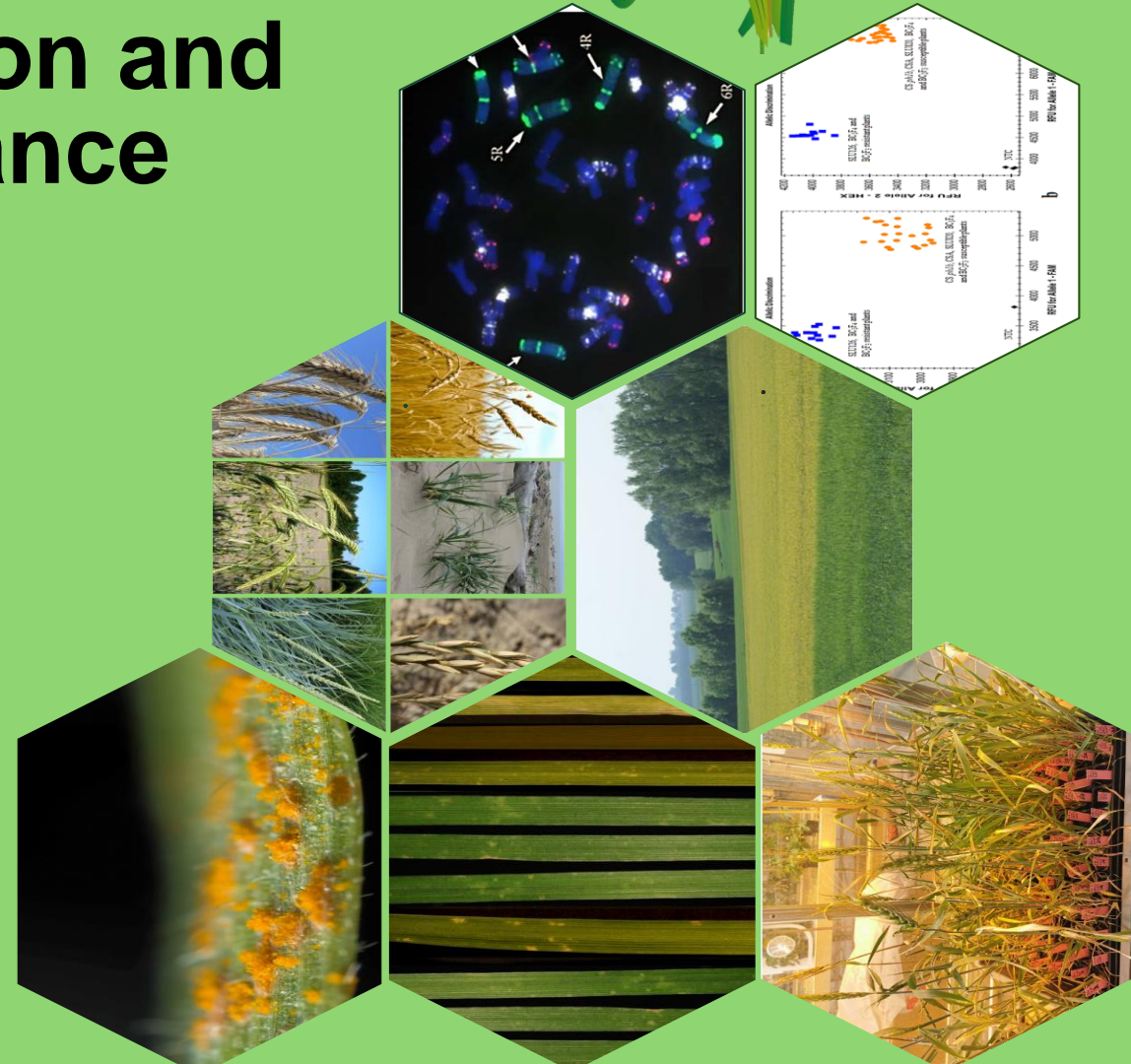


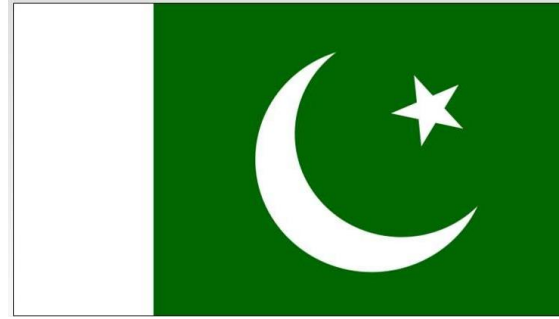
Identification, characterization and combination of novel resistance genes against rust in wheat

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PAKISTAN



M.Sc (Botany)
M.Phil (Plants Genetics and genomics)



SWEDISH UNIVERSITY
OF AGRICULTURAL
SCIENCES

P.hD (Plant Breeding)

Outline



1

Background

2

Aims and objectives

3

Potential of wheat-alien Introgression lines

4

Rye chromosome against Stripe rust (*Yr*)

5

Combination of *Sr* and *Yr* resistance genes

6

Conclusions

7

Future Directions

8

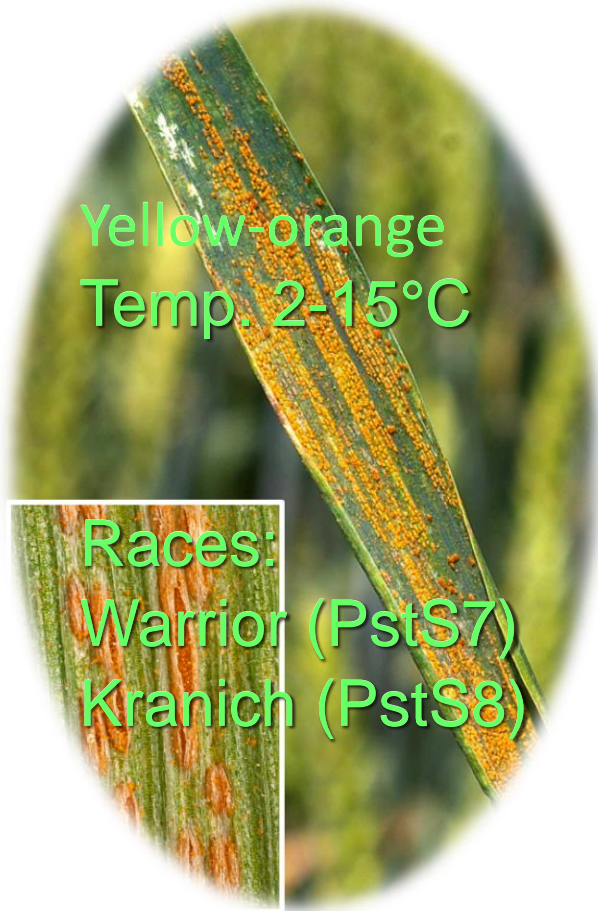
Acknowledgements

Wheat rust 'Cereal killers'

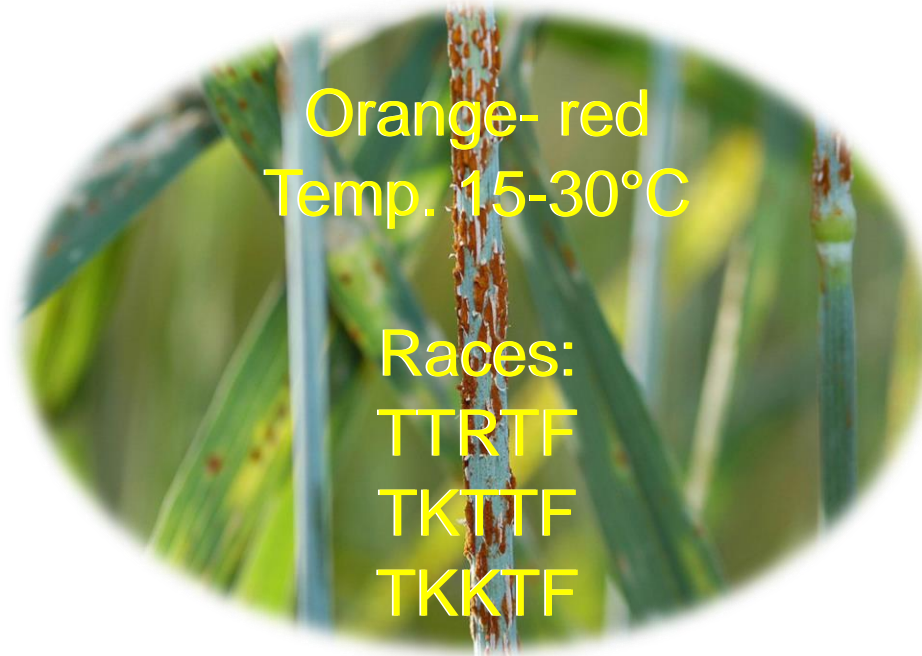
15 million tons of wheat are lost yearly, resulting in a \$3 billion loss.

