

DEVELOPING AN EARLY WARNING SYSTEM FOR WHEAT RUST

Three rust species infect wheat and are distributed globally:

Yellow (stripe) rust (Puccinia striiformis f.sp. tritici) **Leaf (brown) rust** (*Puccinia triticina*) **Stem (black) rust** *Puccinia graminis* f.sp. *tritic*)

These pathogens have a worldwide prevalence and potential for long distance spread. Invasions into new areas pose a serious threat to food security regionally and globally. NIAB is a partner in the European-wide RustWatch project, which is establishing a stakeholder driven early-warning system to improve preparedness and resilience to emerging rust diseases on wheat. Information gathered as part of the project includes:

Understanding pathogen biology and drivers

Figures 1 and 2 show the frequency distribution of yellow rust genetic groups. The Red Group [PstS10, Warrior(-)] has spread across Europe since 2013 and is the dominant genetic group across Europe.

Figure 1. Distribution of yellow rust genetic groups 2013-21



Disease prevention by host resistance

Microscopy is used to distinguish initial adult plant resistance mechanisms involved in the pathogen-host interaction in around 240 wheat varieties.

Figure 3. Yellow rust elongating and infecting host cells

Figure 4. The formation of a yellow pustule



Figure 2. Distribution of yellow rust genetic groups 2021



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Stakeholder networks, shared facilities and case studies

Variety blends have been shown to reduce risk from yellow rust, with 22 trials over three years in Europe. 25% of the wheat area in Denmark was sown with blends in 2022. The next step will be to investigate whether pathogen populations evolve to be more complex on blends vs monoculture.

> Figure 5. 2021 yellow rust scores (% infection) – NIAB, Cambridge

Variety	Untreated	Full fungicide programme	Reduced fungicide programme
Crusoe	0	0	0
KWS Zyatt	35	0	1

 Skyfall
 55
 0.1
 3

 3-way blend
 15
 0.1
 3



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