



High throughput phenotyping to improve genetic gain in wheat – prospects and challenges

March 9th 2023

PD Dr. Andreas Hund and Olivia Zumsteg

Outline

1 Research site and high throughput phenotyping

1.1 Drone

1.2 FIP

2 Projects

2.1 Caterra

2.2 ClimBreed

2.3 Genomic Predictions

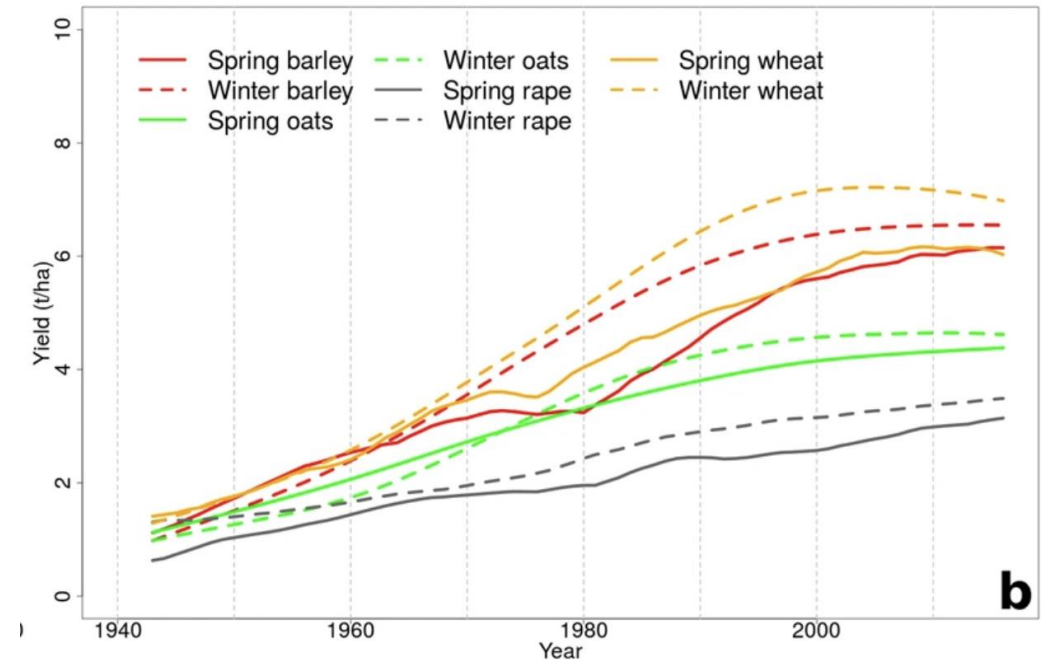
Motivation

Stagnating wheat yield

Increasing global food demand and climate change impacts

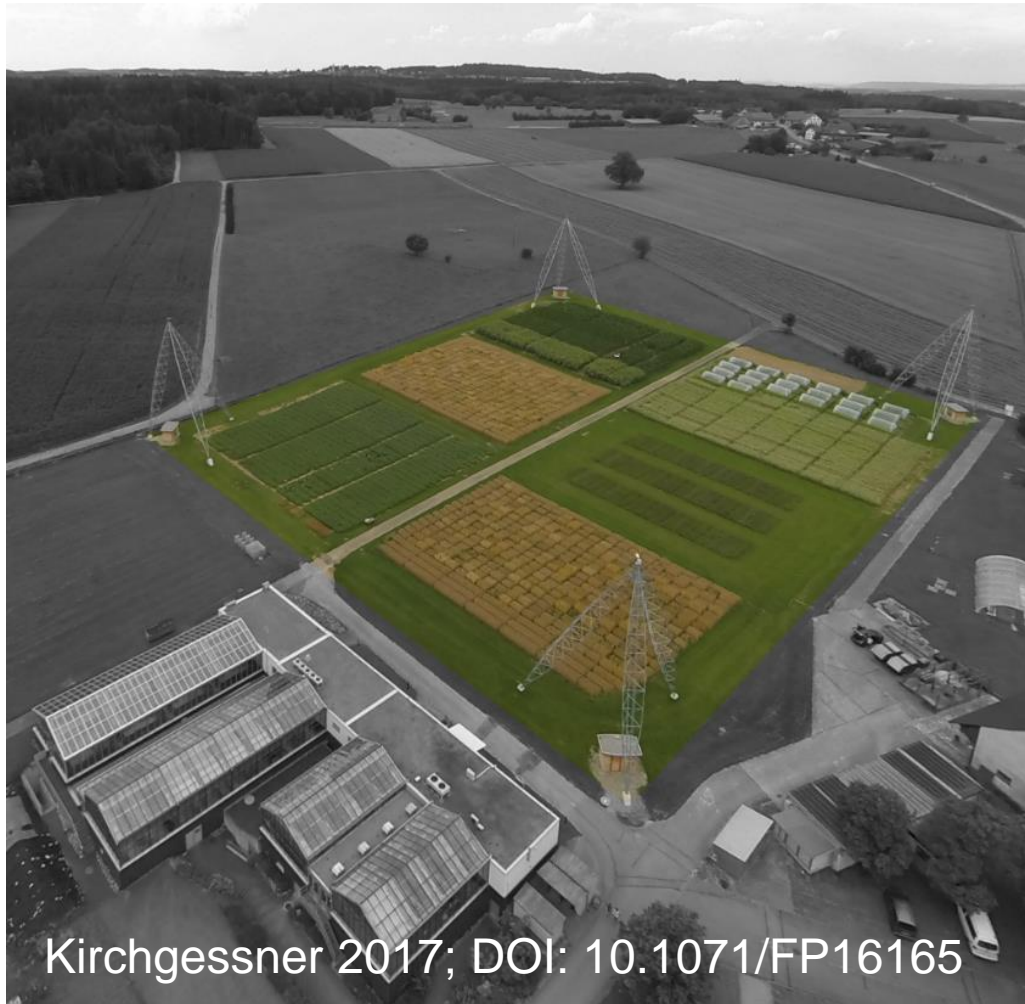
Phenotypic and genomic prediction methods could help to cope with this challenge

Stagnating wheat yield in France



(Bernhard Schauburger et al. (2018). Yield trends, variability and stagnation analysis of major crops in France over more than a century)

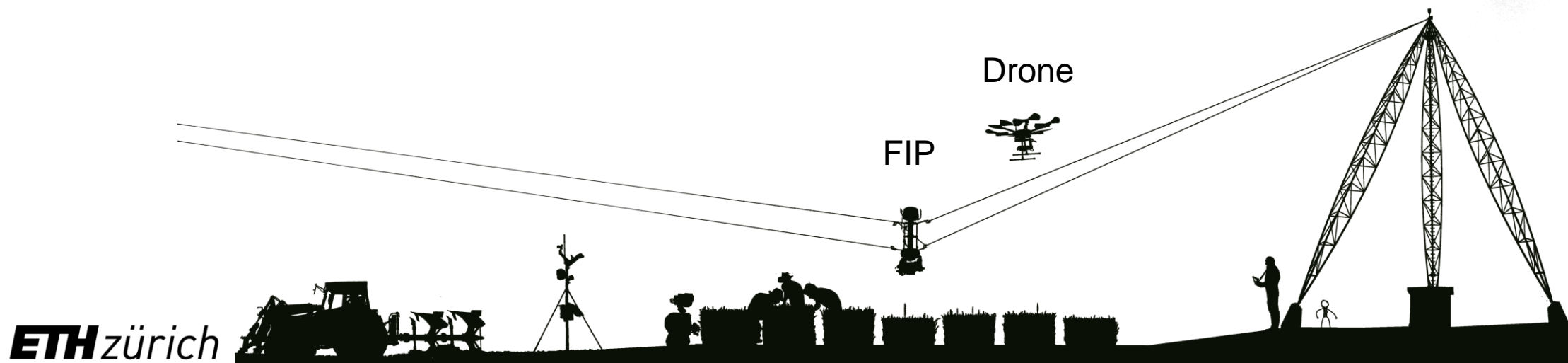
1) Research Site and High Throughput Phenotyping



Hight Throughput Phenotyping (HTP)



Possibility to assess crop growth under realistic conditions with high temporal and spatial resolution



HTP: Drone



Derived Traits from Drone

Height related traits

Final plant height

Start stem elongation

Stop stem elongation

Temperature response parameters:

