













## Searching needles in haystacks

genomics- and phenomics-based exploitation of wheat genetic resources for disease resistance breeding









## GeneBank2.0: Potential and limits of genomics-informed pre-breeding

#### 1. Yield and yield stability

- Grain yield
- Yield components
- Winter hardiness
- Lodging resistance
- Nutrient use efficiency
- 2. Hybrid seed production
- Pollination capability
- Restorer genes

#### 3. Disease resistance

- Septoria tritici blotch
- Fusarium head blight
- Leaf rust
- Powdery mildew
- Yellow rust
- Tan spot
- Eyespot
- Septoria nodorum
- Soil-borne mosaic virus

#### 4. Quality

- Ash content
- Flour yield
- Endosperm texture
- Protein content
- Zeleny sedimentation
- Hagberg falling number
- Water absorption
- Dough quality
- Baking volume



## Challenging logistics to activate an entire collection

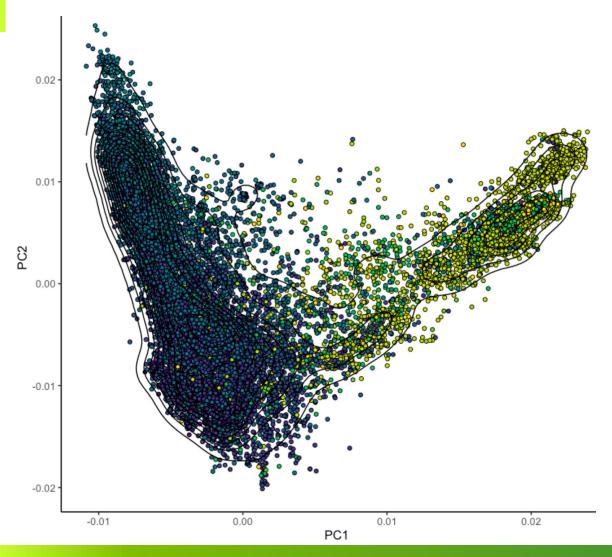


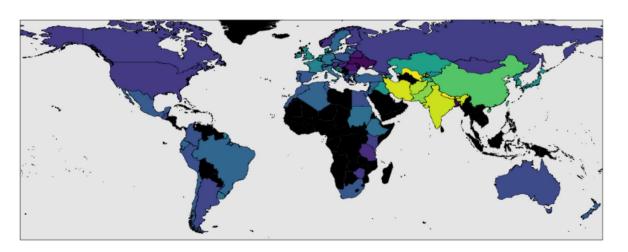






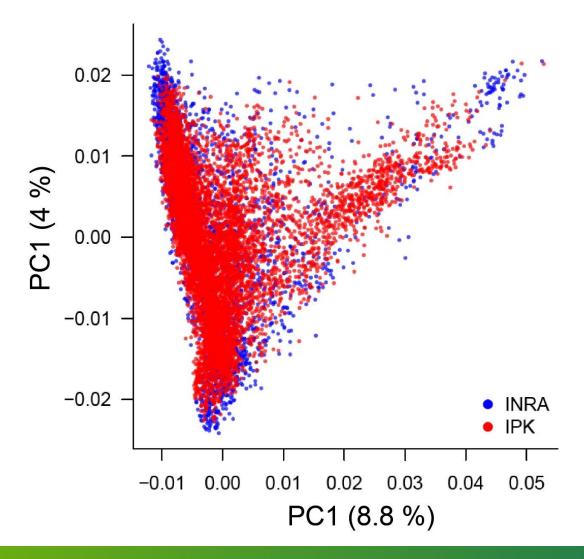
## Genotype atlas for 8,762 winter and 5,861 spring wheat accessions