Enough to bake ~ 25.5 billion loaves of bread
425 million peoples annual bread consumption

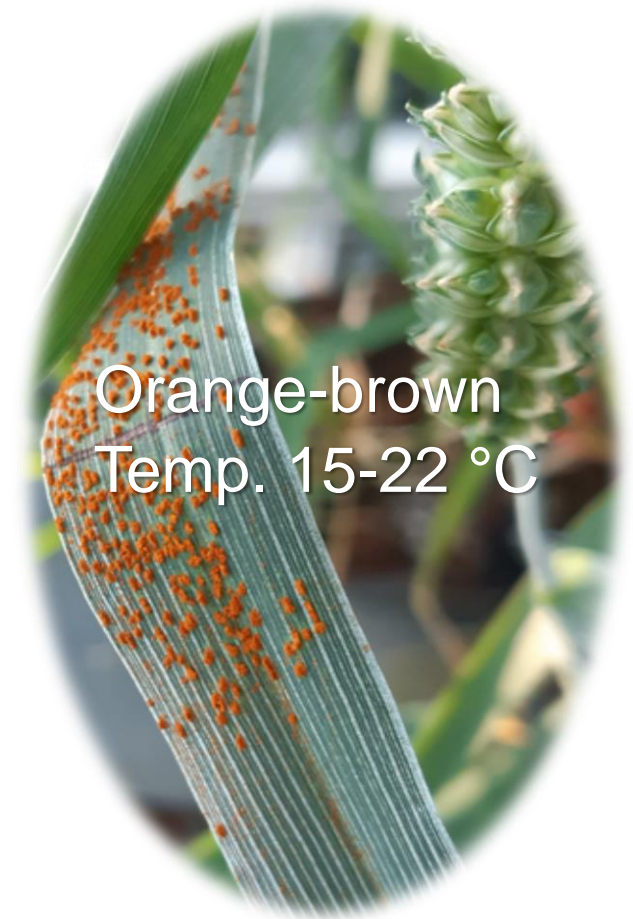
The three wheat rust fungi



Yellow/Stripe rust
Puccinia striiformis
f.sp.tritici/Pst



Black/Stem rust
Puccinia graminis f.sp.
tritici/Pgt



Brown/Leaf rust
Puccinia triticina

Life cycle of Rust fungi (Cereal Killer)



Common barberry removal Campaign (1918 to 1980)



BARBERRY BEE, RED WING, MINNESOTA, SEPTEMBER 20, 1922



Barberry eradication in Western Europe

UNITED STATES
DEPARTMENT OF AGRICULTURE
DEPARTMENT CIRCULAR 269

DESTROYING BARBERRY

BLACK STEM RUST of wheat is controlled in much of western Europe by the eradication of the common barberry. This is not a theory.

DESTROYING BARBERRY

BLACK STEM RUST of wheat is controlled in much of western Europe by the eradication of the common barberry.

This is not a theory. It is a fact. Since the barberry-eradication law was passed in the United States much has been heard about the destructive epidemics of black stem rust in Europe. Less was known about what other countries were doing. In the United States asked whether exterminating barberries would control the rust. The writer therefore advised the Department of Agriculture to go to Europe during the spring and summer of 1922.

Black rust does no appreciable damage in western Europe (Fig. 1) from which barberry has been eradicated.

DENMARK KILLS BARBERRY AND STOPS RUST.

Denmark also has prevented attacks of black stem rust by eradicating barberry bushes. If there ever was an established scientific fact, this is one. For years rust attacks had been severe in Denmark. Barberry bushes had been brought into Denmark and planted some time during the seventeenth century. By about 1805 they had become so numerous that black rust began to be destructive. Many farmers

NO BLACK RUST WITHOUT BARBERRY IN NORWAY AND SWEDEN.

Agriculturists in Sweden are making desperate attempts to secure the passage of an effective barberry-eradication law. And well they may. If anyone doubts the beneficial effect of destroying the barberry he should travel from Denmark to Sweden. There are enormous numbers of barberries in some districts of Sweden, and the attacks of black rust in those regions are terrific. Between Stockholm and Upsala, in the summer of 1922 fields of oats were black

TTTTF, a kind of stem rust, damages tens of



By Chris Hill

@ChrisHill75

Agricultural Editor



Work News Events About Countries 

News > Successful surveillance results in early first detection of Ug99 in South Asia

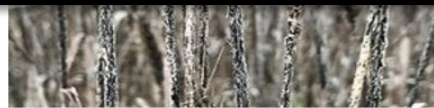
NEWS DISCOVERY ENVIRONMENTAL HEALTH AND BIODIVERSITY INNOVATIONS

Successful surveillance results in early first detection of Ug99 in South Asia

The detection of a Ug99 race in Nepal highlights the effectiveness of the wheat rust surveillance and monitoring systems developed by CIMMYT and partners.

By CIMMYT

April 3, 2024



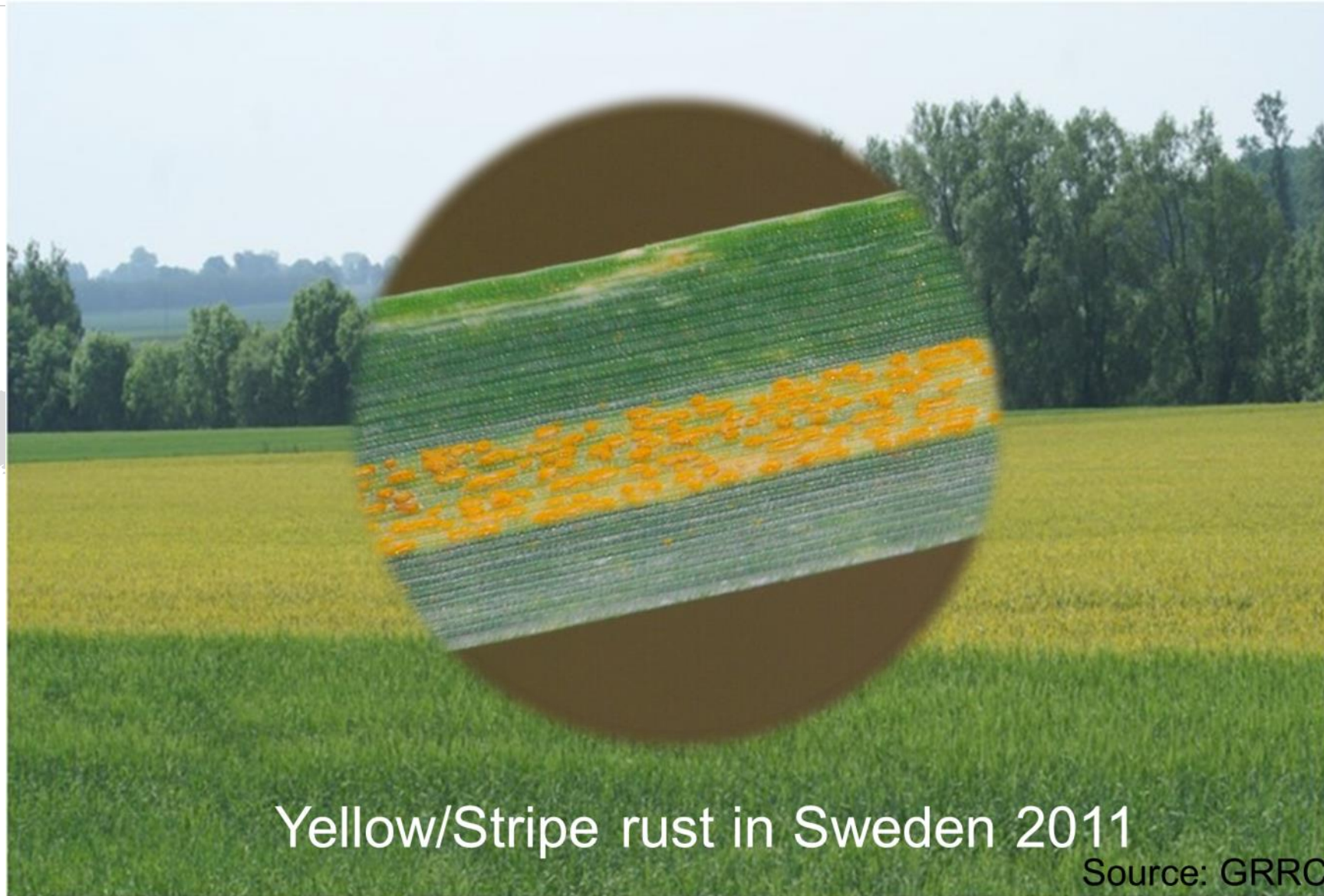
Grass has, for millions of years, been attacked by rust fungi. When man began farming and breeding various types of grass, such as our common grains, the

Picture: Paul Fenwick, Limagrain UK.
(Image: Paul Fenwick, Limagrain UK)

crop
lls
ther

k.

Stripe Rust is spreading rapidly across the globe.



How to respond once disease becomes established?

Agronomic practices



Modern Fungicides



Resistance breeding



European Union Farm to Fork strategy aims to reduce pesticide use and risk by 30% by 2030

Resistance breeding

- Environmental friendly
- Economical effective

Wild relative to
wheat

Durable source
Biotic/abiotic



(*Secale cereale*)