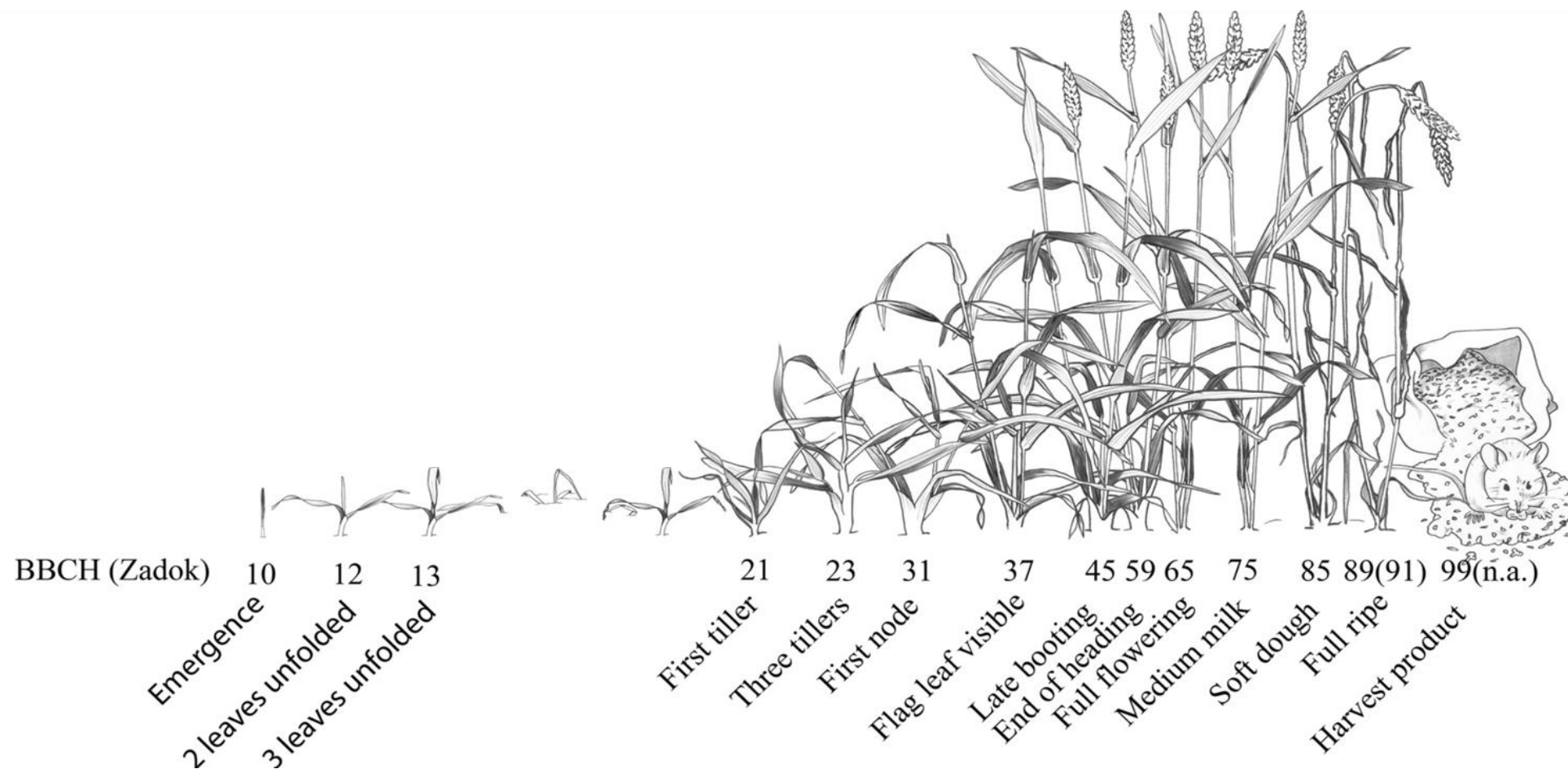
Maximum growth rate

Steepness ofn response

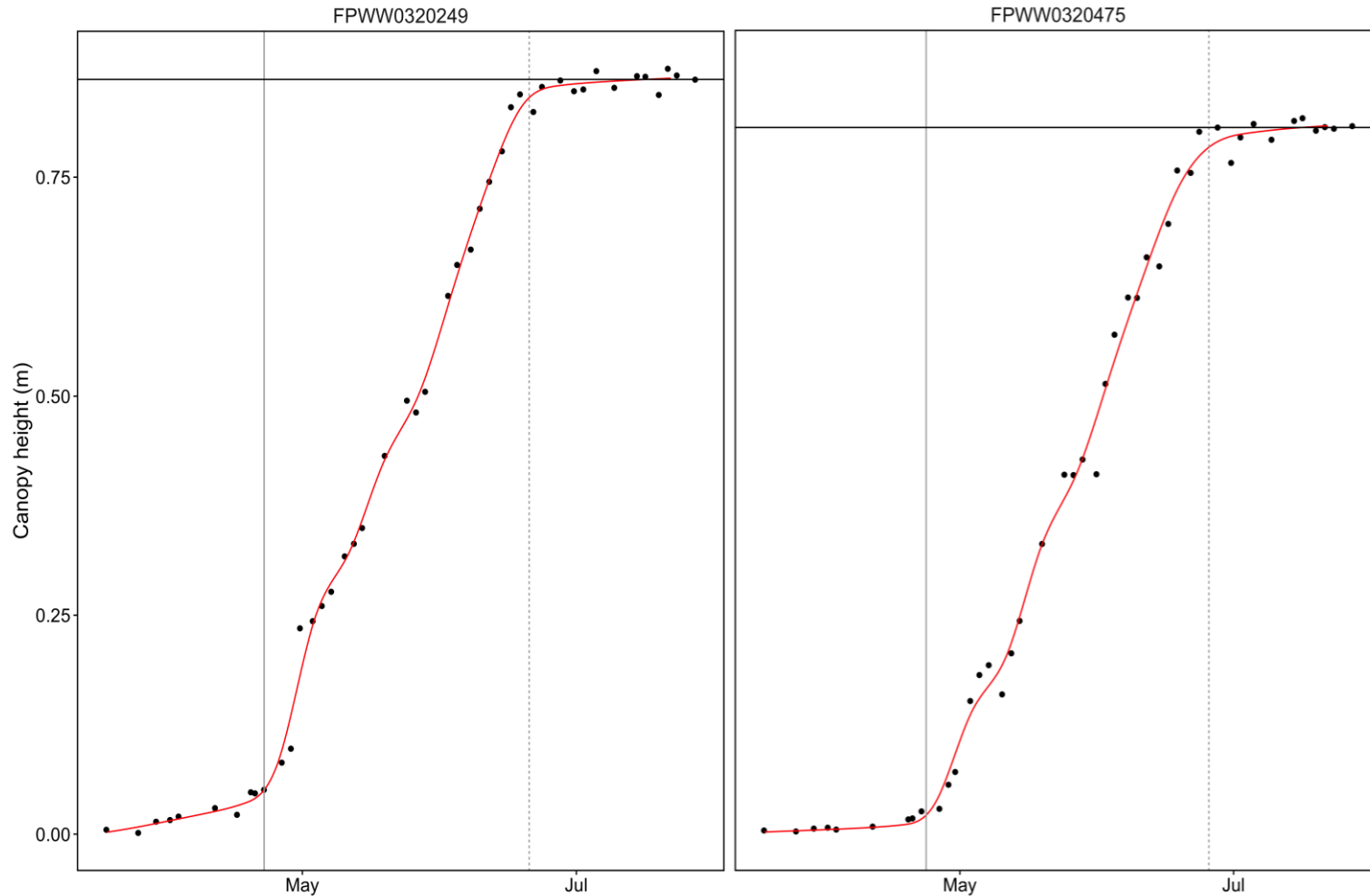
Base Temperature



Crop development: Height



Extraction of Height and Temp. Related Traits

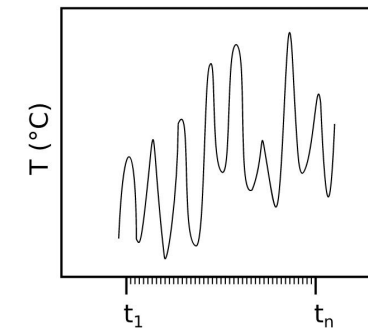


1) Height Related Traits:

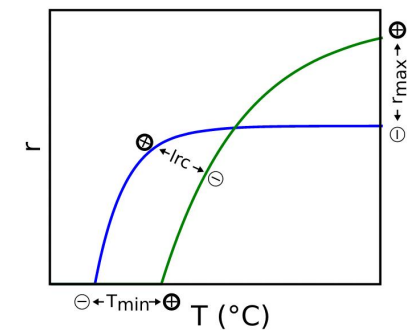
- final height (PH_{max})
- | start growth phase ($t_{PH_{start}}$)
- ⋯ stop growth phase ($t_{PH_{stop}}$)

2) Temp. Related Traits:

(b) Covariate measurement



(c) Dose-response curve



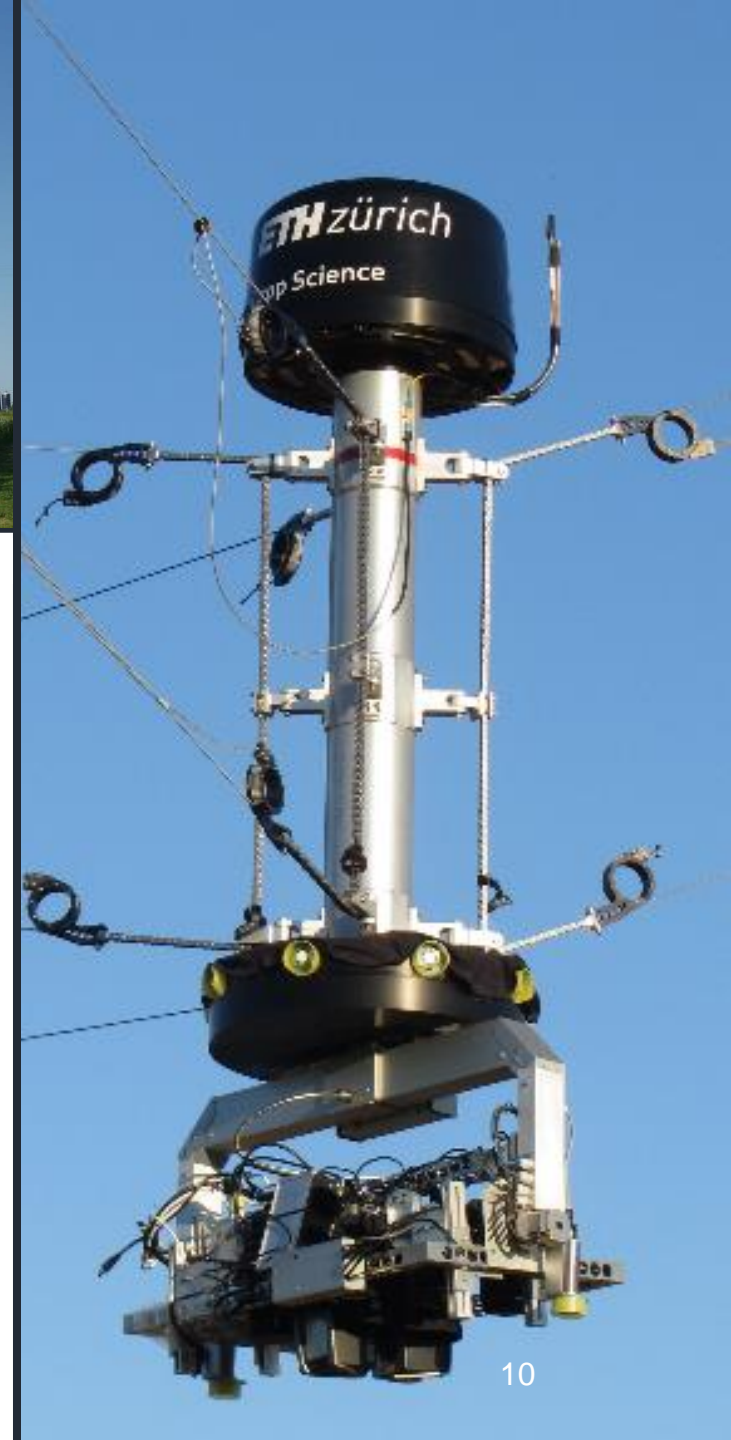
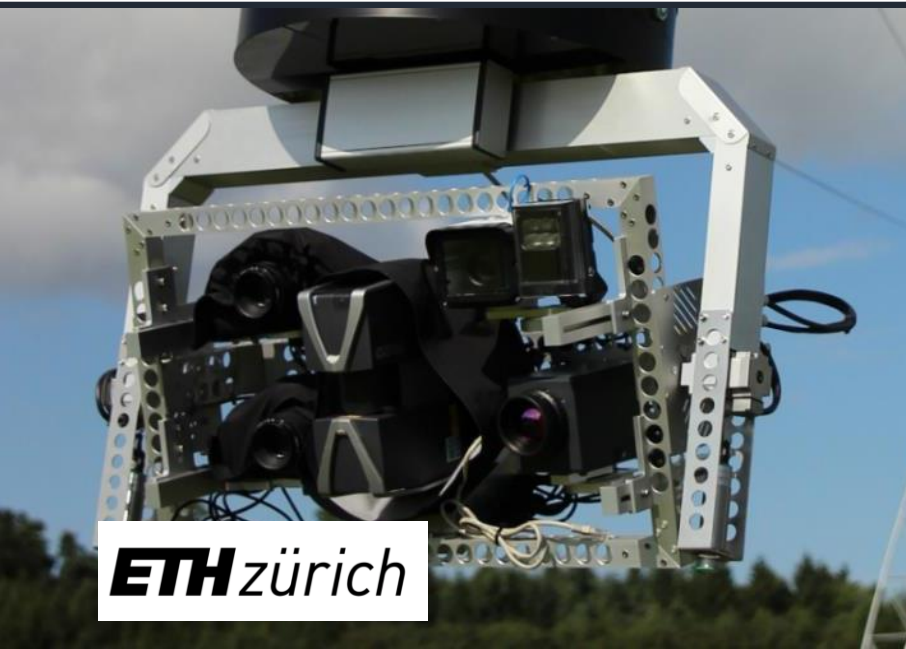
HTFP: Field phenotyping platform (FIP)



- 4 poles of 24 m height
- 4 lots
- Operating on regular basis
- 3 m distance to the ground

Sensor:

- High-resolution RGB images



Derived Traits from FIP

Height related traits



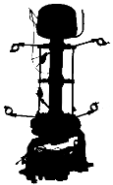
Final plant height

Start stem elongation

Stop stem elongation

Temperature response parameters

Canopy cover related traits



Time point when apparent leaf area increase is maximized

Maximum apparent leaf area after canopy closure

Leaf area at 15 % final height

Time of canopy closure



HTFP: Field phenotyping platform (FIP)

The new FIP sensor head: Bringing multi-view imaging into the field

13 machine-vision camera triggered synchronously



Additional Traits to Establish

Height related traits



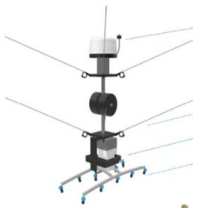
- Final plant height
- Start stem elongation
- Stop stem elongation
- Temperature response parameters

Canopy cover related traits



- Time point when apparent leaf area increase is maximized
- Maximal apparent leaf area after canopy closure
- Leaf area at 15 % final height
- Time of canopy closure