## ... towards a Pan-European genotype atlas



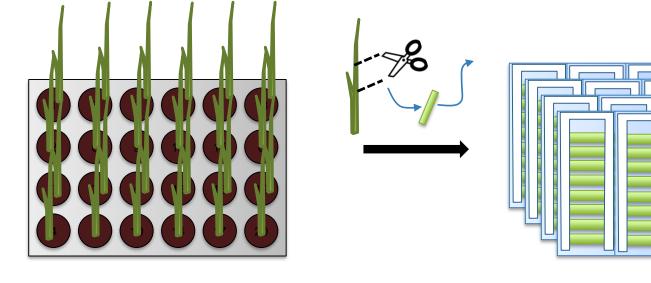


## Disease resistance phenotyping

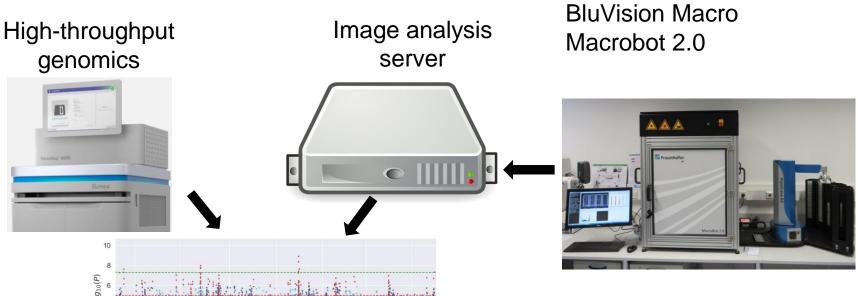


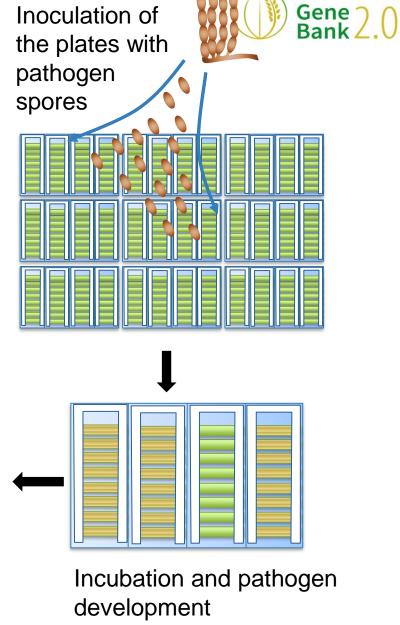






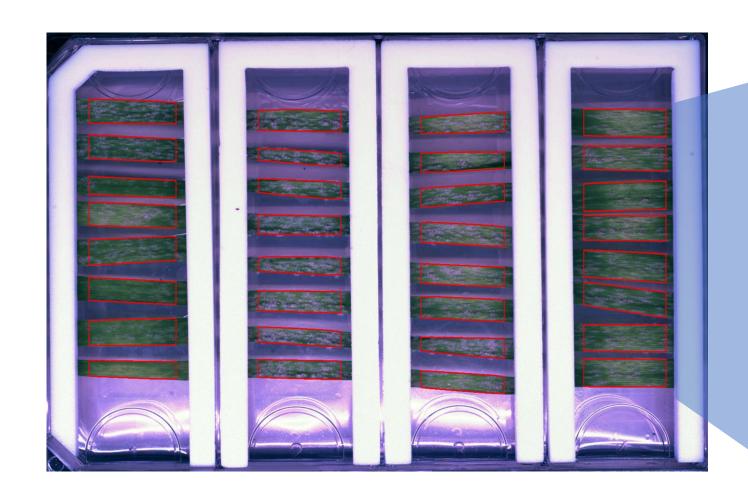
## Macrophenotyping platform for disease resitance





