Emerging fungal threats to UK brassica crop health ROTHAMSTED

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RESEARCH

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Focus on two major UK brassica diseases that can co-occur





Identifying and managing emerging pathogen threats: pathogen population biology research

- Researching the biology and genetics of pathogen populations can help identify, prevent and if needed manage emerging pathogen threats that may cause substantial economic and/or environmental damage
- We demonstrate this concept using two case studies based on major UK brassica crop diseases
 - Case study #1: Light leaf spot
 - Case study #2: Phoma leaf spot / stem canker (blackleg)
- These principles can be applied to many other pathogens; e.g. humans, animals, forests etc.

Case study #1 : Light leaf spot (Pyrenopeziza brassicae)



LLS long been reported throughout Europe and Oceania (>80 years)

Major UK disease of OSR + vegetables



An emerging international threat

- North America: Oregon + Washington States
- Absent in disease surveys until 2013
- First detection 2014
- Since: rapid, invasive spread across western Oregon + Washington. Speciality brassicas + OSR



Carmody SM*, King KM*, Ocamb CM, West JS, du Tout LJ, 2020. Plant Pathology 69, 518-537.

Population biology studies reveal two evolutionary lineages

of Pyrenopeziza brassicae



Based on alignments of three housekeeping genes (ITS, beta tubulin and alpha elongation factor)

- P. brassicae lineage 1 in Europe and Oceania long resident
- P. brassicae lineage 2 in North America emerging invasive

The lineages cause two distinct diseases on brassicas

Lineage 1 Light Leaf Spot (LLS)



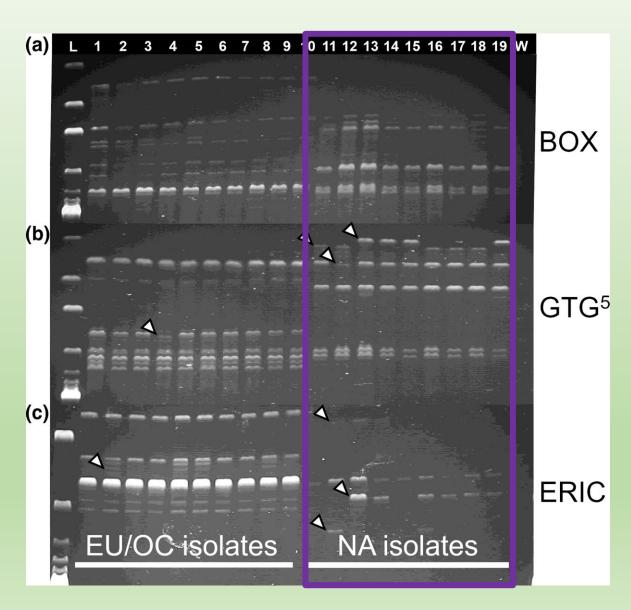
- LLS in Europe + Oceania
- White conidiomata may be visible
- Pale green / bleached lesions that extend slowly, later becoming necrotic
- Individual lesions may merge causing complete leaf necrosis



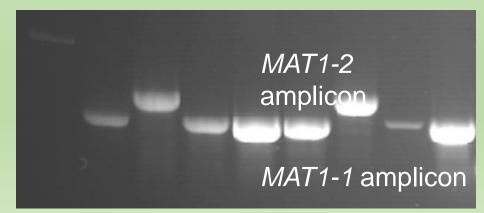
Lineage 2 Chlorotic Leaf Spot (CLS)

- <u>CLS in North America</u>
- No white conidiomata
- Bright yellow chlorotic spots that expand rapidly, later remain chlorotic
- Individual lesions may merge causing complete leaf chlorosis followed by early senescence

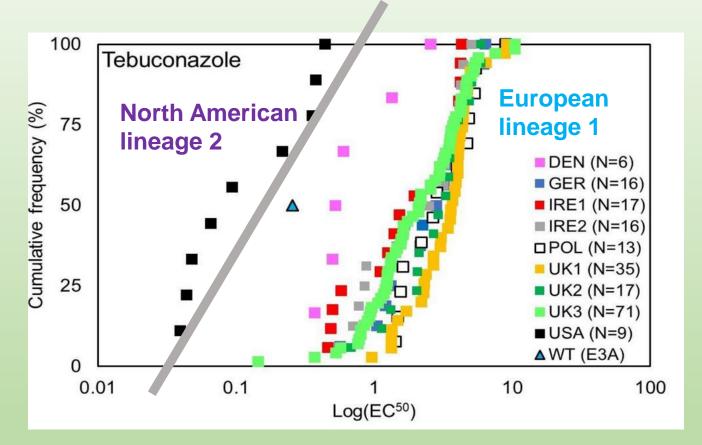
Population genetics: North American lineage 2