Traits to establish



Number of plants / tillers and spike volume

Plant height and biomass

Architecture

ClimBreed

Outline

1 Research site and high throughput phenotyping

1.1 Drone

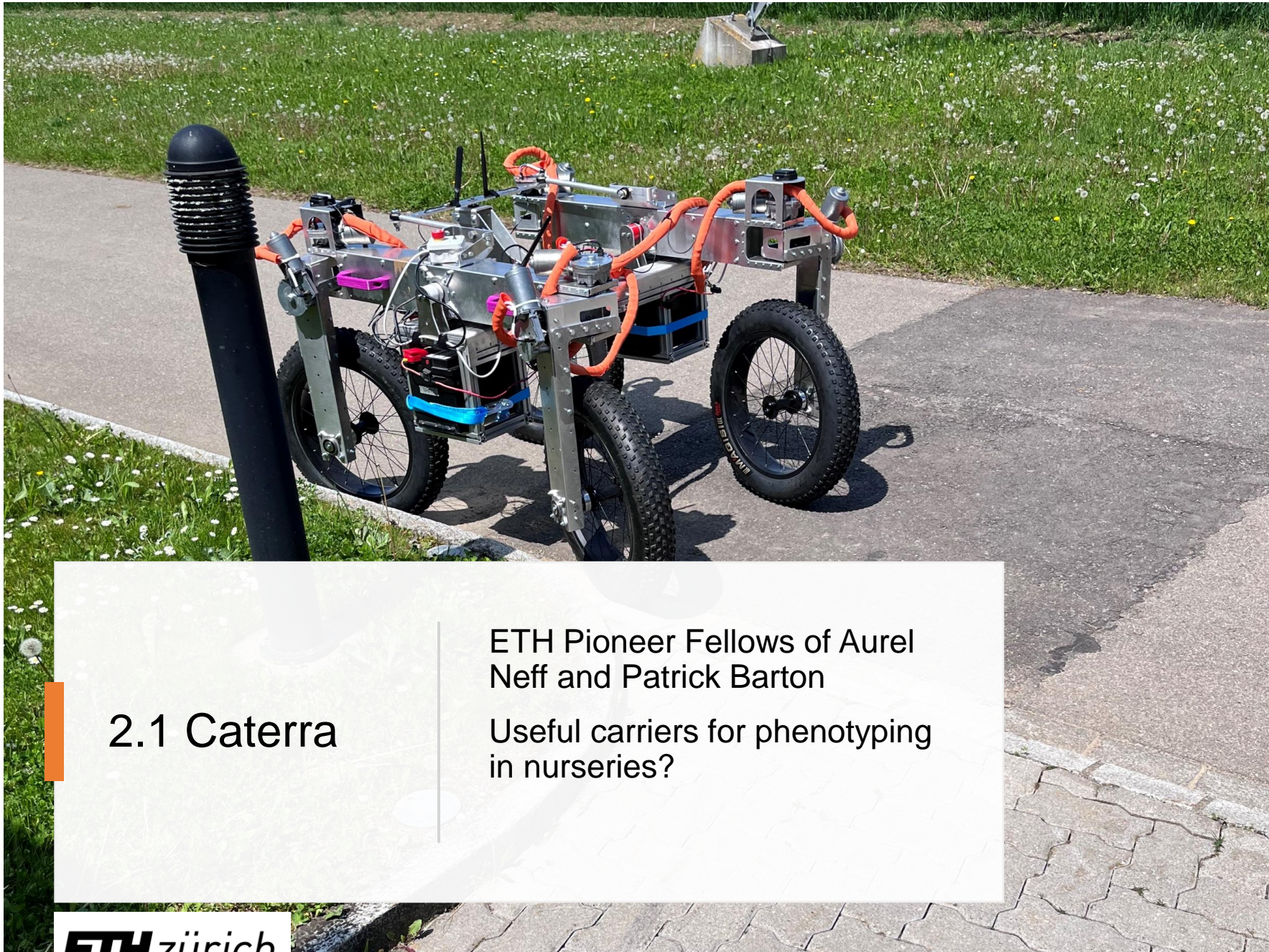
1.2 FIP

2 Projects

2.1 Caterra

2.2 ClimBreed

2.3 Genomic Predictions



2.1 Cattera

ETH Pioneer Fellows of Aurel Neff and Patrick Barton
Useful carriers for phenotyping in nurseries?



Laser-based weeding robot

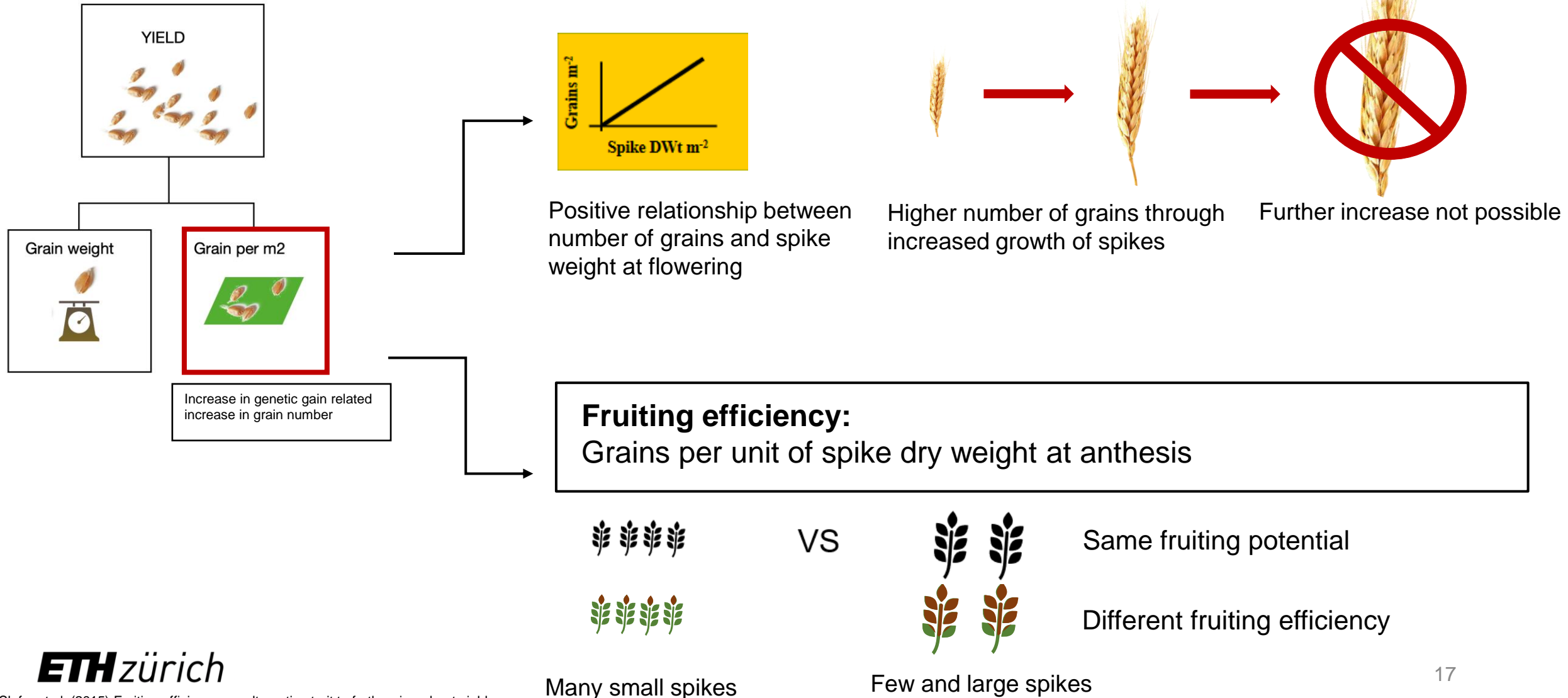


Stefan G. with Literal device

2.2 ClimBreed

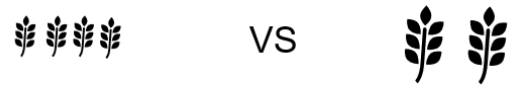


WP1: Estimate Fruiting Potential and Efficiency



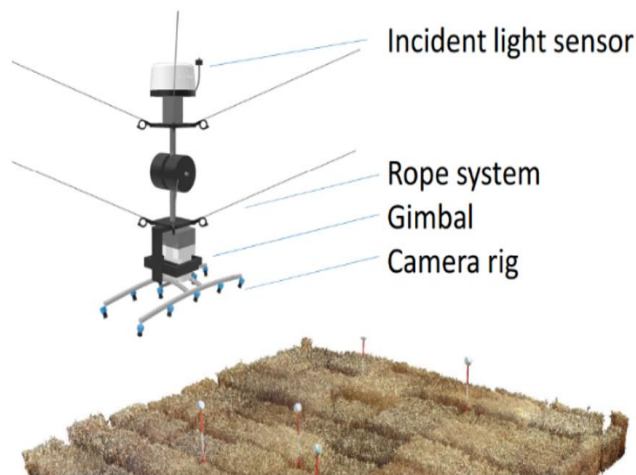
WP1: Measuring Fruiting Potential and Efficiency

Estimating fruiting potential:



Number and volume of spikes

Biggest challenge!

