#### SCIENCE MEETS LIFE

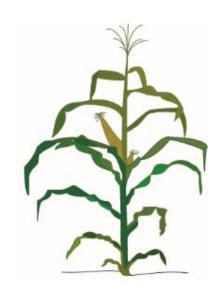
### **Rainbow Crops**

Hilde Nelissen





#### Maize plants are extremely suited for genetics



... and there is a lot of knowledge on genetic variation

#### Plants grow continuously

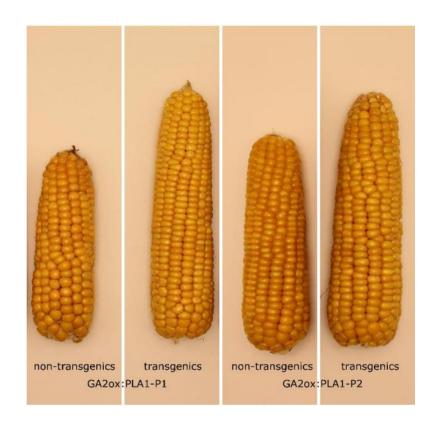


... and that is what interests our research team, Plant Growth Dynamics

# Maize growth processes: yield and architecture: multigenic traits



Transgenic Non-transgenic



Ectopic expression of *PLASTOCHRON1* (GA2OX::PLA1): increased leaf and stem biomass and seed yield



GA20- control GA
OXIDASE bios
OE mu

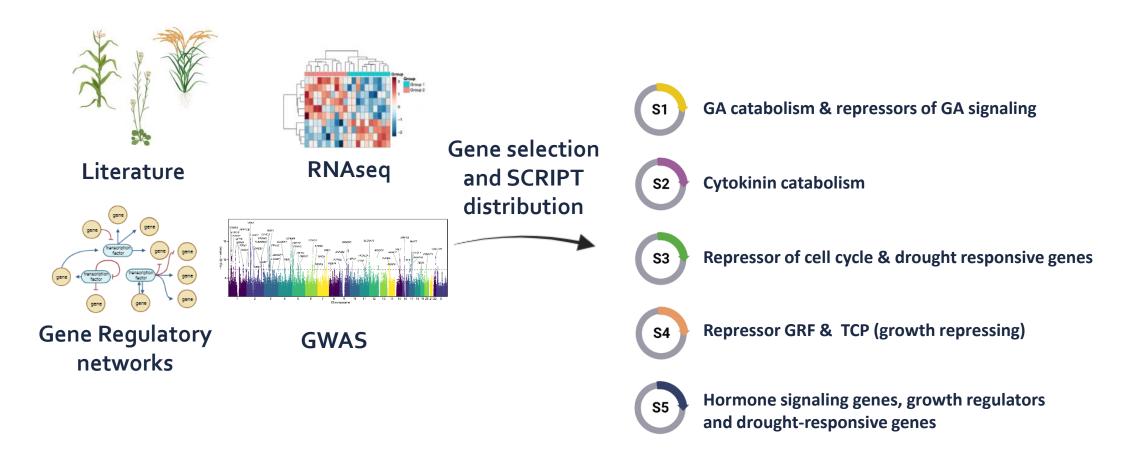
GA biosynthesis mutant

#### BREEDIT = Breeding + editing

- Most traits (including growth) are multigenic traits: thus variations at different locations are needed
- ▶ Redundancy
- Gene regulatory networks
- Linked loci

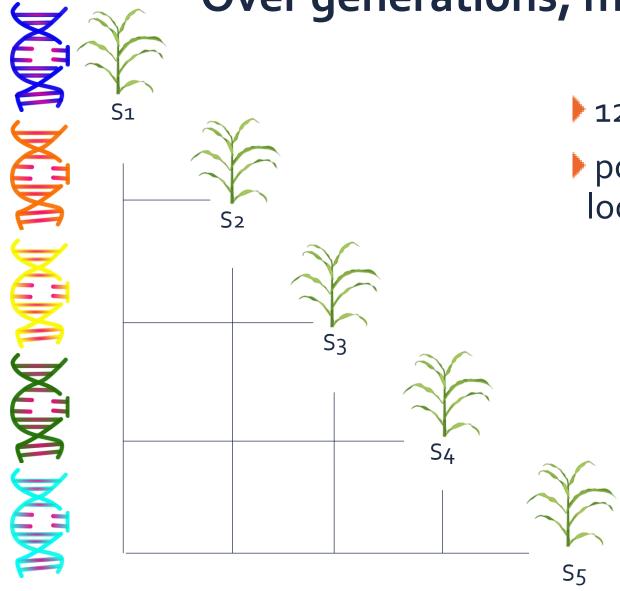
Growth as an example for multiplex genome editing