Rye chromosomes

1R

2R

3R

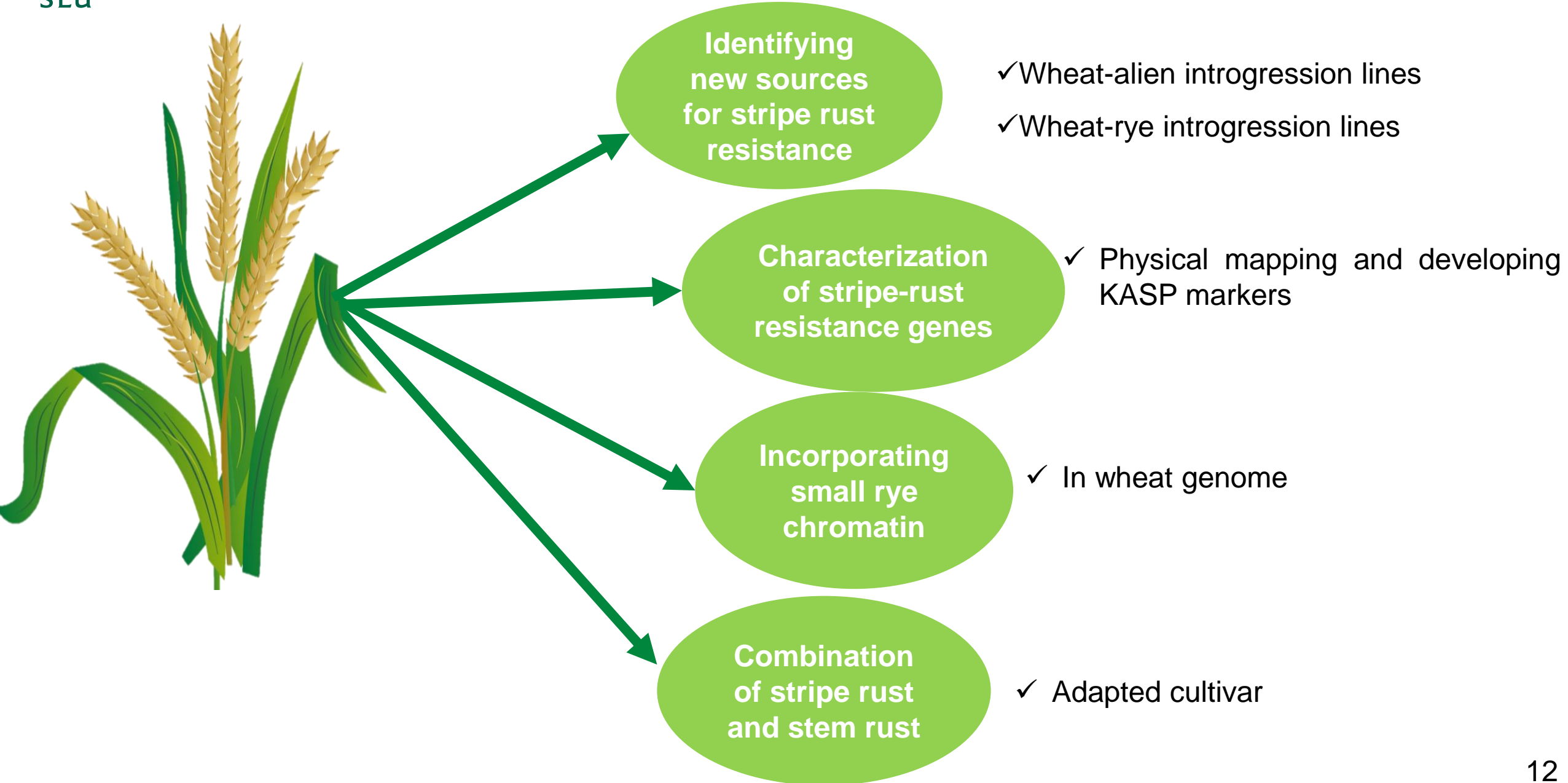
4R

5R

6R

7R

Study Aim and Objectives





Wheat-alien introgression lines



Plant materials

- ✓ **Wheat-rye introgression lines i.e., SLU124, SLU126, SLU128 etc.**
- ✓ **Wheat-*Leymus racemouses* introgression lines**
- ✓ **Wheat-*Leymus mollis* introgression lines**
- ✓ **Wheat-*Thinopyrum junceiforme* introgression lines**



Materials and Methods

- ✓ A set of winter and spring wheat-alien introgression lines
- ✓ Agronomic performances, diseases and pests screenings
- ✓ Allelopathic Potential
- ✓ End-use quality analysis (Grain Samples for Micronutrients Concentration and Protein Composition)



Results



- ✓ wheat–rye introgression lines highly resistant
- ✓ Resistance towards the Hessian fly
- ✓ High levels of zinc and iron
- ✓ Very low cadmium concentrations
- ✓ Large variation in bread-making quality
- ✓ Improve production, resistance and quality



Basis for Durable Resistance and Quality Characteristics in Bread Wheat

Eva Johansson^{1*}Tina Henriksson²Maria Luisa Prieto-Linde¹Staffan Andersson¹Rimsha Ashraf¹Mahbubjon Rahmatov¹

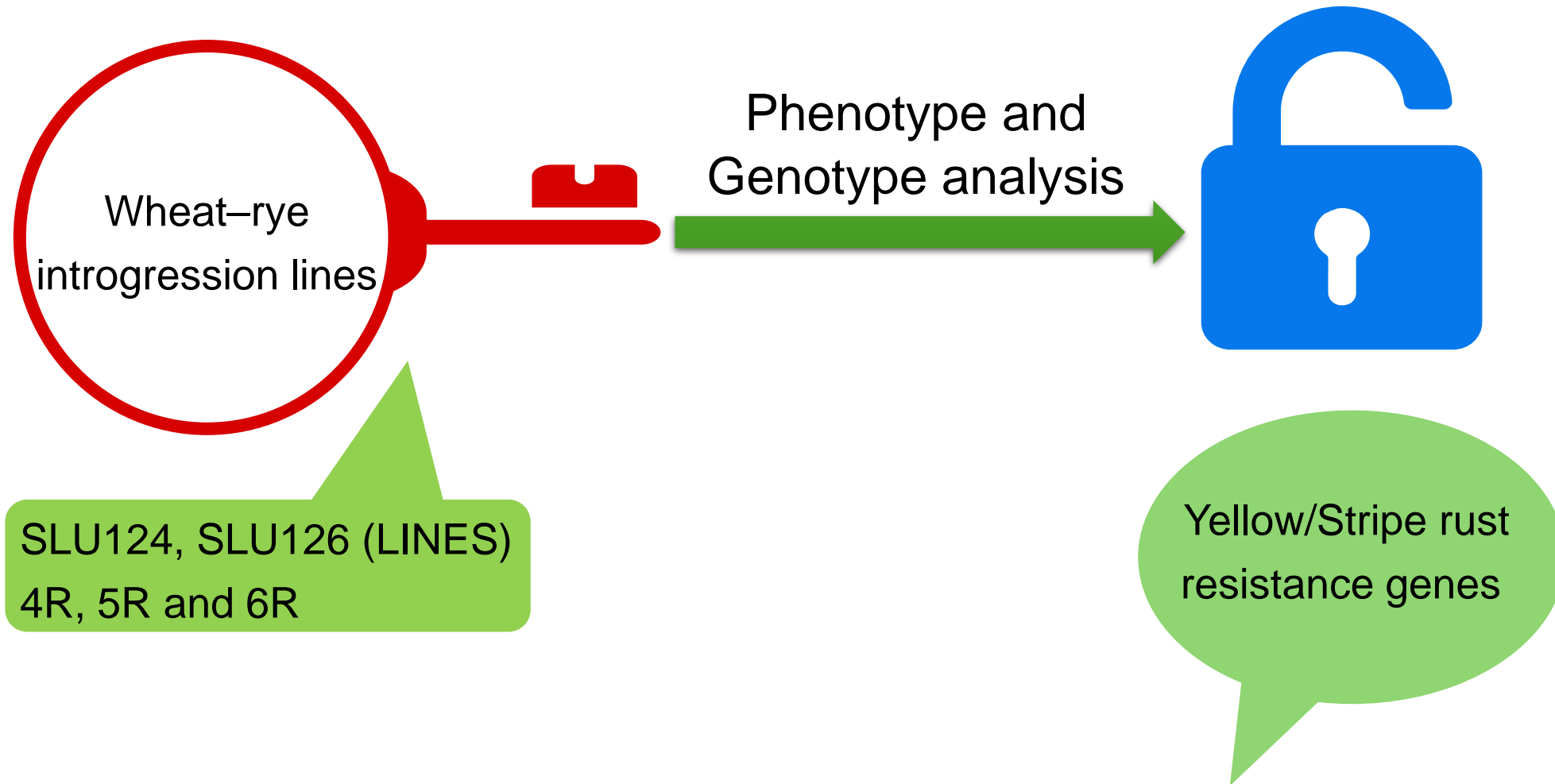
¹ Department of Plant Breeding, The Swedish University of Agricultural Sciences, Alnarp, Sweden

² Lantmännen Lantbruk, Svalöv, Sweden

Wheat productivity has been significantly improved worldwide through the incorporation of novel genes from various gene pools, not least from wild relatives of wheat, into the commonly cultivated bread and durum wheat. Here, we present and summarize results obtained from a diverse set of wheat-alien introgression lines with mainly introgressions of rye, but also of

Rye chromosome (6R) for stripe rust resistance gene (*YrSLU*)