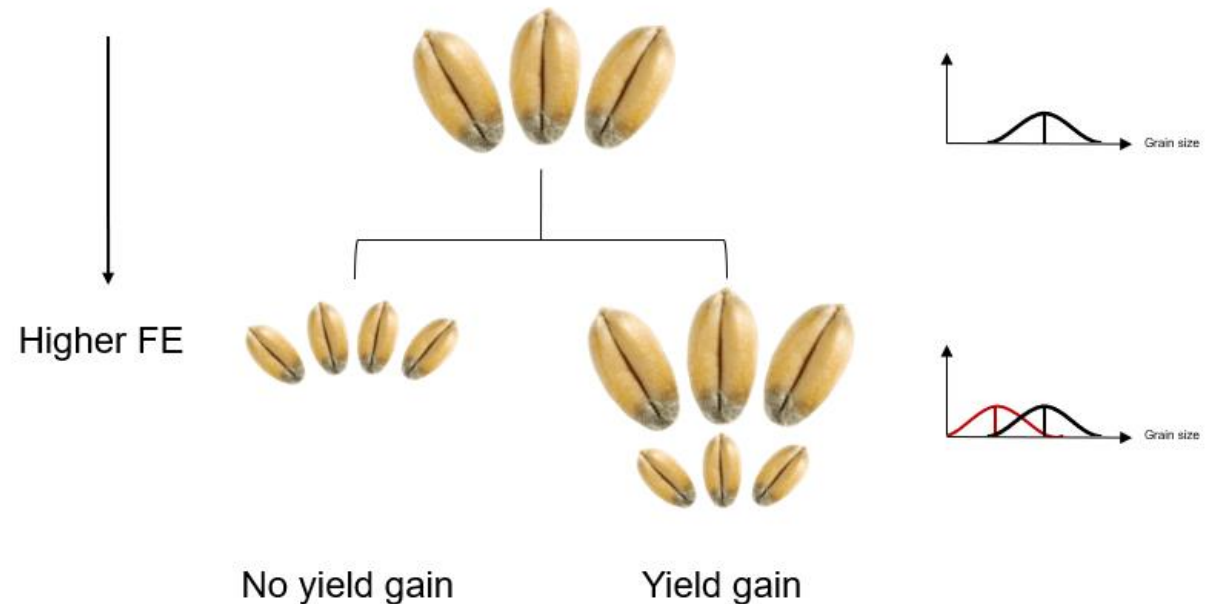


Estimating fruiting efficiency:



Grain number

Potential trade-off:

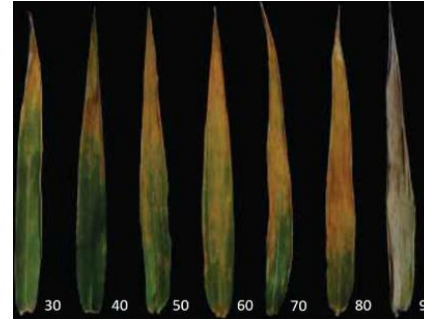


Protein content



<https://www.indiamart.com/proddetail/wheat-grains-19327420562.html>

Senescence



Chapmann et al. (2021) Capturing and Selecting Senescence Variation in Wheat

WP2: Traits related to Fruiting Efficiency

Heading



<https://site.extension.uga.edu/wilcoxcoag/2019/03/wheat-heading/>



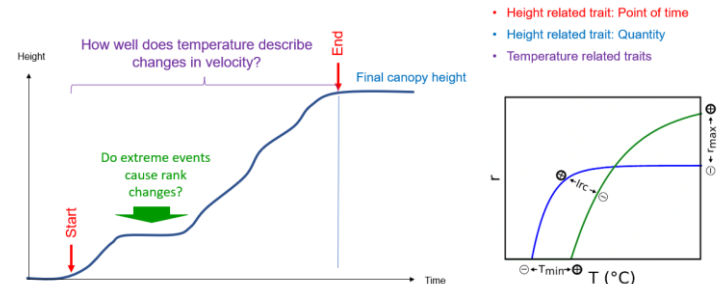
<https://www.freepnglogos.com/pics/wheat>

Grain size



<https://jasoncholewa.com/2013/02/08/early-humans-wheat-cultivation-and-your-health/>

Height and temperature related traits



Lukas Roth et al. (2021) Phenomics data processing: A plot-level model for repeated measurements to extract the timing of key stages and quantities at defined time points, visualized by Andi Hund.

WP3: How to Deliver New Traits to Breeders: Genomic Prediction

Which traits are heritable?