#### Data driven gene selection of negative growth regulators



## CAS<sub>9</sub> BREEDIT = Breeding + edit **12**g S1 Generation of populations in which different alleles of 12 **12**g loci segregate S<sub>2</sub> **12**g S3 **12**g

#### Over generations, more variation



- 12, 24, 36, etc...
- populations of different alleles of 60 loci segregating



Up to 60 gene edits segregating

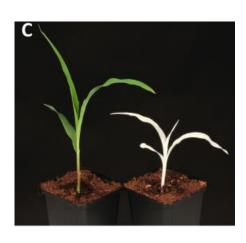
# Expected, unexpected but interesting and undesired outcome



Increased growth



**Compact plants** 

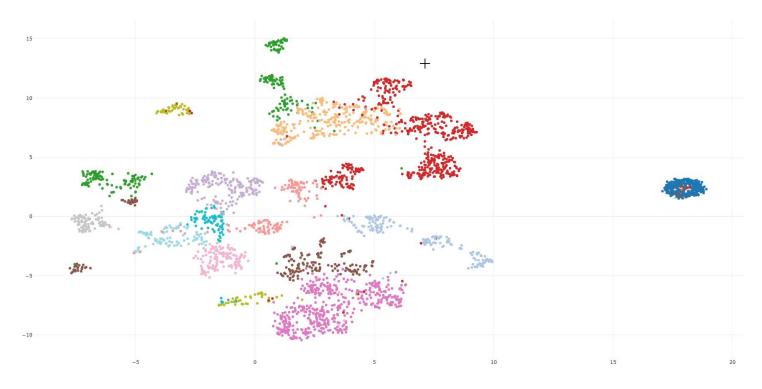


Albino plants

- The BREEDIT strategy shows which edits to combine, or which combinations are detrimental
- but how to identify the underlying edits?

#### Populations of individuals that are genetically unique

All plants are phenotyped and genotyped by multiplex amplicon sequencing

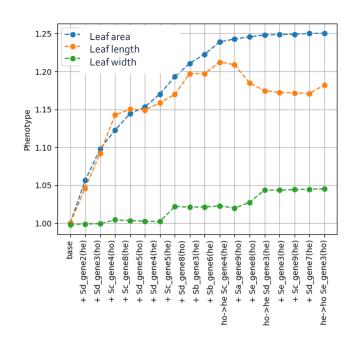




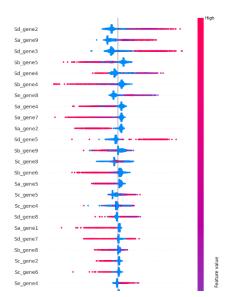
#### Train a model on genotype – phenotype data