## Precise phenotyping for disease resistance (powdery mildew)



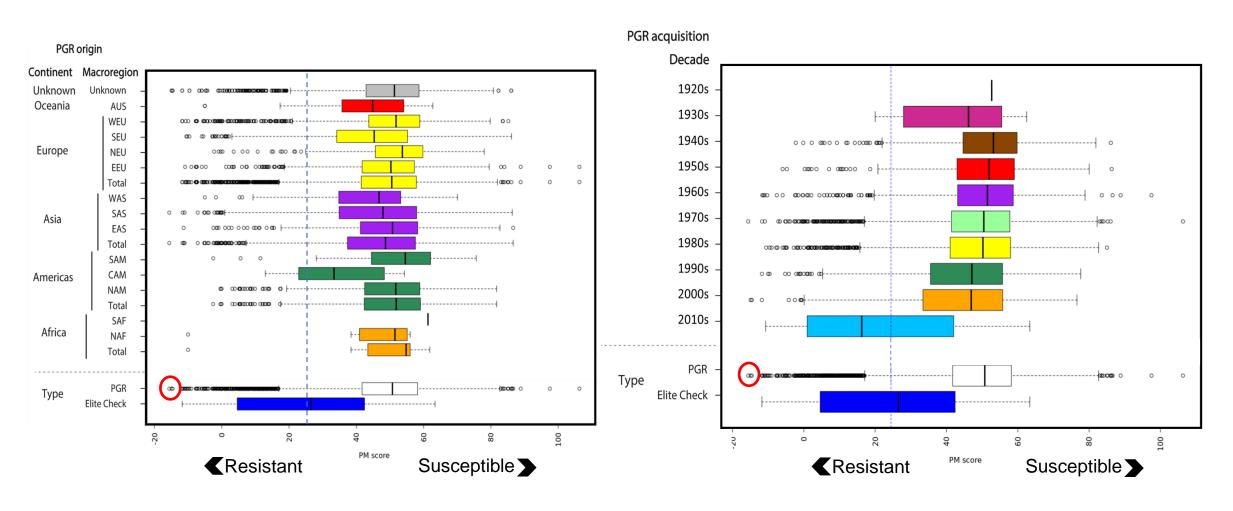




## Searching needles in a haystack



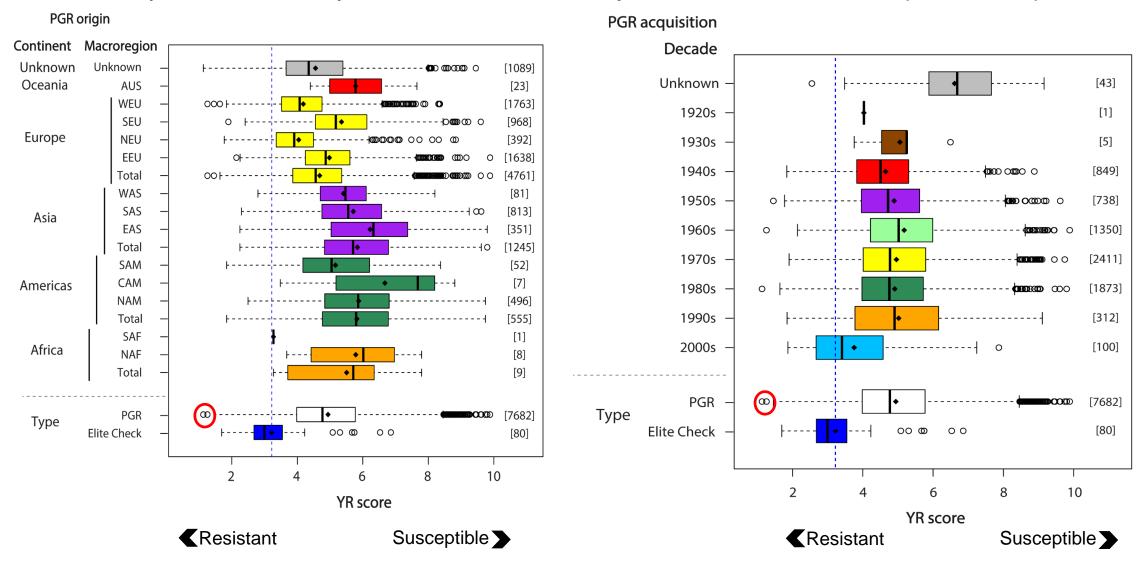
#### Spatial and temporal distributions of powdery mildew resistance