Index to predict low / high fruting efficiency

Select these genotypes and
confirm genomic prediction

Cultivation in France as a validation site

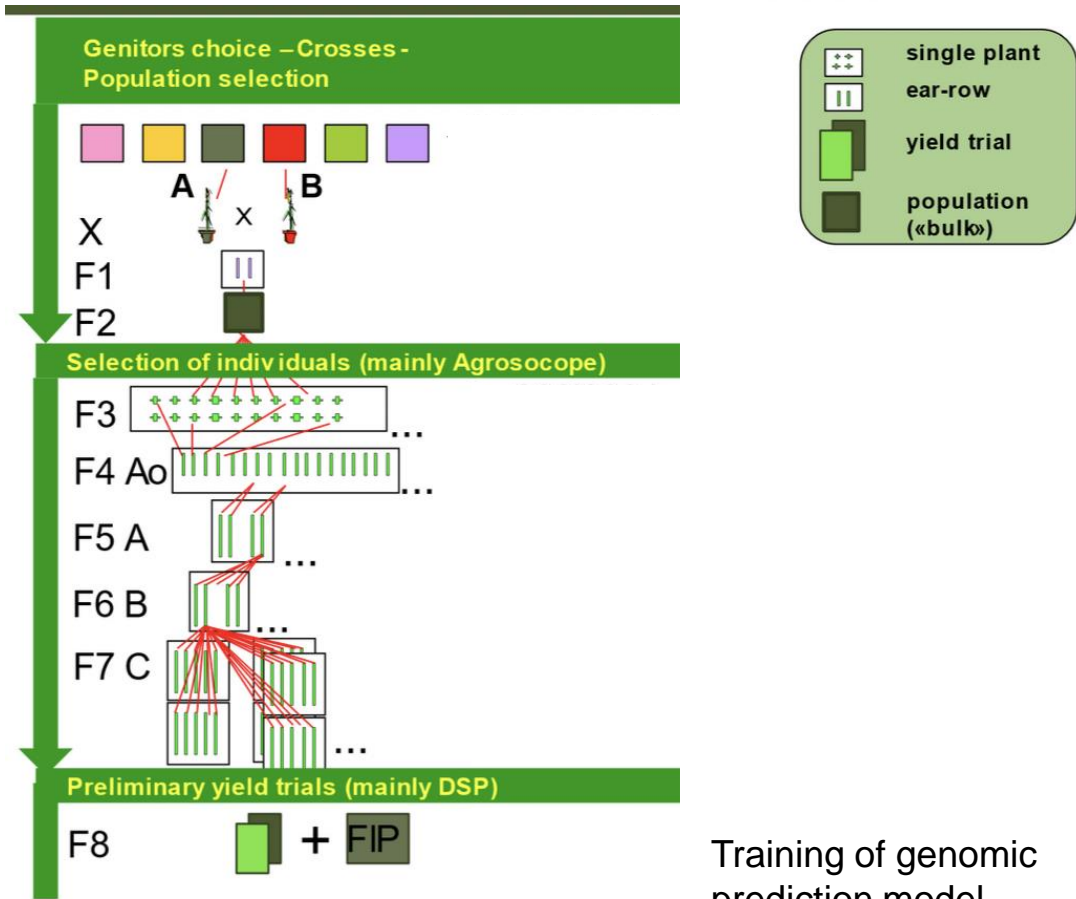


<https://www.freepnglogos.com/pics/wheat>



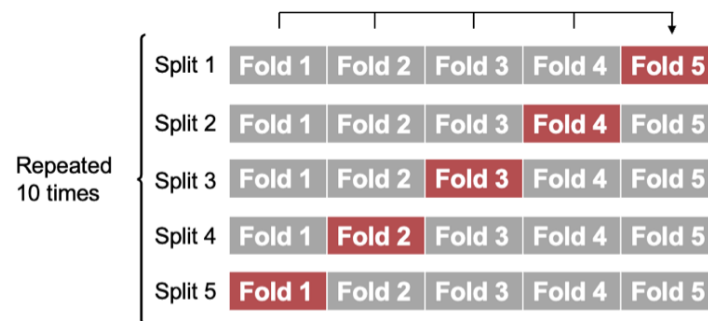
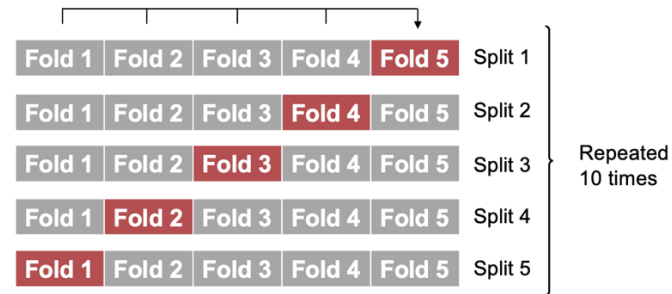
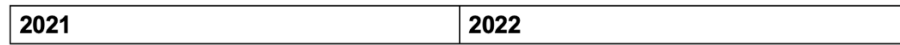
Future-climate site at *Gréoux-les-Bains*

2.3 Genomic Prediction

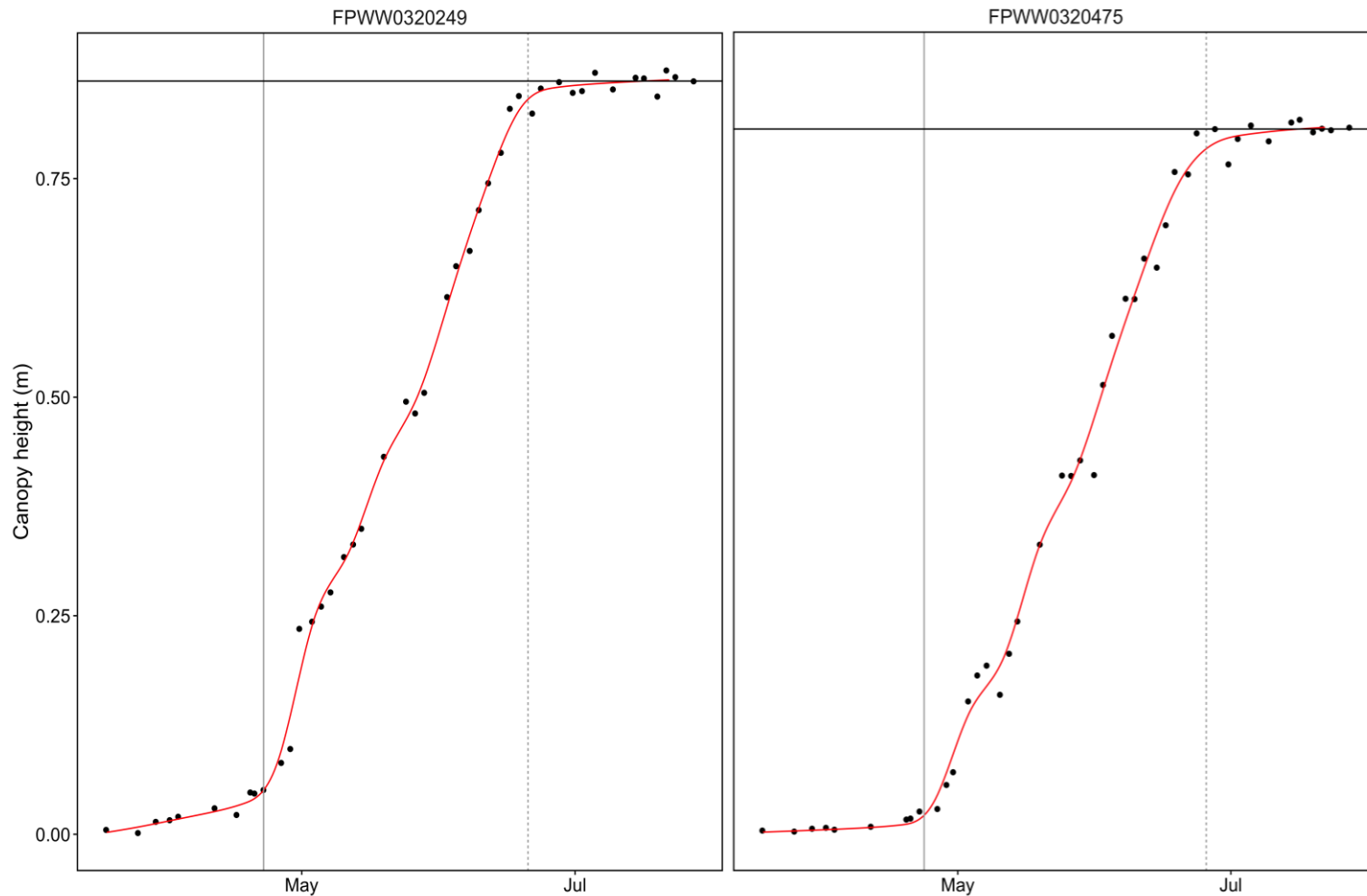


(Dario Fossati, Agroscope)

F8: Cross-Validation and Prediction of Unseen Years



Extraction of Height and Temp. Related Traits

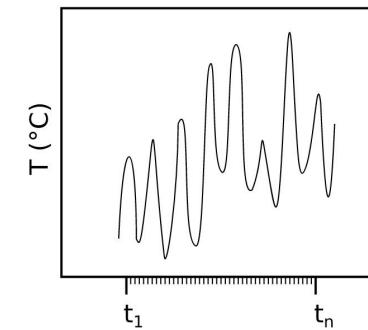


1) Height Related Traits:

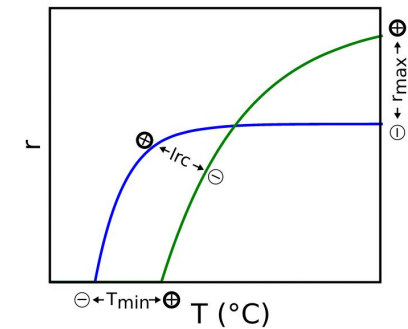
- final height (PH_{max})
- | start growth phase ($t_{PH_{start}}$)
- ⋯ stop growth phase ($t_{PH_{stop}}$)

2) Temp. Related Traits:

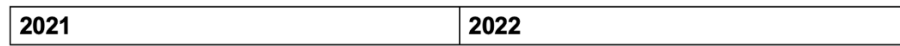
(b) Covariate measurement



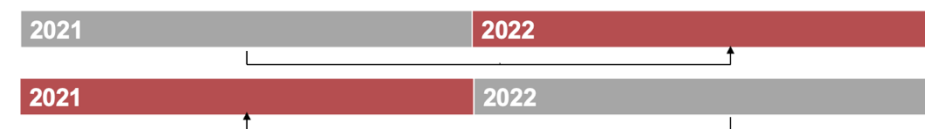
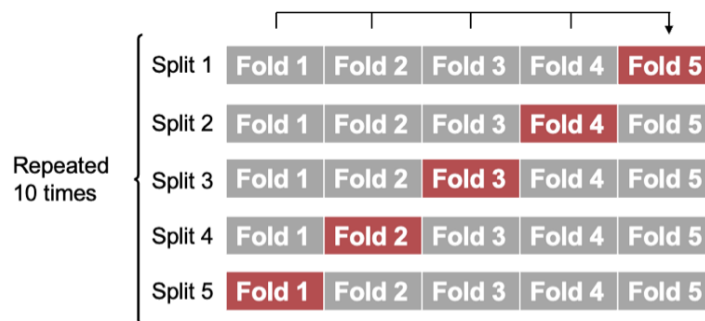
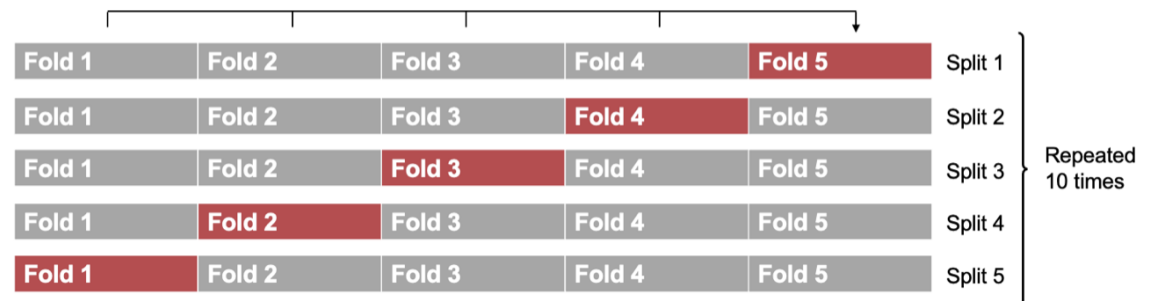
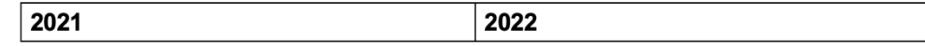
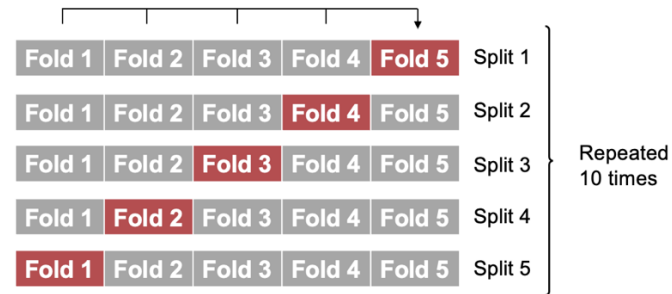
(c) Dose-response curve