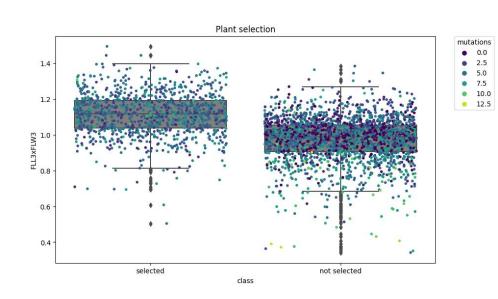
In silico trait design



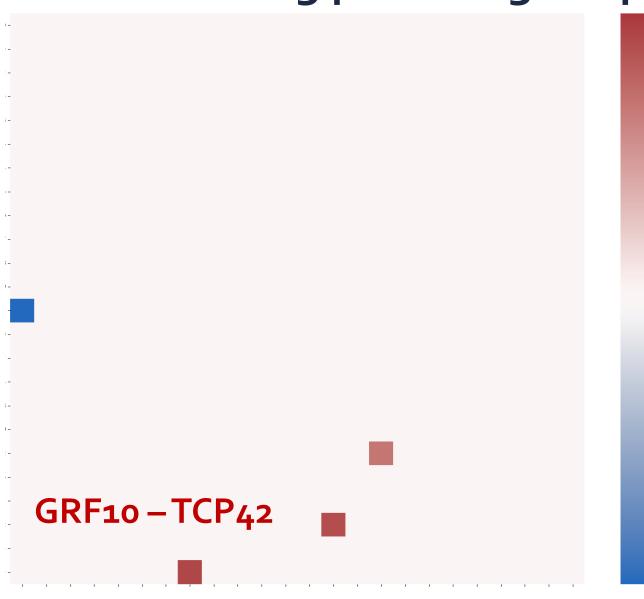
Al-based **gene** selection



Al-based **plant** selection

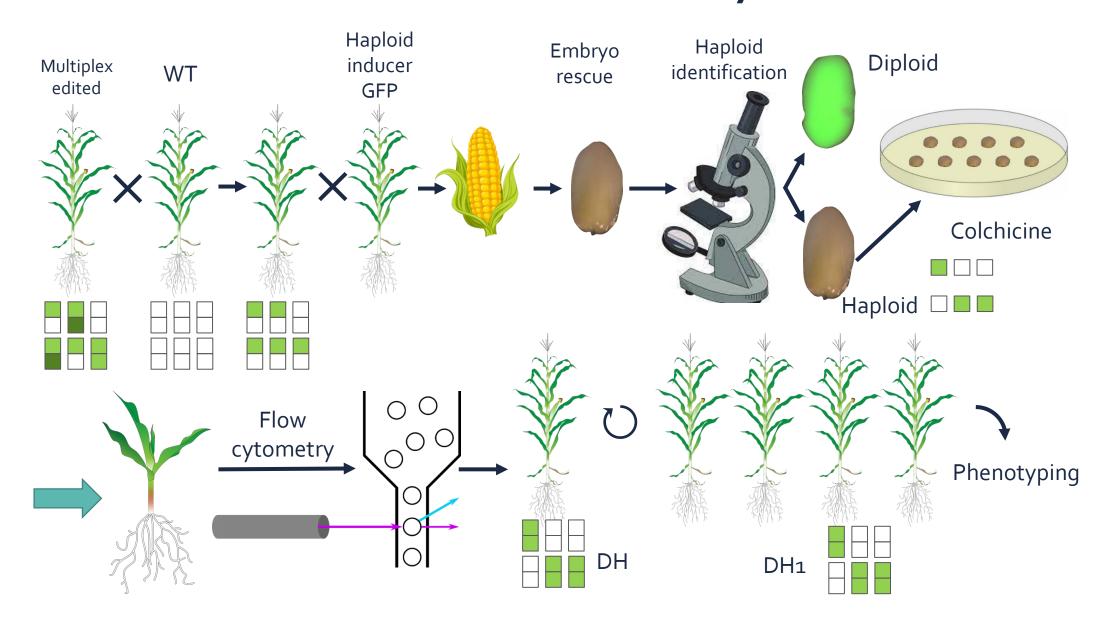


#### Al-based modeling predicts gene pairs

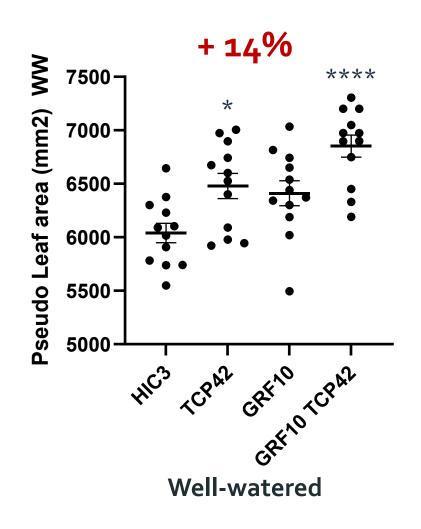


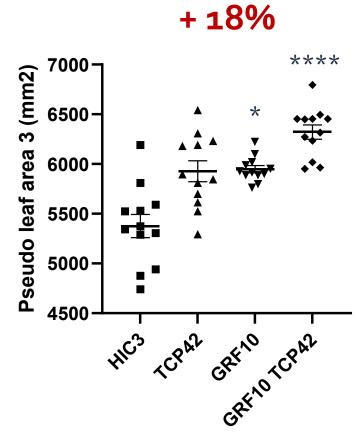
Gene pairs that are major contributors to leaf size, for example **GRF10-TCP42** 