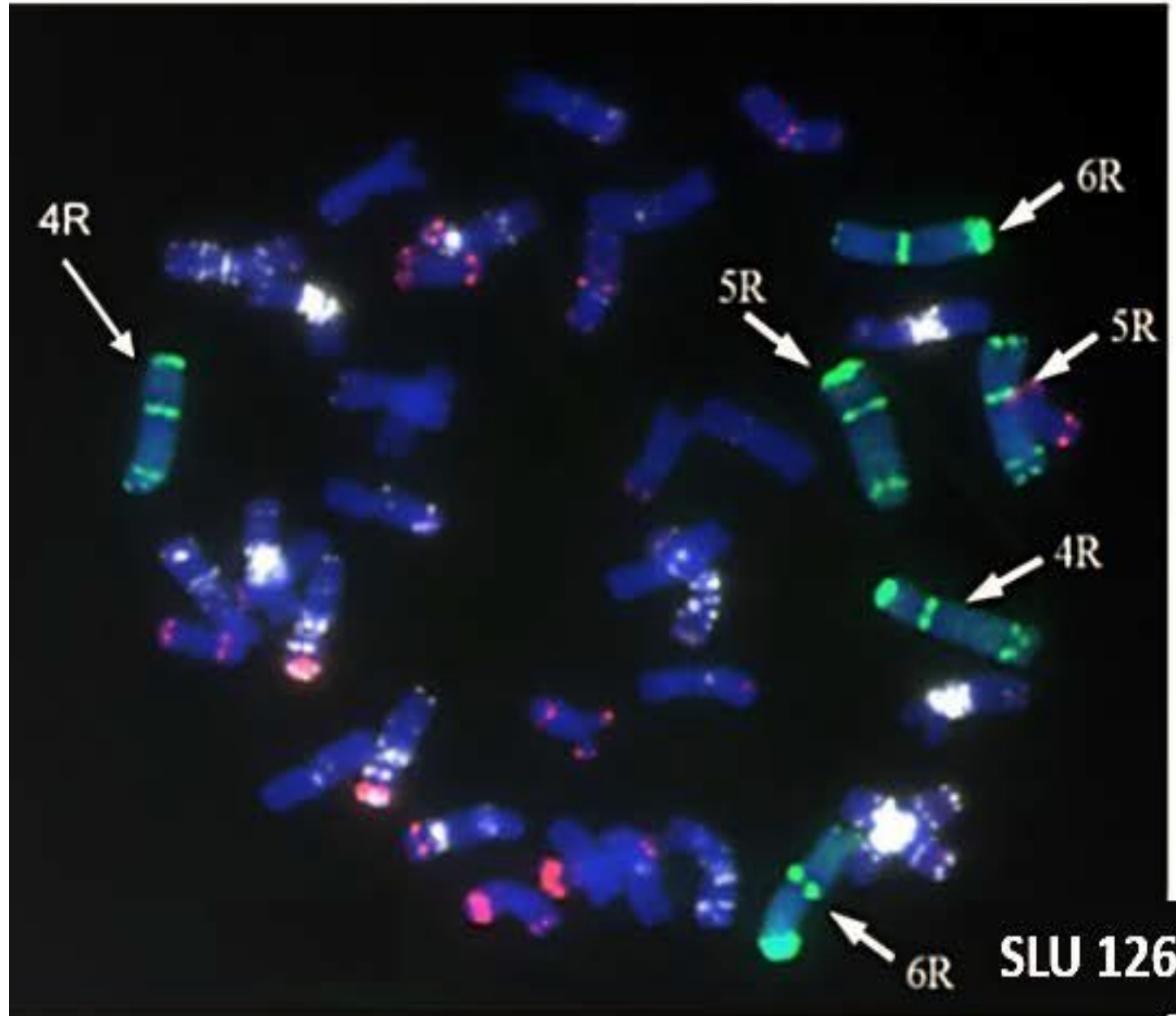
Line SLU126



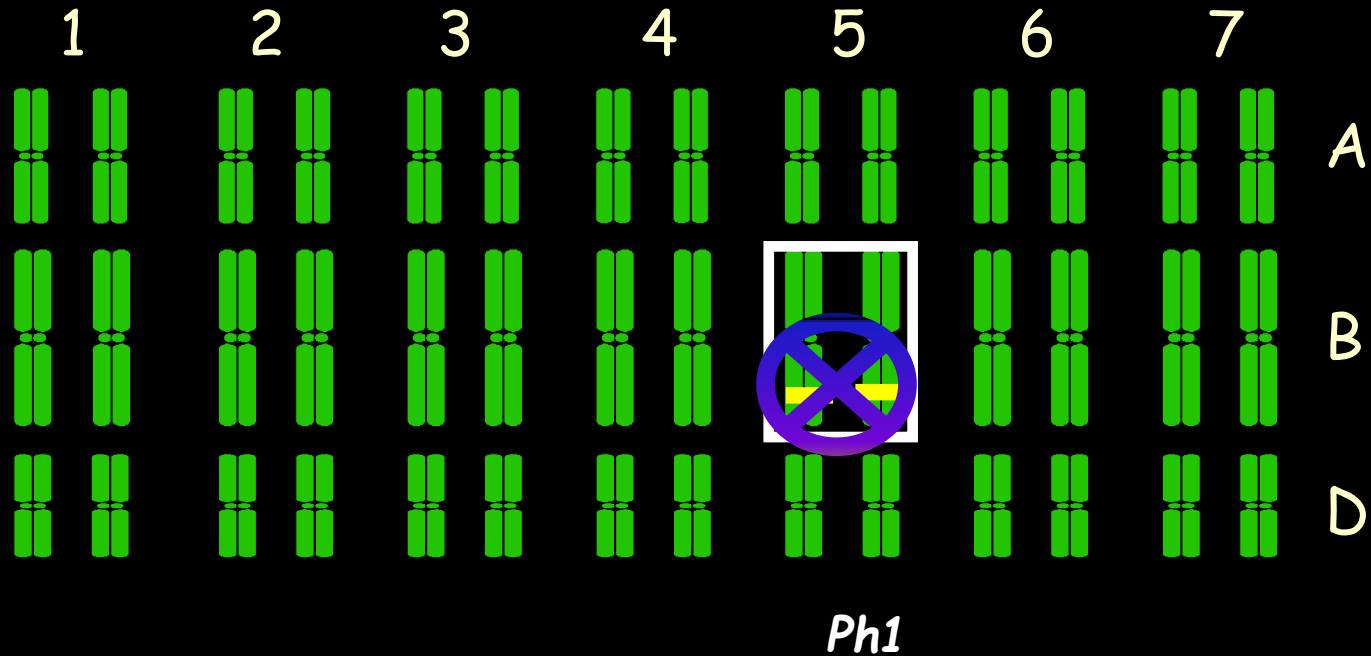
KRANICH UK94/519 DK66/02 Taj01a/10 ER02/03 DK11/09 DK71/93 AF87/12 DK09/11 SE100/09 TR34/11

30 *Yr* races

Fluorescent in situ hybridization (FISH)



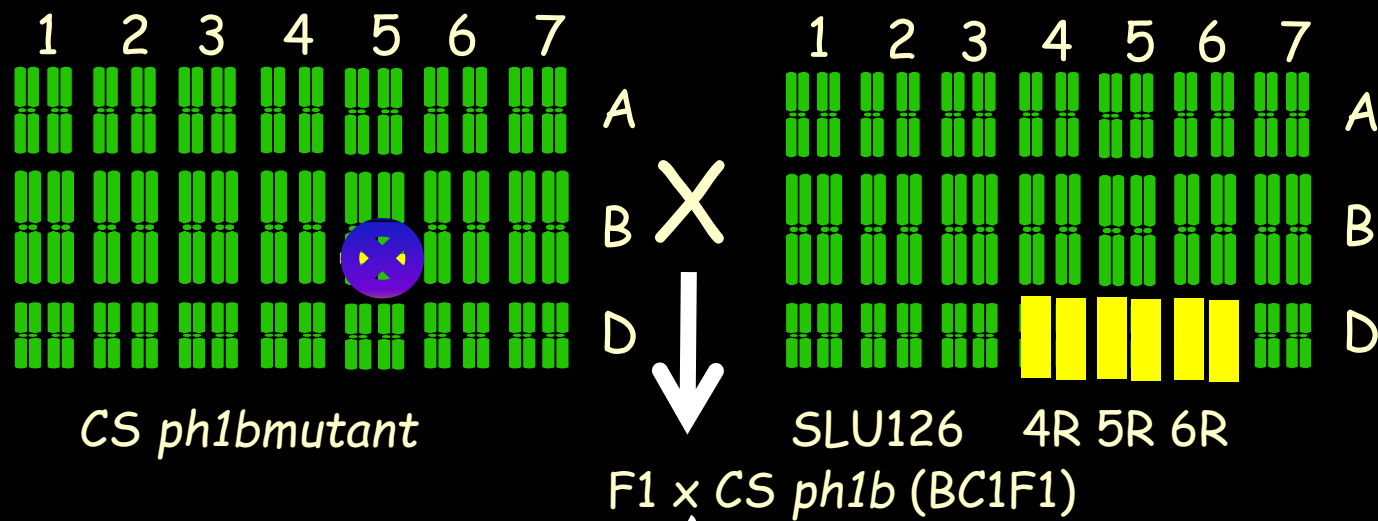
The role of *Ph* (*Ph1*) locus



Pairing homoeologous loci (*Ph*)

(Sears, 1977)

Induce Homoeologous Recombination Using CS *ph1b* Mutant

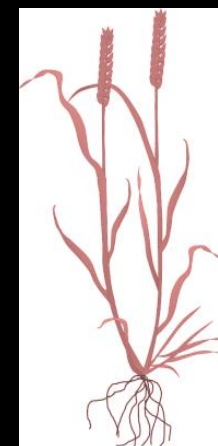


➤ Identify recombinants

- Yellow rust seedling resistance tests PSTv-14, PSTv-37, PSTv-40, PSTv-218 and PSTv-221
- Molecular markers analysis for 4R, 5R, 6R rye chromosomes