- Neutral markers: genetically diverse
 (compared to European lineage 1)
- Both mating types (*MAT1-1* and *MAT1-2* present) required for sexual reproduction are present, and in a 1:1 ratio!
- A sexually reproducing population?
- Evolutionary potential + disease management implications....



Differences in fungicide sensitivity between the lineages...



- Fungicide sensitivity testing has shown:
- Differences in azole + MBC sensitivity between European lineage 1 and North American lineage 2
- Need to monitor North American lineage 2 population: early detection of resistance
- Resistance management strategies

New European isolates deposited into OREGIN culture collection. A genetic resource for future studies

King KM, Bucor DE, Ritchie F, Hawkins NJ, Kakzmarek AM, Duan Y, Kildea S, West JS, Fraaije BA, 2021. Plant Pathology 70, 2086-2103.

P. brassicae lineage-specific PCR diagnostics developed



Implications of the two P. brassicae lineages...

- Lineages vs. species...
- UK Plant Health authorities alerted
- Geographic distribution of the lineages: nationally + internationally
- Origins of *P. brassicae*: other lineages?; sources of host resistance?
- Brassica crop health implications requires further research:
 - But, rapid invasive spread of lineage 2 poses threat to US Brassica growers (and Canadian OSR crops)
 - Avoid introductions of the lineages between continents: molecular diagnostics now available
 - Implications of lineage introductions:
 - Individually: distinct plant health risks
 - Physical contact between lineages: interlineage recombination / hybridization

Case study # 2: Phoma leaf spot

Plenodomus lingam = *Leptosphaeria maculans* Two genetic subclades: 'brassicae', 'lepidii'



- Evidence that the species and subclades pose distinct plant health risks: require different management strategies?
 In Europe, only *P lingam* ('brassicae') and
- In Europe, only *P. lingam* ('brassicae') and *P. biglobosus* ('brassicae') reported so far, but no recent surveys...

Plenodomus biglobosus = Leptosphaeria biglobosa Seven genetic subclades (so far!): 'americensis', 'australensis', 'brassicae', 'canadensis', 'erysimii', 'occiaustralensis' and 'thlaspii'

Phoma on wasabi (Eutrema japonicum) crops in the UK?





- Three UK sites surveyed in 2021 Northern Ireland, Southern England & West Midlands
- Lesions on leaves and also some petioles and stems (along vascular bundles) suggesting systemic infection
- Symptoms consistent with Phoma leaf spot. Is this correct? If so, which *Plenodomus* species and genetic subclades are present?

King & West (2021) European Journal of Plant Pathology. In press.