## Searching needles in a haystack

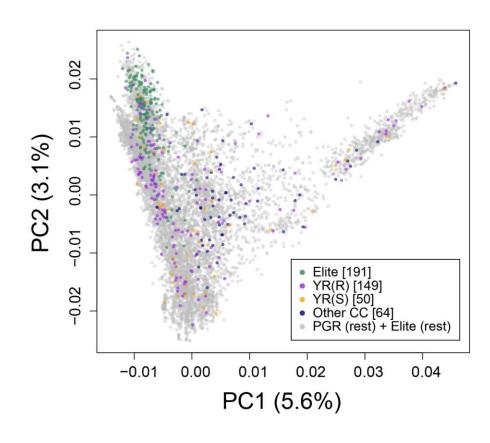


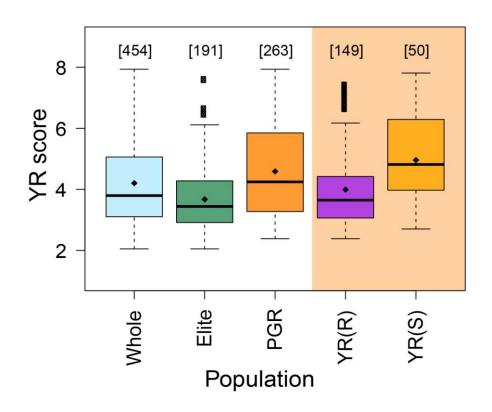
Spatial and temporal distributions of yellow rust resistance (field data)





## Designing a trait-customized core collection

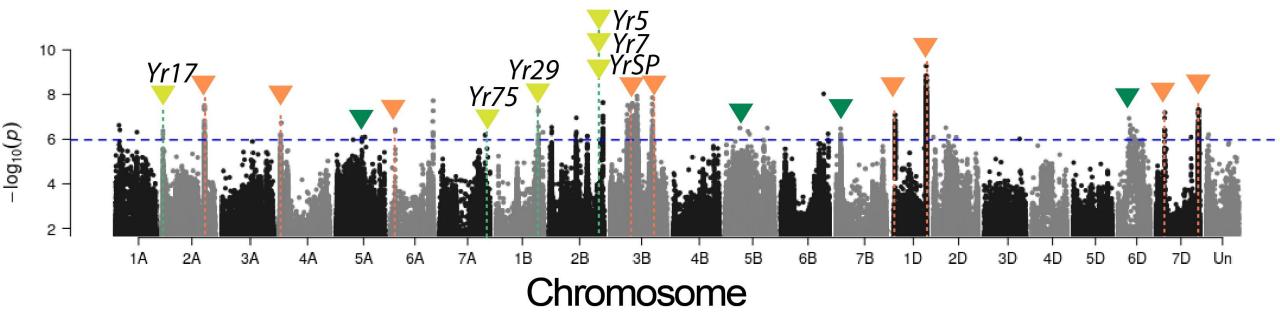






## Mining for novel resistance genes

- Known YR resistance loci
- Resistance alleles are lost in elite pools
- Resistant alleles are almost fixed in elites