BC1F2
BC1F3
BC1F4



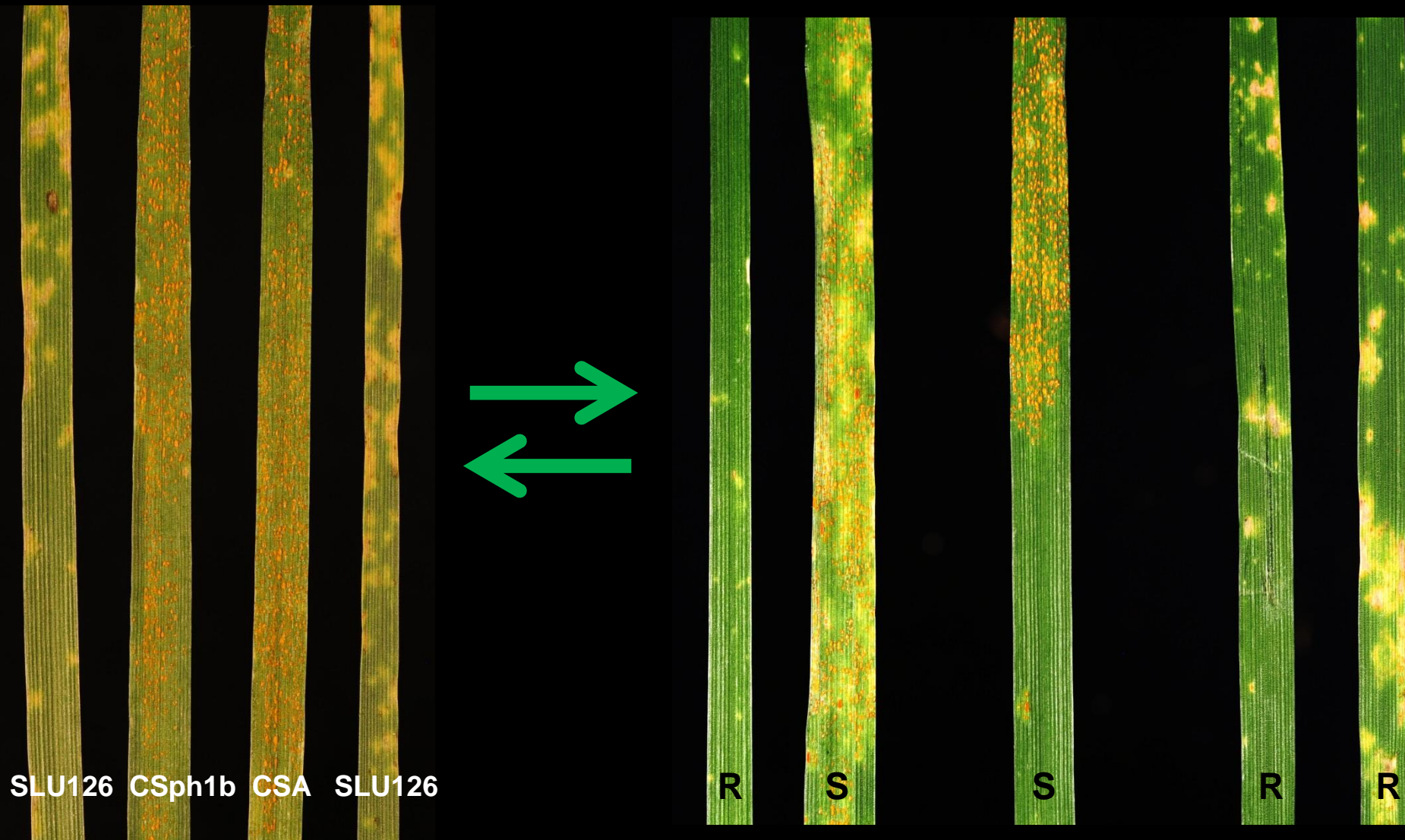
BC2F1
BC2F2
BC2F3

Yr seedling resistance tests in BC₁F₁

Mix of PSTv14 and PSTv37 races

Parental Lines

BC₁F₁



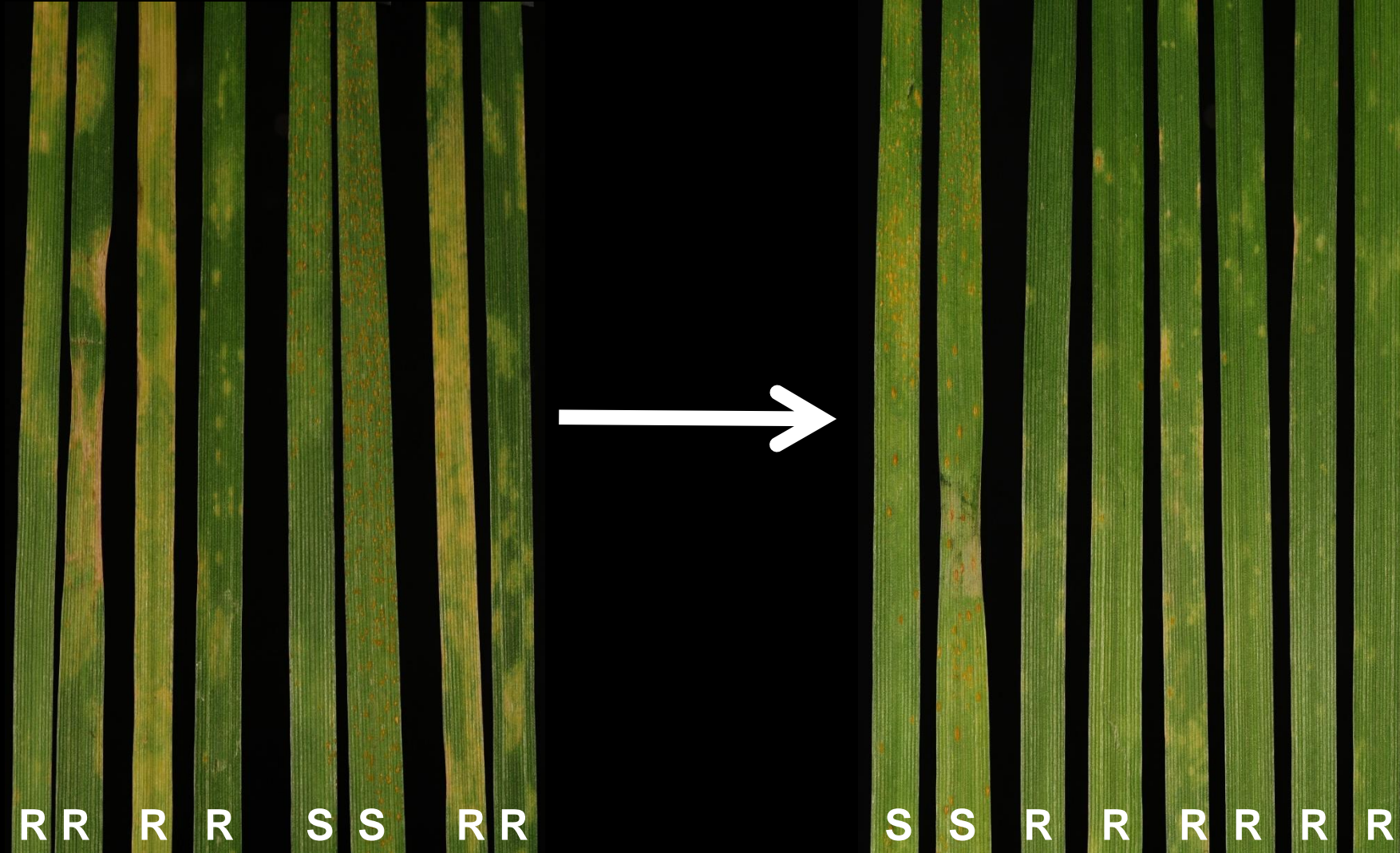
Yr seedling resistance tests in BC1F4 and BC2F3 (Family N3-5)

Line 245
BC₁F₃

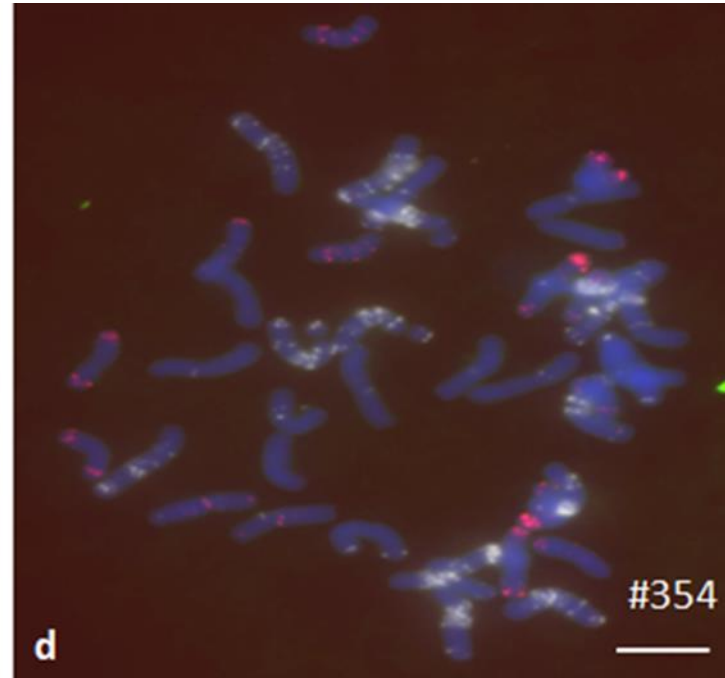
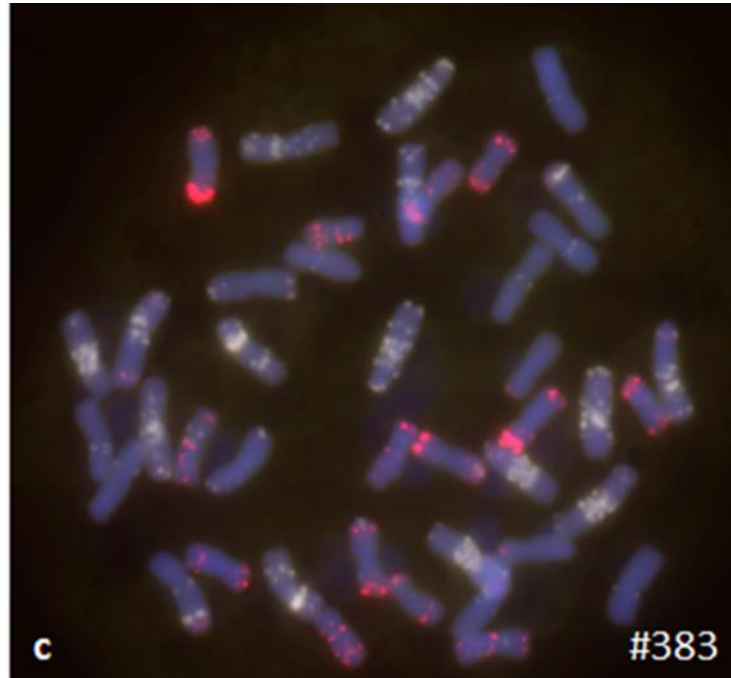
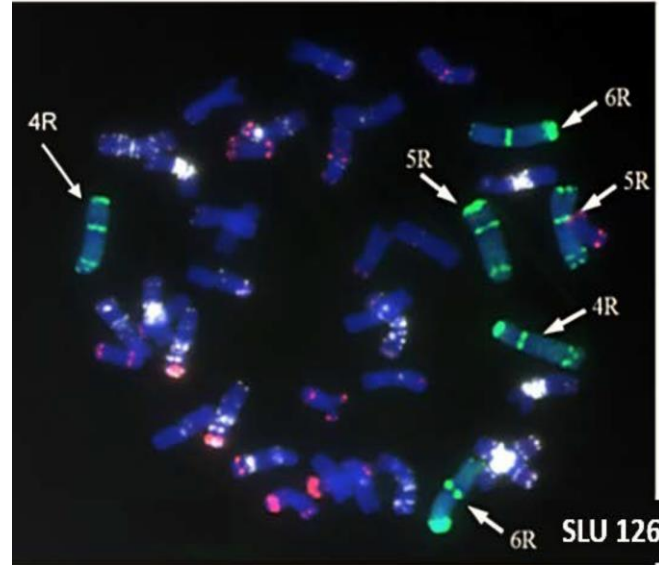
Line 180
BC₂F₂