Fungal isolation from phoma-like diseased wasabi leaves

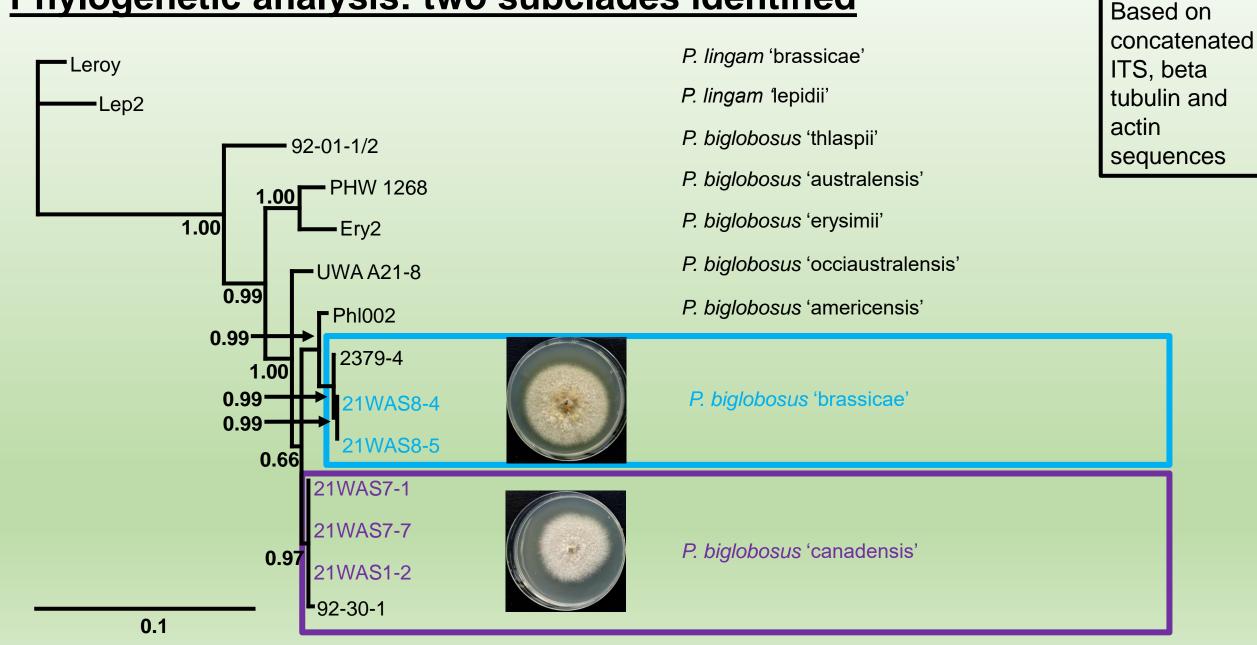
Two distinct isolate phenotypes identified:

Non



Bright yellow pigment

Phylogenetic analysis: two subclades identified

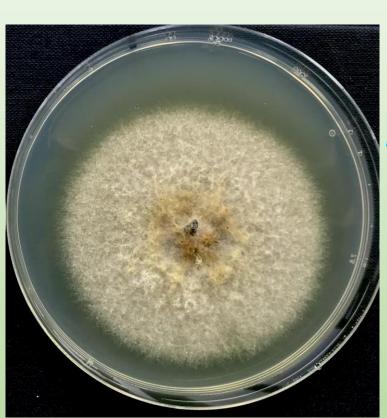


Sequence analysis confirmed genetic identities of isolates

Plenodomus biglobosus subclade 'canadensis'



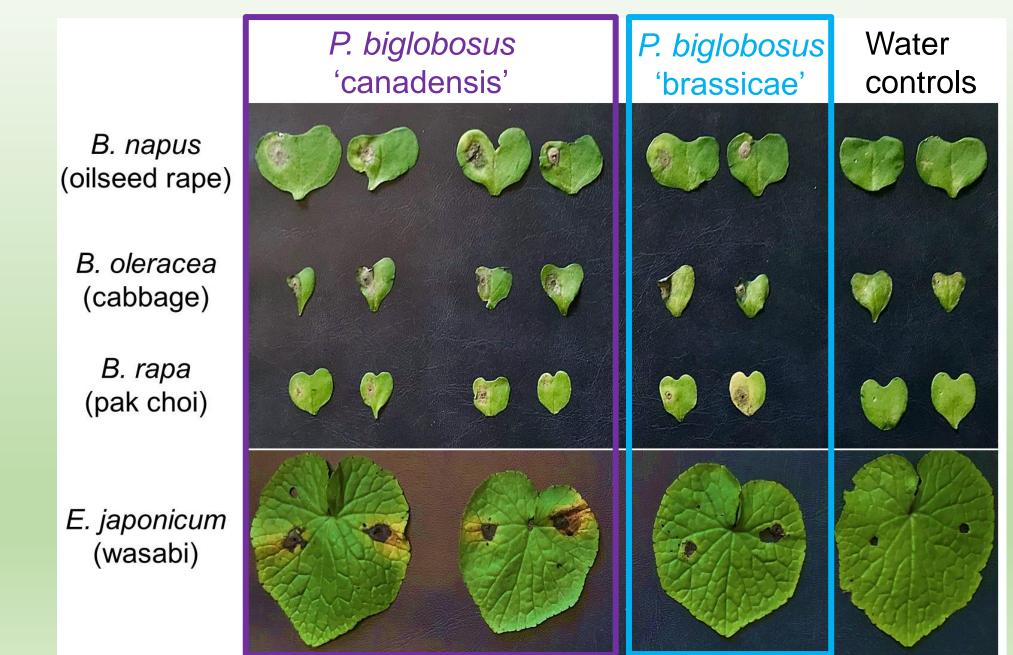
- First report of 'canadensis' subclade in Europe and on wasabi host
- Southern England & Northern
 Ireland