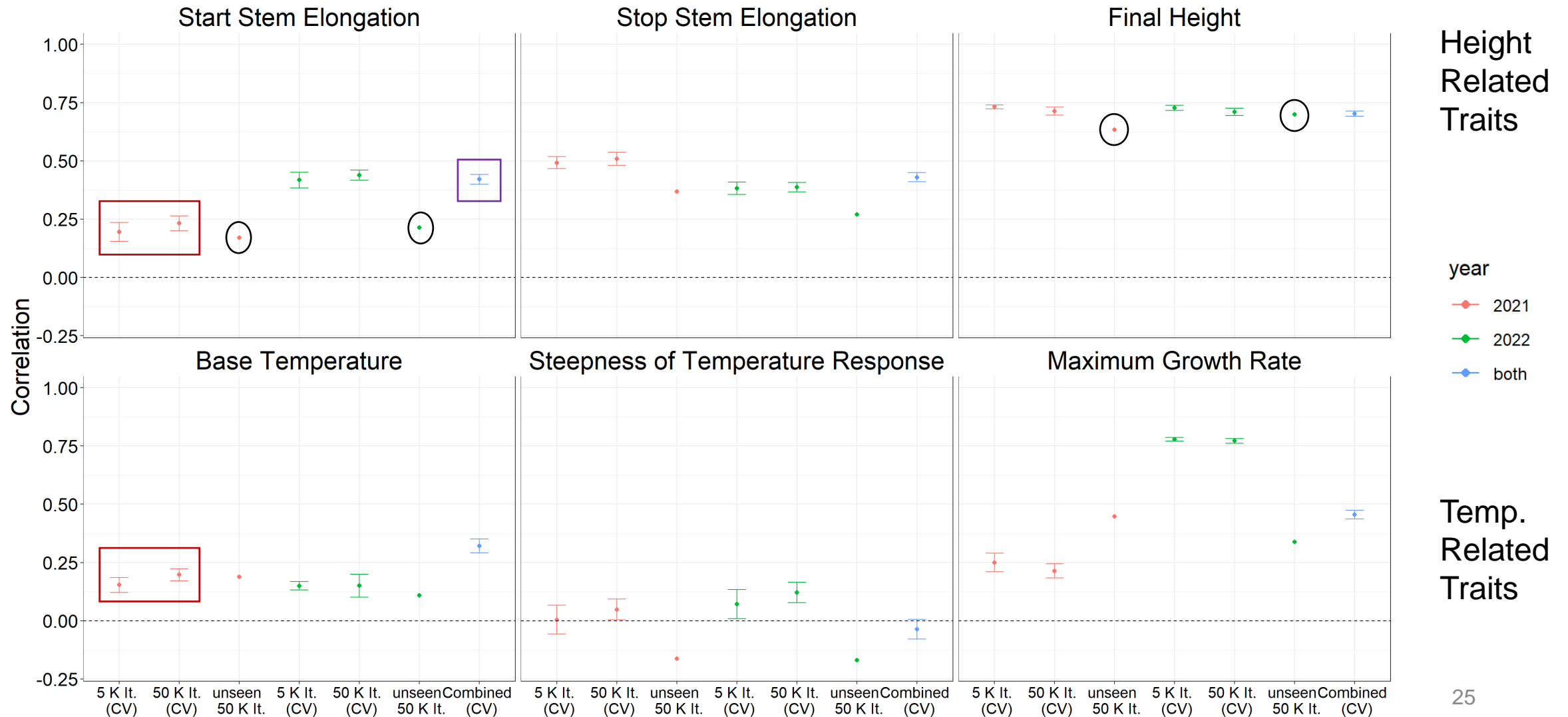
F8: Cross-Validation and Prediction of Unseen Years