#### Stabilize mutations for statistical analysis

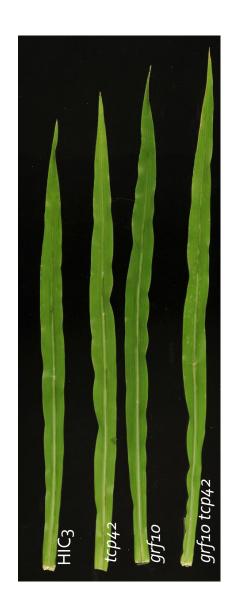


#### Haploid induction to fix the mutations

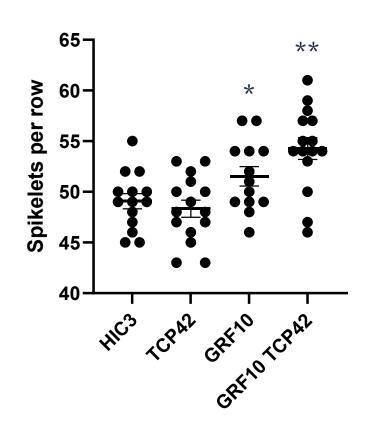


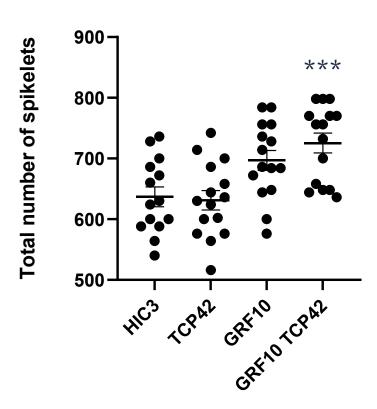


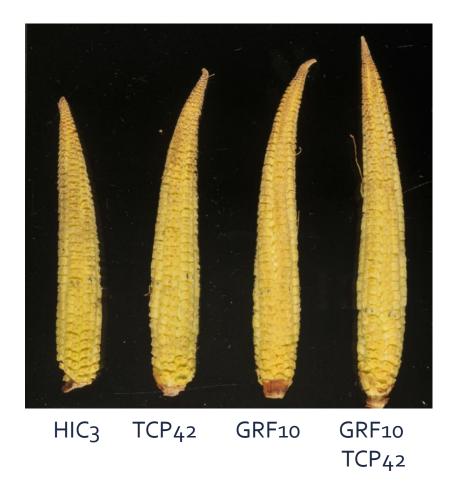




#### Mature plants with increased seed yield potential



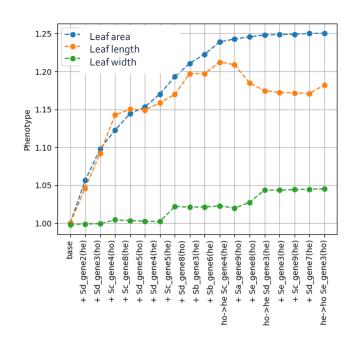




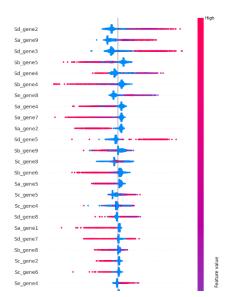
#### Train a model on genotype – phenotype data