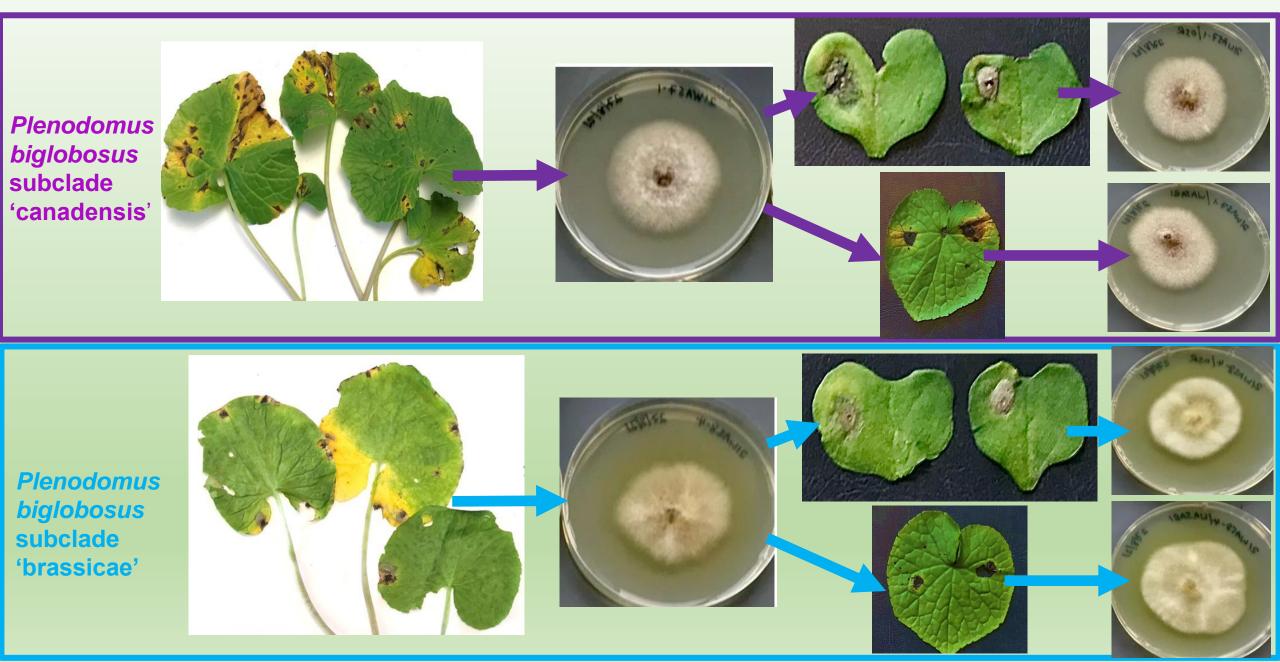
Plenodomus biglobosus subclade 'brassicae'

- First report of 'brassicae' subclade on wasabi host
- West Midlands

Pathogenicity testing on multiple hosts



Confirmation of Koch's postulates



Implications of findings

- Distribution of *P. biglobosus* subclades canadensis vs brassicae on OSR + veg
- Other subclades already present???
- Highlights importance of molecular diagnostics
- Existing PCR-based diagnostics and colony morphology may not accurately detect/discriminate
- Other brassica crop health implications of the subclades:
 - Distinct plant health risks
 - Different responses to management strategies e.g. fungicides?
 - Sexual or asexual recombination?
 - Hybridisation?

Concluding remark

- As demonstrated for two major OSR diseases:
- Pathogen population biology research, particularly on major current pathogens, can:
 - Advance knowledge
 - Identify future likely plant health risks (horizon scanning)
 - Improved molecular diagnostics (detection of pathogen genotypes):
 - Reduce the risk of accidental introduction events by improved targeting of finite resources
 - Improved disease management strategies

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