Line 180
BC₂F₃

Line 245
BC₁F₄



Cytogenetic analysis (FISH and GISH) in BC1F4 and BC1F3



Genotyping by Sequencing (GBS)

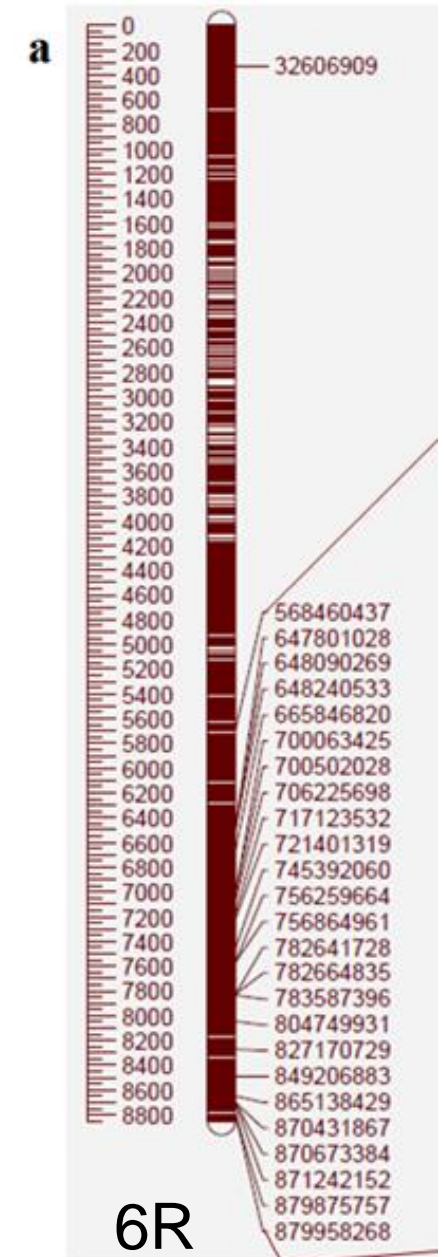
- ✓ GBS in parental lines (SLU126, *CSph1b*, CSA)
- ✓ Aligned the GBS to wheat and rye reference genomes

| Chromosome | High-quality SNP (after filtering) |
|------------|------------------------------------|
| 6R | 10675 |

NLR (Nucleotide binding site leucine rich repeats) sequences for Rye

✓ NLR genes for Rye by using NLR parser software

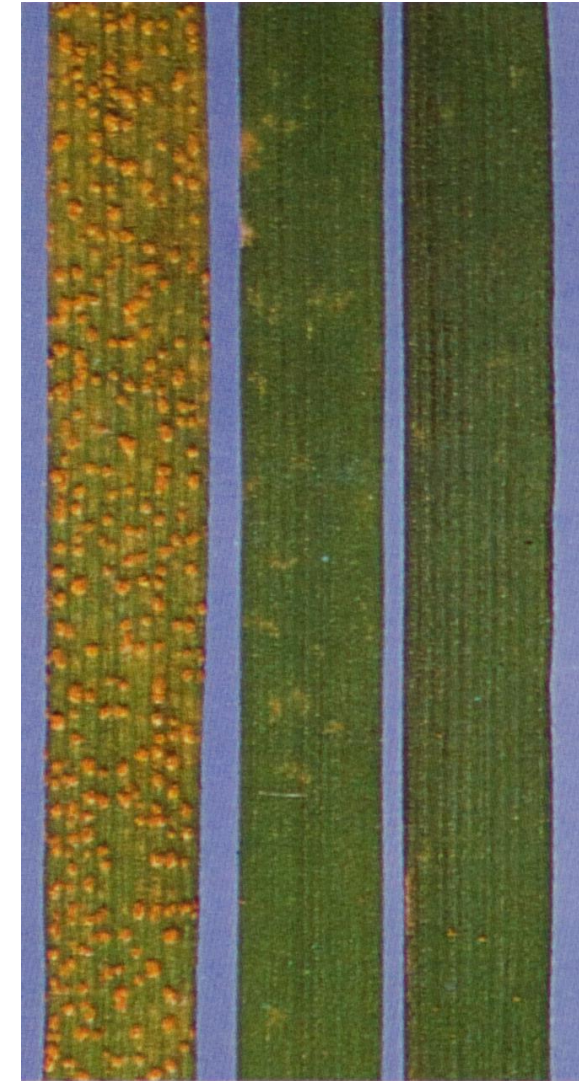
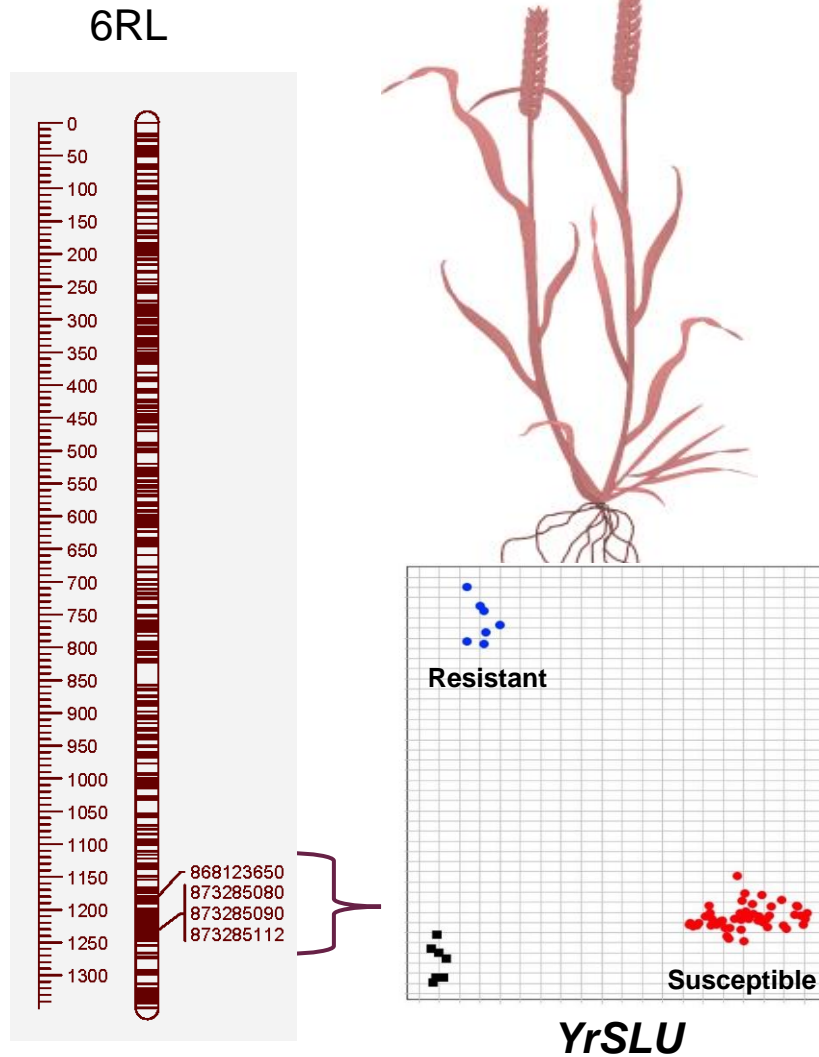
| Rye Chromosome | NLR genes | Designated NLR to SLU126 (6R) |
|----------------|-----------|-------------------------------|
| 6R | 184 | 26 |



