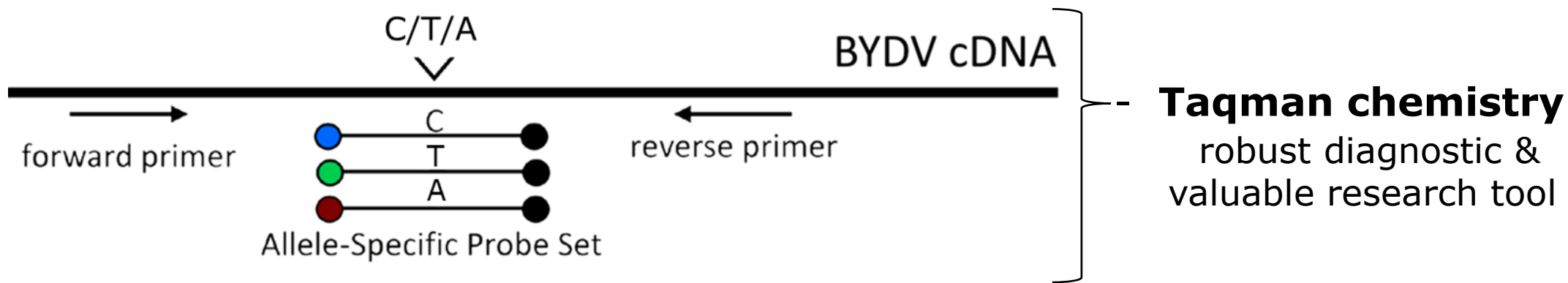
BYDV-PAV



Sequencing data anonymised but representative



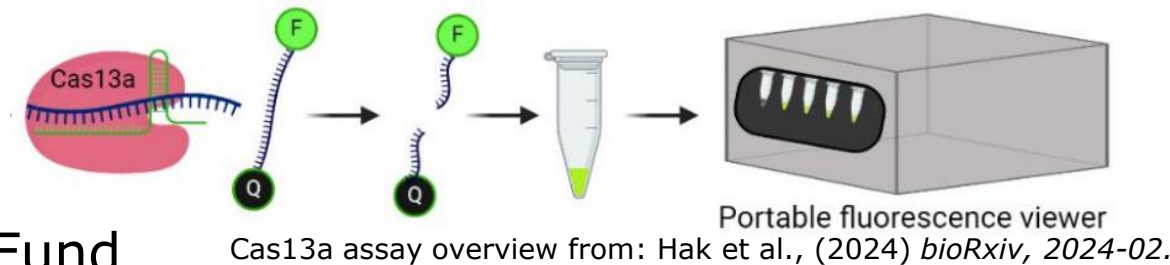
# Better diagnostics



- Strain-specific Taqman, KASP, PCR & qPCR assays = **established**
- RT-LAMP & CRISPR-Cas13a assay = in progress, for **"in-field" testing**



BSPP Small Project Fund



- Can now process up to 24 discrete molecular assays vs 192 samples
- **4,608** qPCRs in ~2h
- **Massive potential for higher-throughput exploration**