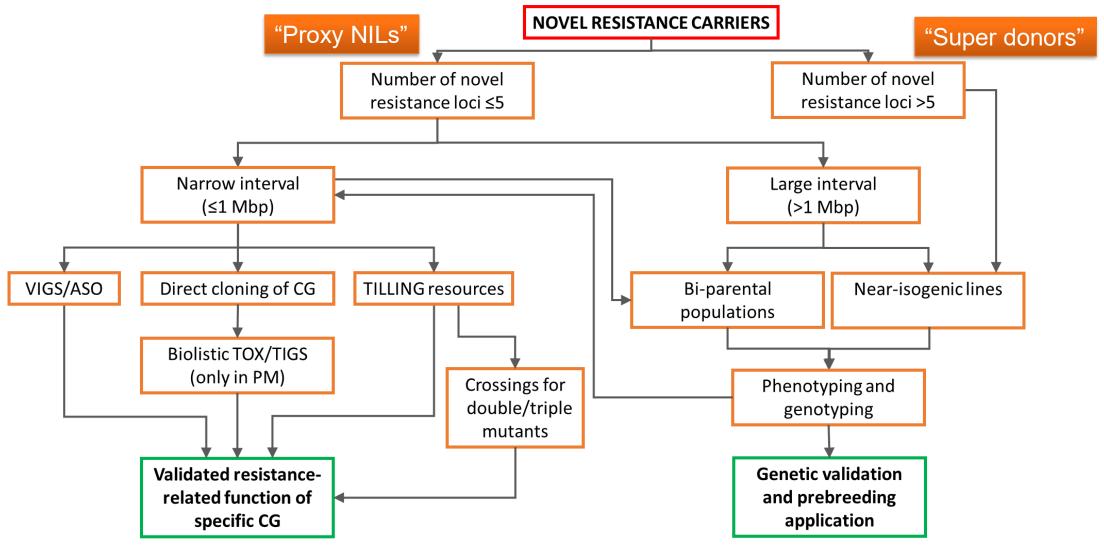
# Graphics removed In publication

Comparison of chromosomal regions in near-isogenic lines (NILs) harboring known and novel YR genes and loci

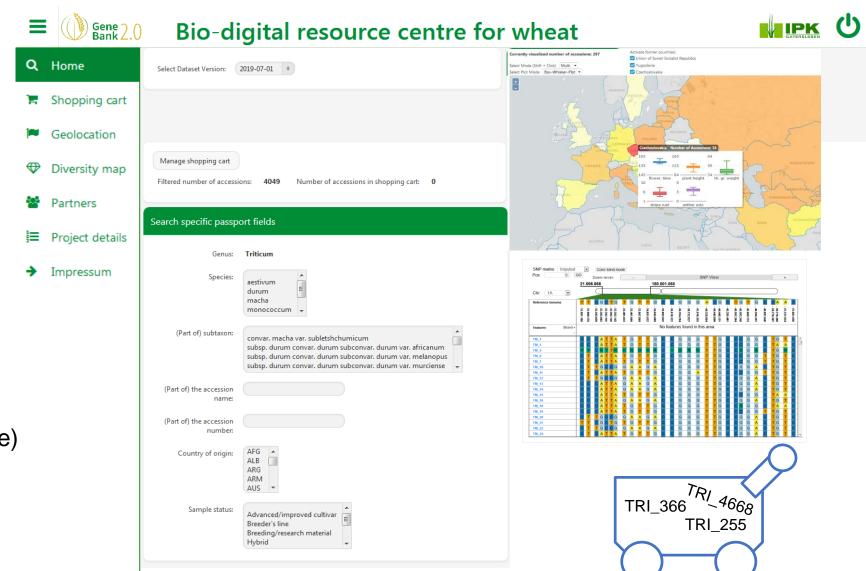
- Introgressions from known resistance donors into the susceptible genetic background.
- Novel resistance-conferring haplotypes
- Already cloned *Yr* genes

## Outlook: Resistance gene candidates validation





## https://wheat-dw.ipk-gatersleben.de/



Test account
Username: demo
Password: demo

For registration please contact Stefanie Lück (lueck@ipk-gatersleben.de)

## Thank you very much!

#### Julius Kühn Institute

Ulrike Beukert Frank Ordon Andreas Stahl Albrecht Serfling

#### **University of Hohenheim**

C. Friedrich H. Longin

#### **IPK MPE**

Bihal Takar Nicolaus von Wirén

## **PLANT 2030**

#### **KWS Lochow**

Monika Spiller Nina Pfeiffer Sonja Kollers

#### Limagrain

Philipp Boeven Johannes Schacht

#### GFPi/ProWeizen

Tanja Gerjets

#### **IPK GGK**

Marion Röder

### Plant Breeding Research for the Bioeconomy

#### **IPK GGR**

Nils Stein Sandip Kale

#### **IPK DG**

Max Haupt Martin Mascher

#### **IPK BIT**

Stefanie Lück Sebastian Beier Uwe Scholz



#### **IPK BIM**

Ruben Betz

#### **IPK QG**

Albert W. Schulthess Valentin Hinterberger Jelena Perovic Christoph Martin Johannes Schneider Jochen C. Reif

GEFÖRDERT VOM