New resistance genes *YrSLU*

6DS.6DL.6RL.6DL

wheat-rye cryptic translocation



CSph1b *YrSLU1* *YrSLU2*

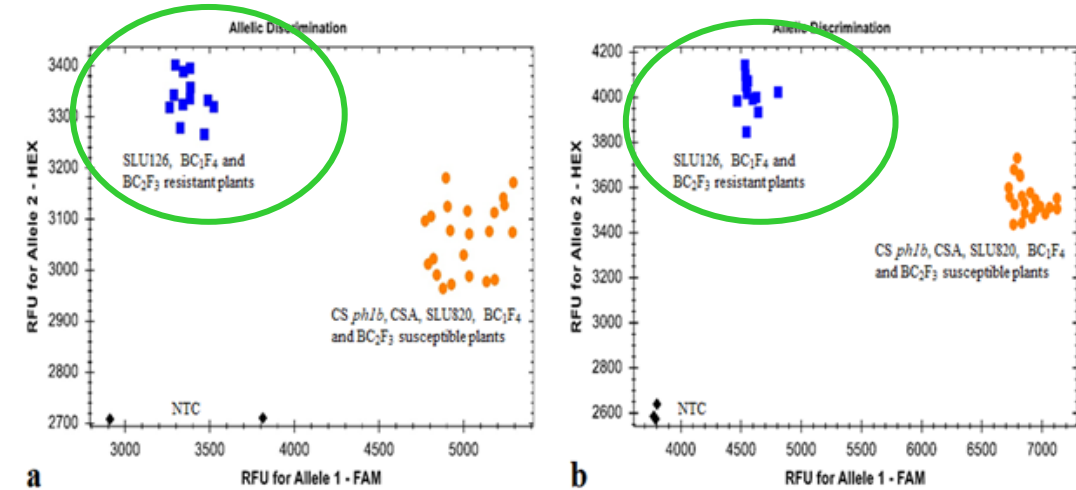
Further analysis



Resistant to PSTv-14, PSTv-37, PSTv-40 and PSTv-221



Segregating against Pstv-218

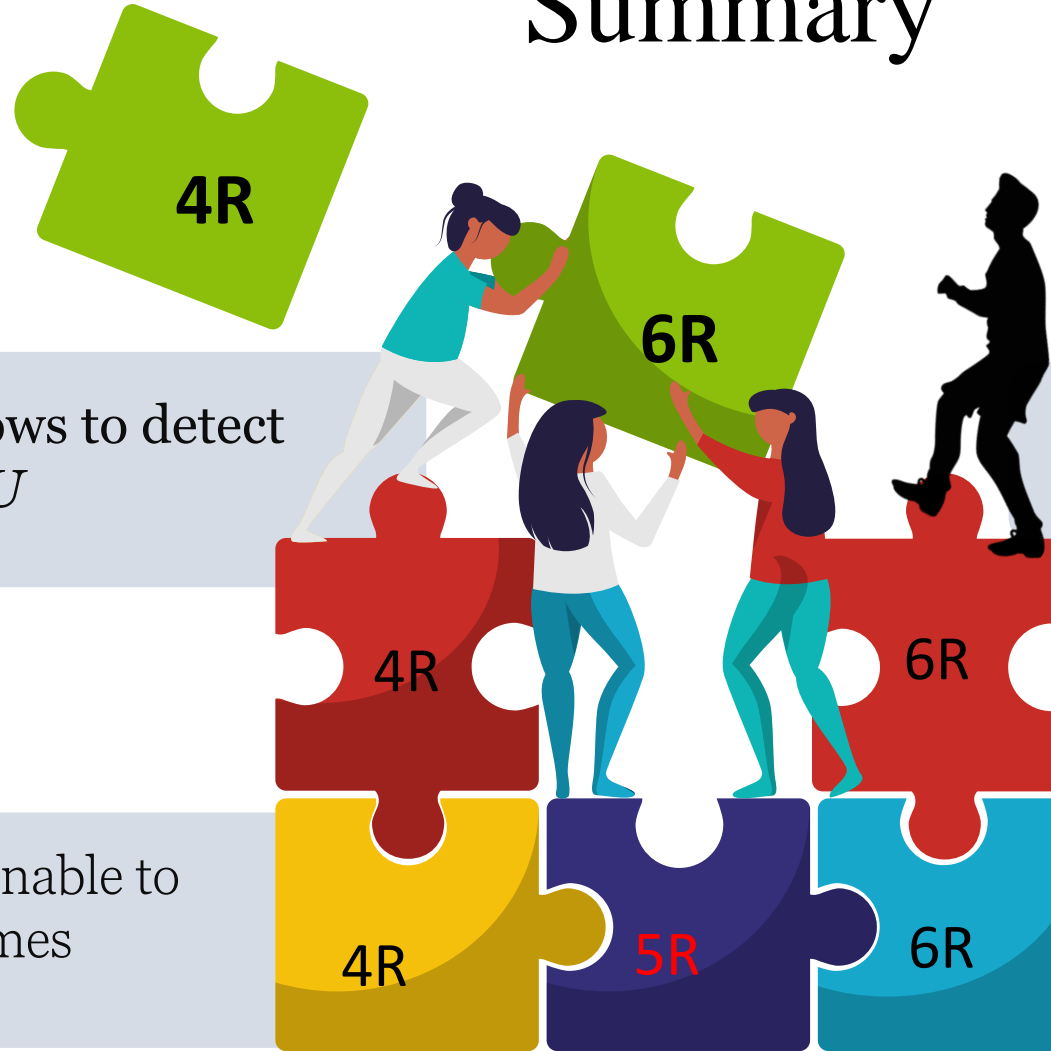


YrSLU1

YrSLU2

Yellow/Stripe rust resistance genes

Summary



GBS analysis allows to detect
YrSLU

Yes, its 6R Cryptic translocation


Molecular markers unable to
detect rye chromosomes

Cytogenetic also confirms no
rye chromosome in resistant
lines

Wheat-rye introgression lines
SLU126 (6R)

RESEARCH

Identification of a Small Translocation from 6R Possessing Stripe Rust Resistance to Wheat


Rimsha Ashraf, Eva Johansson, Pernilla Vallenback, Brian J. Steffenson, Prabin Bajgain, and Mahbubjon Rahmatov 