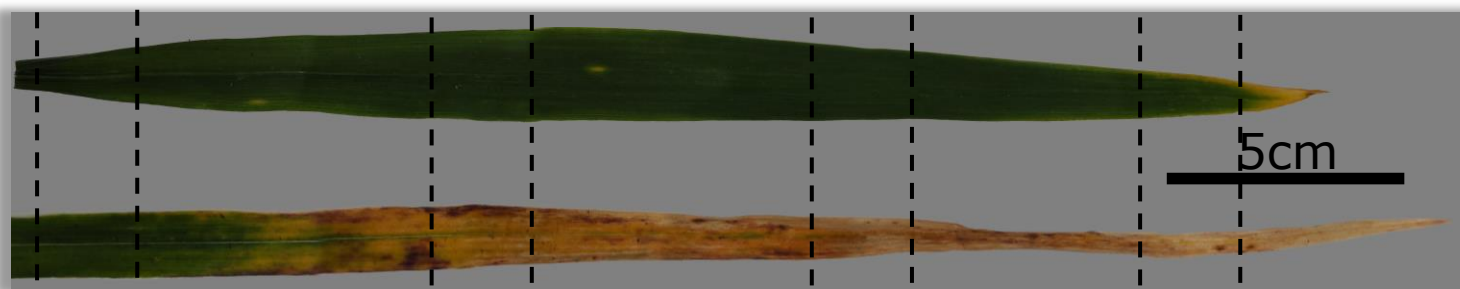
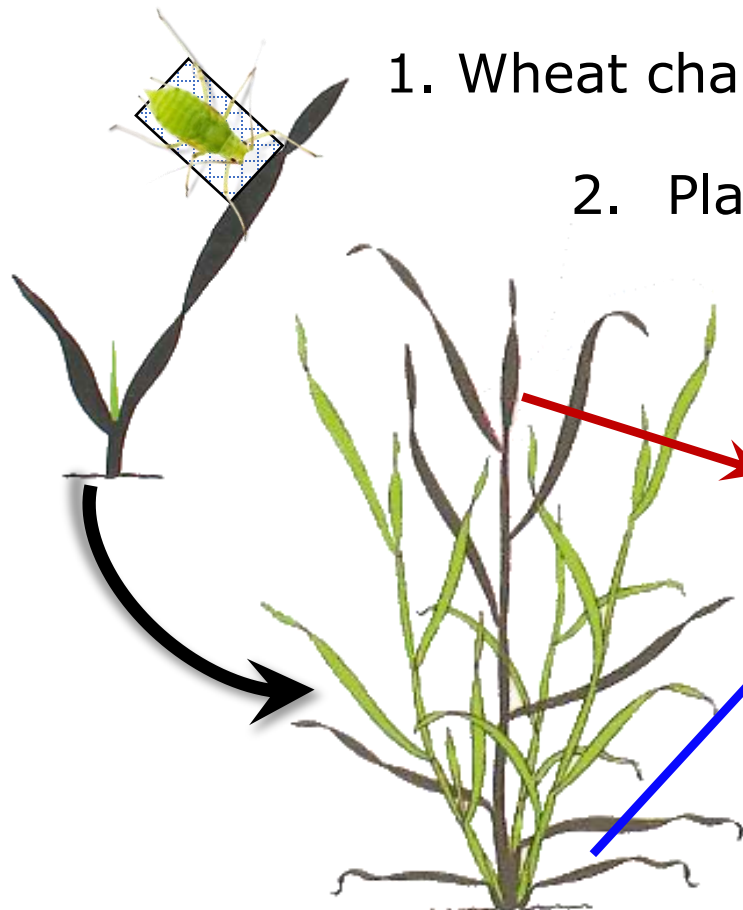


## Does sampling location impact BYDV detection?

1. Wheat challenged with BYDV

2. Plant maintained until flag leaf developed

3. **Flag leaf** and **BYDV-challenged leaf** sampled



BYDV qPCR Ct values	Sampling Locations			
	Location 1	Location 2	Location 3	Location 4
Column-based RNA extraction	20.8	n/a	21.0	22.5
Crude RNA extraction	33.6	33.3	31.4	37.3
	36.4	35.2	36.7	30.9



- Challenges do exist
  - Shifting BYDV strains, aphid populations, climate...
  - Robust resistance deployment and ensuring current resistances last requires ongoing characterisation
- Opportunities
  - The resources exist for modelling changes to this important pathosystems
  - New tools (e.g. infectious clones, higher throughput multiplexed assays) are on the horizon

Aphid vectors are one element, any and all potential wheat samples with BYDV symptoms would be gratefully received





- For your attention and the UKCPVS Stakeholders event organisers
- Rothamsted Wheat Pathogenomics Team (PI, Kim Hammond-Kosack)
  - Wheat-Insect-Virus Interactions Subgroup (PhD student, Izayana Sandoval-Carvajal)
- Continued assistance from the wider wheat and cereals community
  - DSW & WGIN and associated funding (UKRI & Defra, respectively)



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Biotechnology and  
Biological Sciences  
Research Council



BSPP Small Project Fund  
- new 'in-field' assay development



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Insect Survey