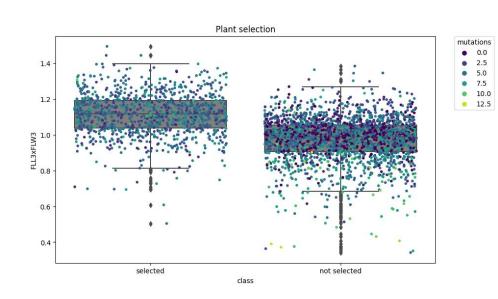
In silico trait design



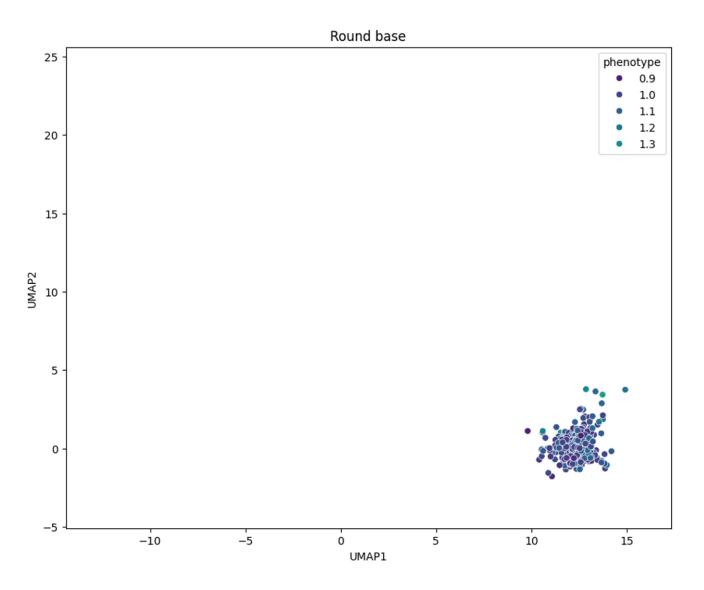
Al-based **gene** selection



Al-based **plant** selection

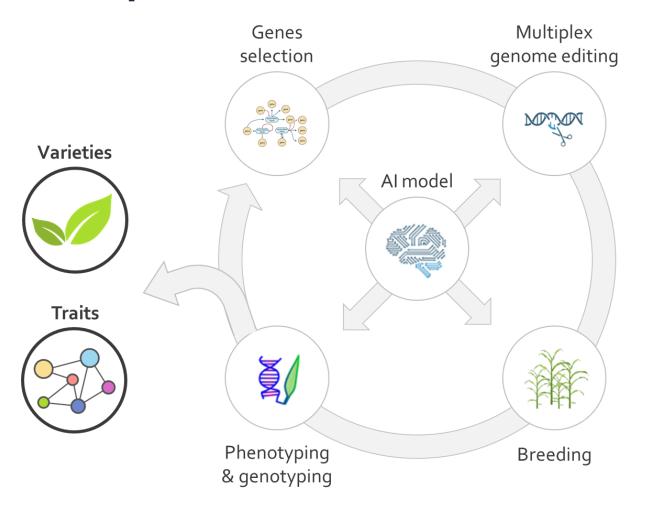


### From exploratory to exploitation: AI-guided breeding



#### From fundamental research to spin-off

- Strong support from VIB Innovation&Business and New Venture teams
- Initial round of funding: launch in April 2025
- CEO: Giacomo Bastianelli
- "Engineering complex traits to develop resilient crops"



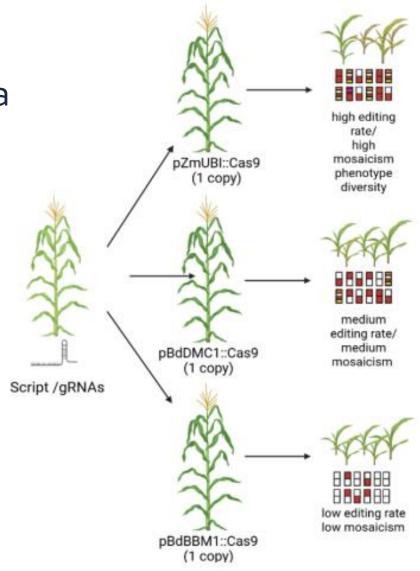


#### **Continued developments**

▶ Select different nucleases: Cas9 and Cas12a

Drive expression of the nucleases by different promoters

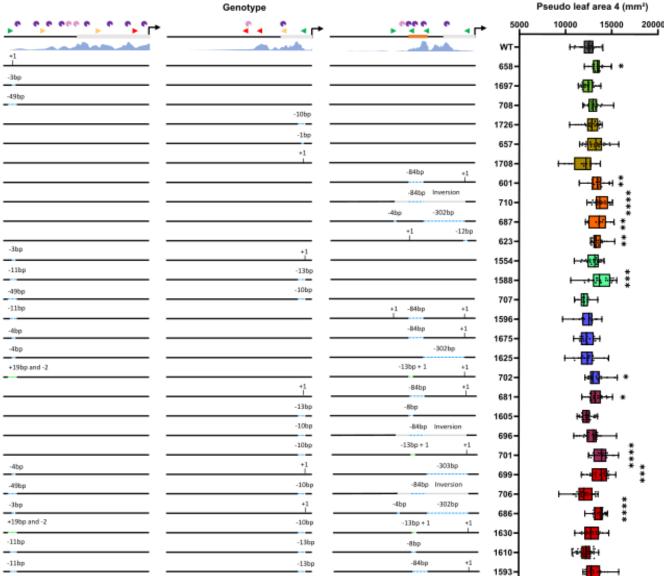
▶ Target regulatory sequences



Editing of regulatory sequences

Finetuned expression instead of constitutive knock-out

(semi-)dominant



Relative expression

## Acknowledgements



engineering complex traits

















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Stijn Seynnaeve

Giacomo Bastianelli

#### **Collaborators:**



Thomas
Jacobs



Laurens Tom Pauwels Ruttink



**Agro-Incubator** 



Plant growth dynamics group



Tom Viaene