Affiliations 



Published Online: 23 Mar 2023 | <https://doi.org/10.1094/PDIS-07-22-1666-RE>

 SECTIONS

 ABSTRACT

 PDF

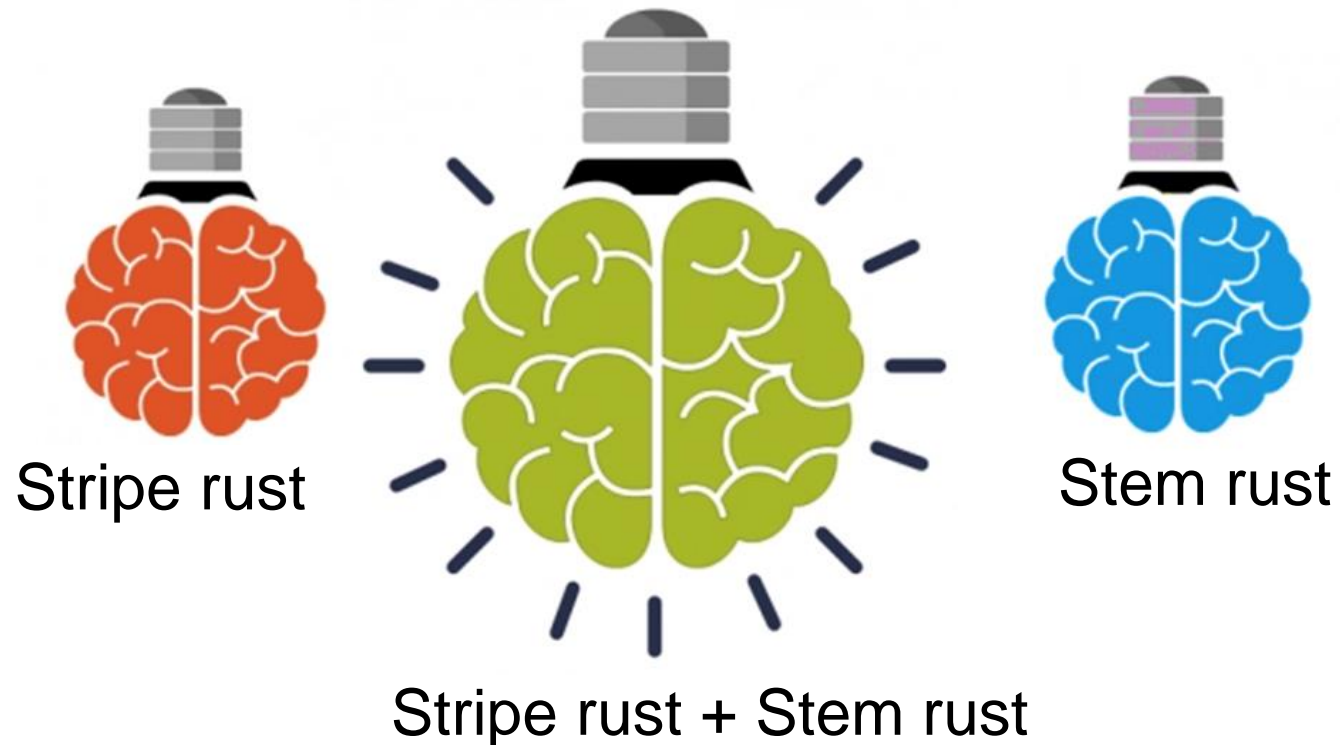
 e-Xtra

 TOOLS  SHARE

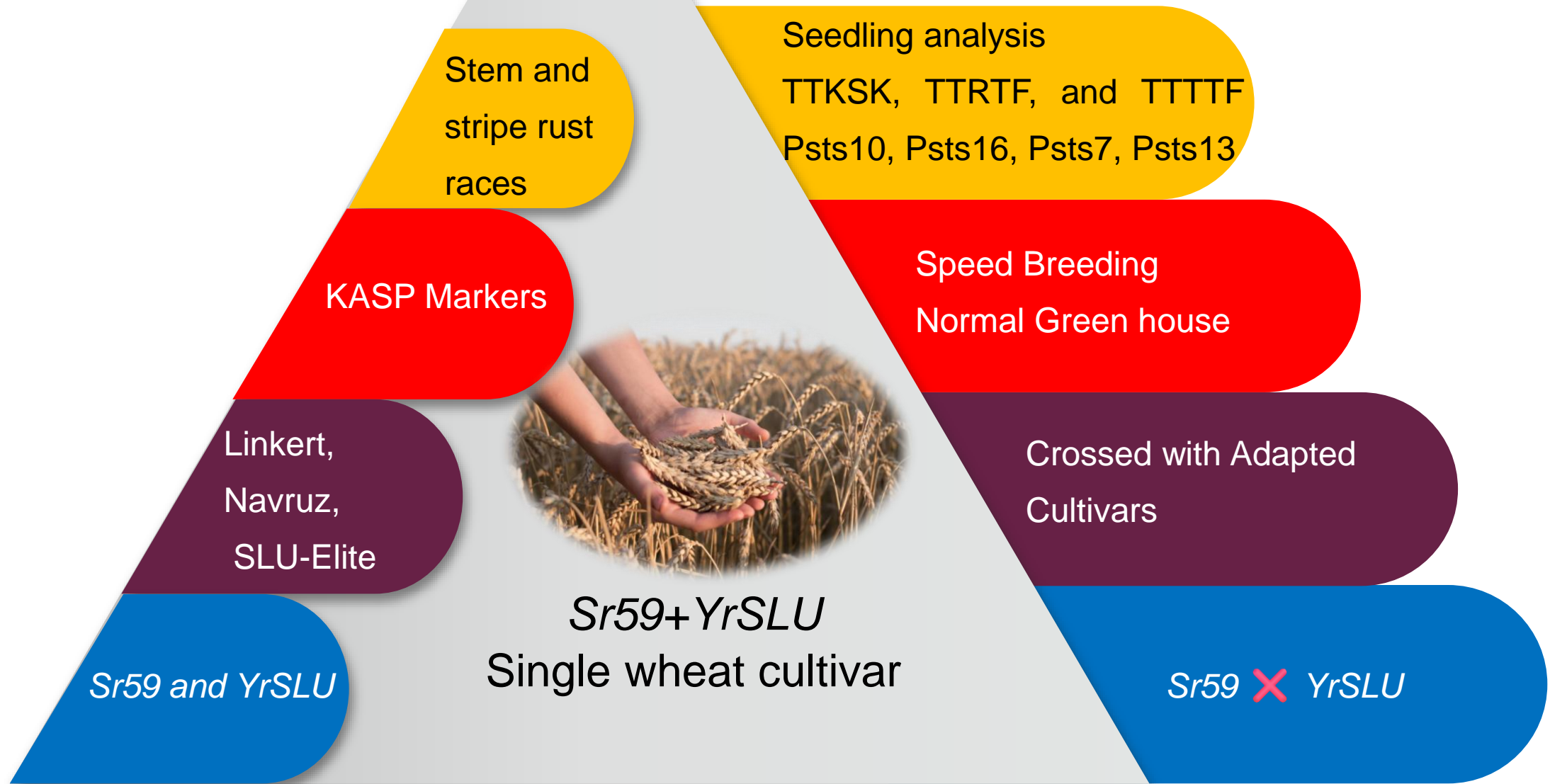
Abstract

Wheat stripe rust, caused by *Puccinia striiformis* f. sp. *tritici* Eriks. & E. Henn, is the most devastating fungal disease of bread wheat. Here, a wheat-rye multiple disomic substitution line, SLU126 4R (4D), 5R (5D), and 6R (7D), possessing resistance against

Opportunities and challenges with the combining of novel stem and stripe rust resistance genes in the same genotype



Marker-assisted gene pyramiding



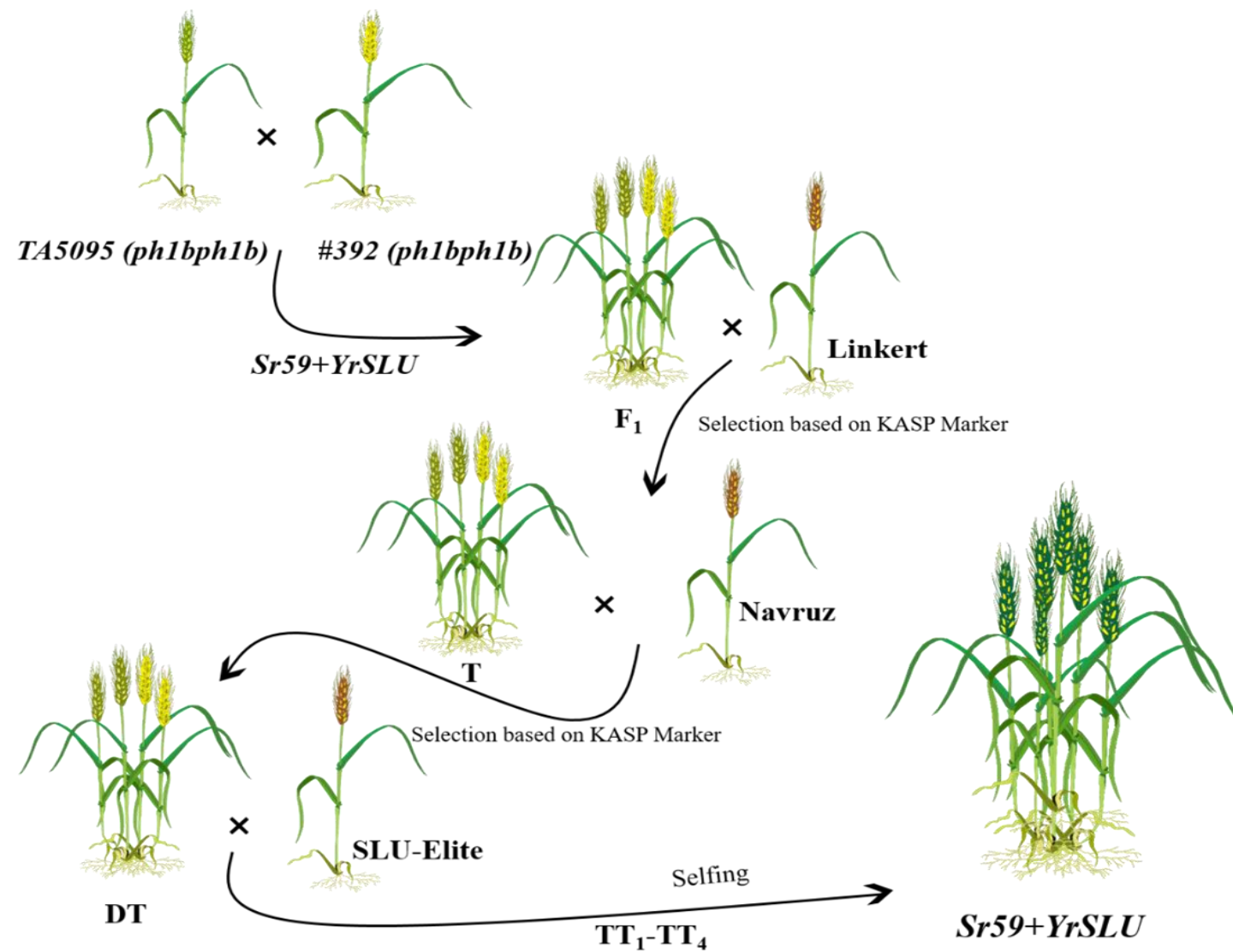


Figure . Scheme of the performed marker-aided gene pyramiding for stem rust and stripe rust resistance, Top-cross (T), Double Top-cross (DT), Triple Top-cross 1 (TT₁), Triple Top-cross 4 (TT₄)

Speed Breeding

21 April 2021



Planting

2 June 2021



Grain Filling

7 June 2021



Dough Development

• Within 6-7 weeks

Greenhouse condition

- Humidity: 40-80%
- Temperature: 18⁰-24⁰°C (20 hours Photoperiod)
- High pressure sodium lamps

Main Results

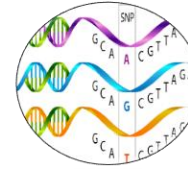
- ✓ Successful transfer of *Sr59* and *YrSLU*
- ✓ Complete resistance to stem rust races
- ✓ Segregated against stripe rust races
- ✓ Confirm genes of interest



Conclusions



Wheat-alien introgression lines has distinct potential for wheat improvement



Validation of KASP markers for *YrSLU* genes transferred from 4R and 6R



Wheat-rye substitution lines possess resistance to 30 races of stripe rust.



Cryptic translocation 6DS.6DL.6RL.6DL temporarily designated *YrSLU* genes



NLR genes for 4R, 5R and 6R rye chromosomes

MAS gene pyramiding provide durable disease resistance against stripe and stem rust



Phenotype and genotype analysis, effective in identifying the source of resistance to stripe rust



End-use quality analysis

05

06

Gene pyramiding into Swedish wheat cultivar

Agronomic Performances, Diseases and Pests

04

Evaluation with additional *Yr* races

03

Functional Analysis

02

Gene Designation for *YrSLU*

01

Acknowledgements



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wheat breeder
Lantmännen Lantbruk



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