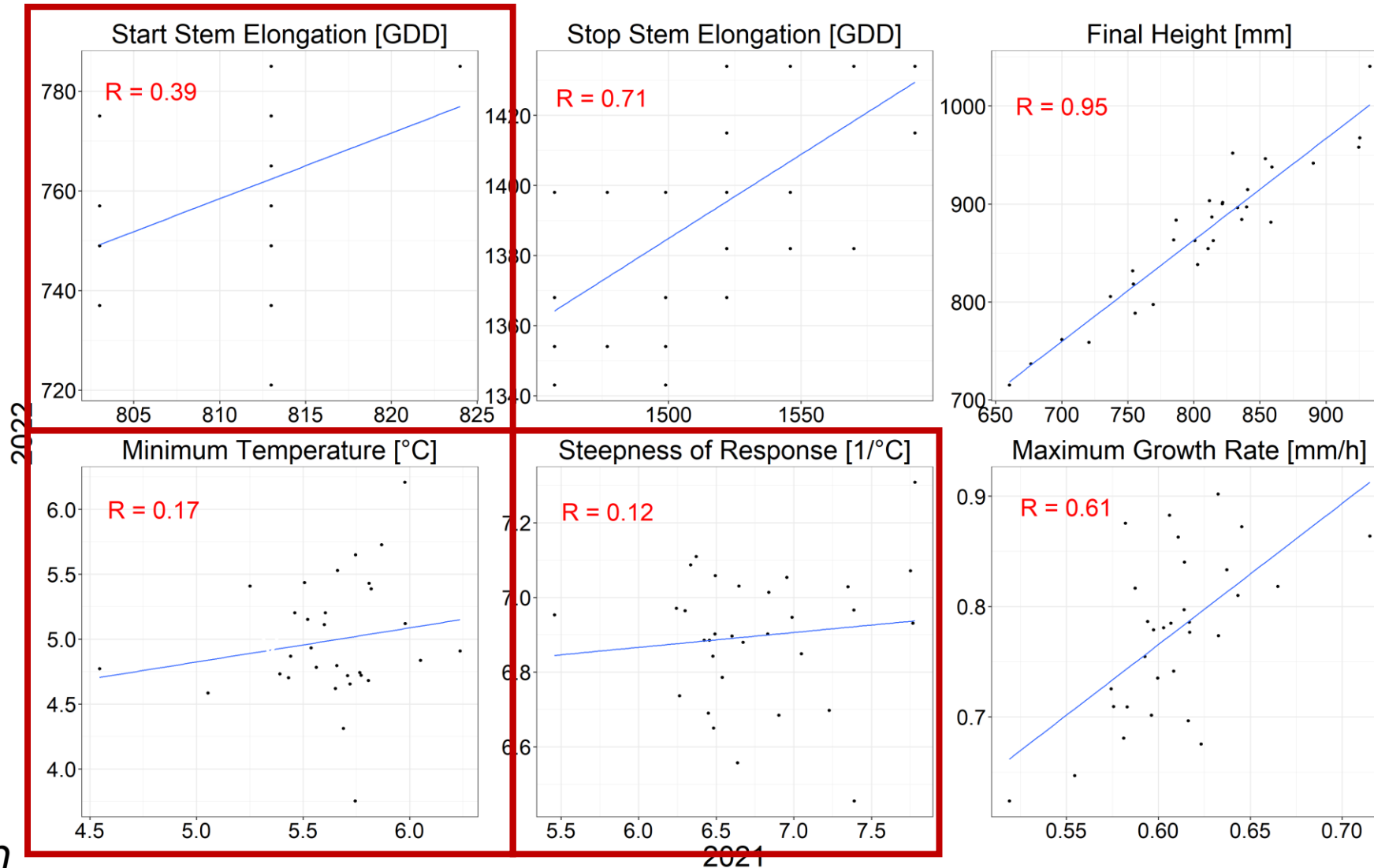
Training data Test data



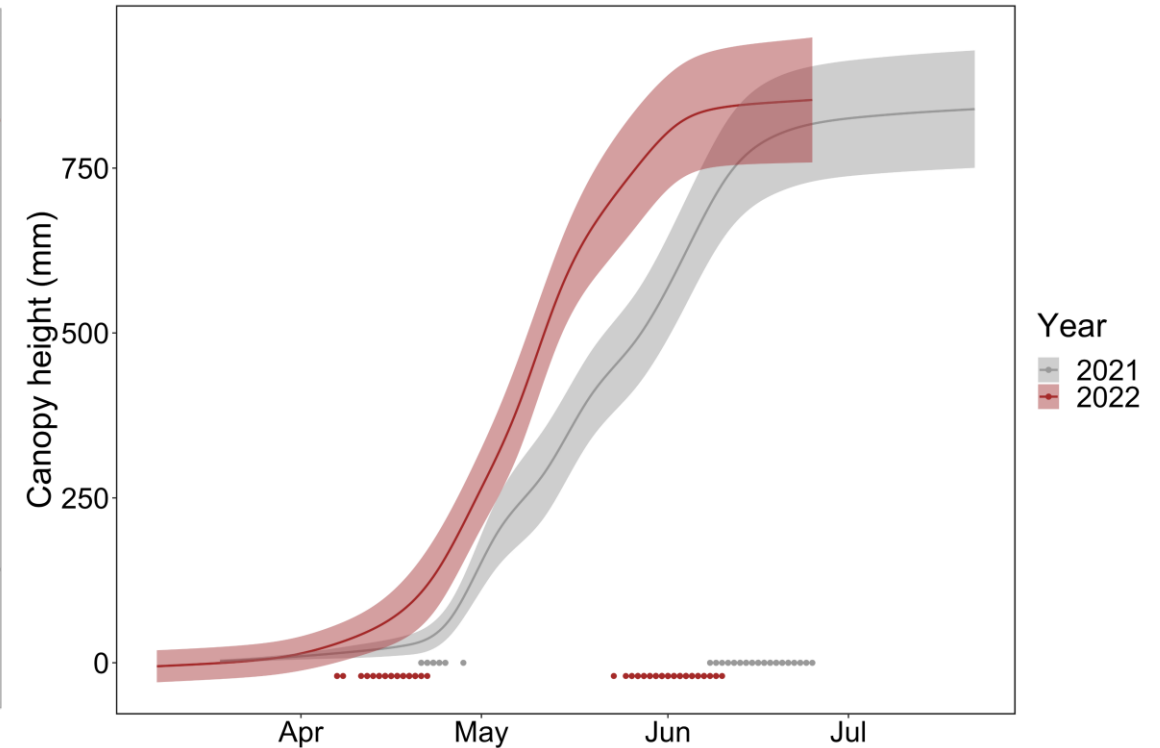
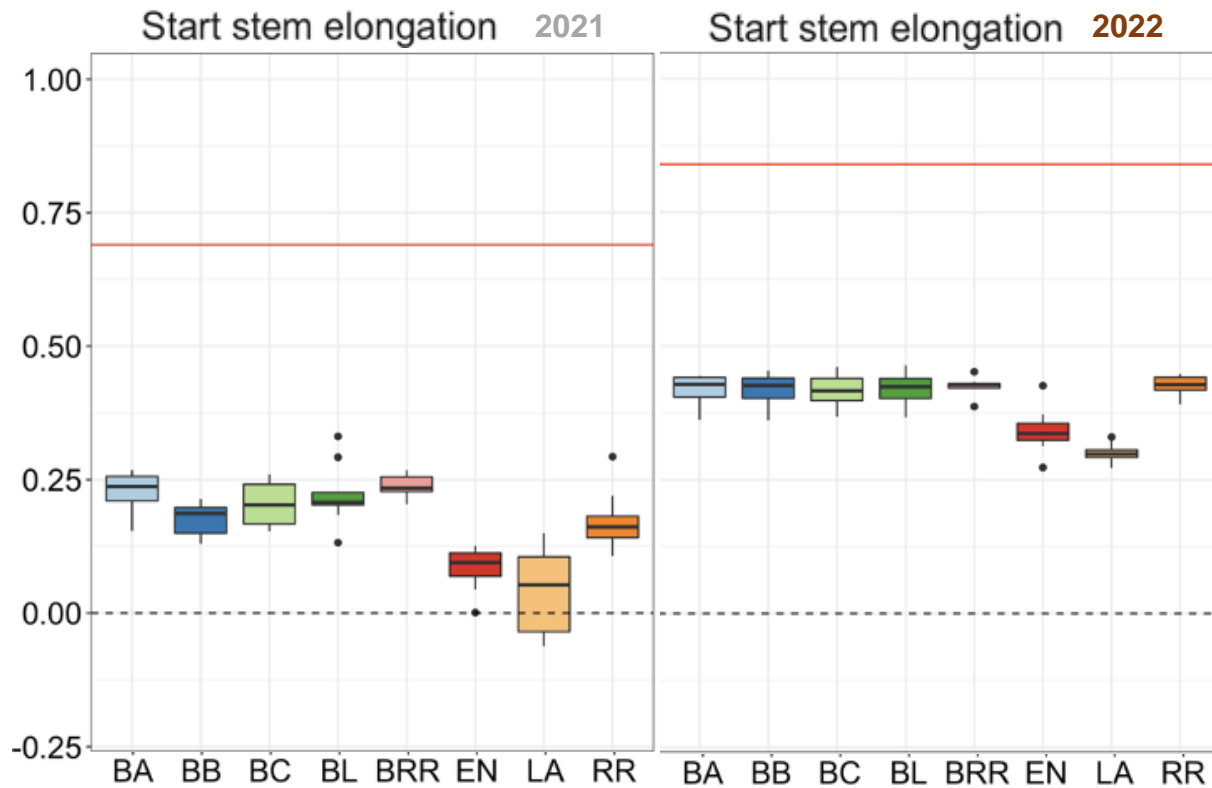
F8: CV and Prediction of Unseen Genotypes 2021 and 2022 with BayesB



F8: Correlation Between 2021 and 2022



F8: Results and Discussion: Start of Stem Elongation in 2021 and 2022



Conclusion

Aim: To use dense data of single season to gain as much information as possible about varieties with non-destructive methods

GP: Differences in prediction accuracies between traits and years

Improving trait extraction, modelling GxE interactions, or including environmental parameters could help to increase prediction accuracy

Genomic Prediction has the potential to speed up breeding process, e.g. with predicting start of stem elongation

Thank you!

Questions?
Feedback?

...



Data 2021 and 2022

Number of genotypes / in GP / number of checks

346, 373 / 254 , 242 / 4, 3

Plot size:

1.5 m x 2 m

Drone: 28 m, 2.3 m/s,

Markers: 25 k snp array: most common markers in European wheat varieties, and additional markers from QTL studies

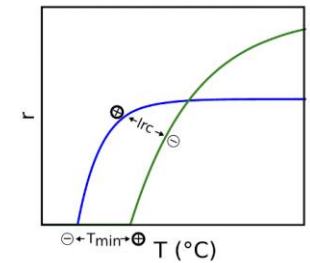
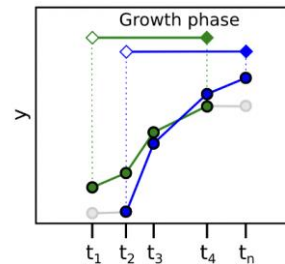
Heritability

Amount of phenotypic variance that can be explained by the genotypic variance

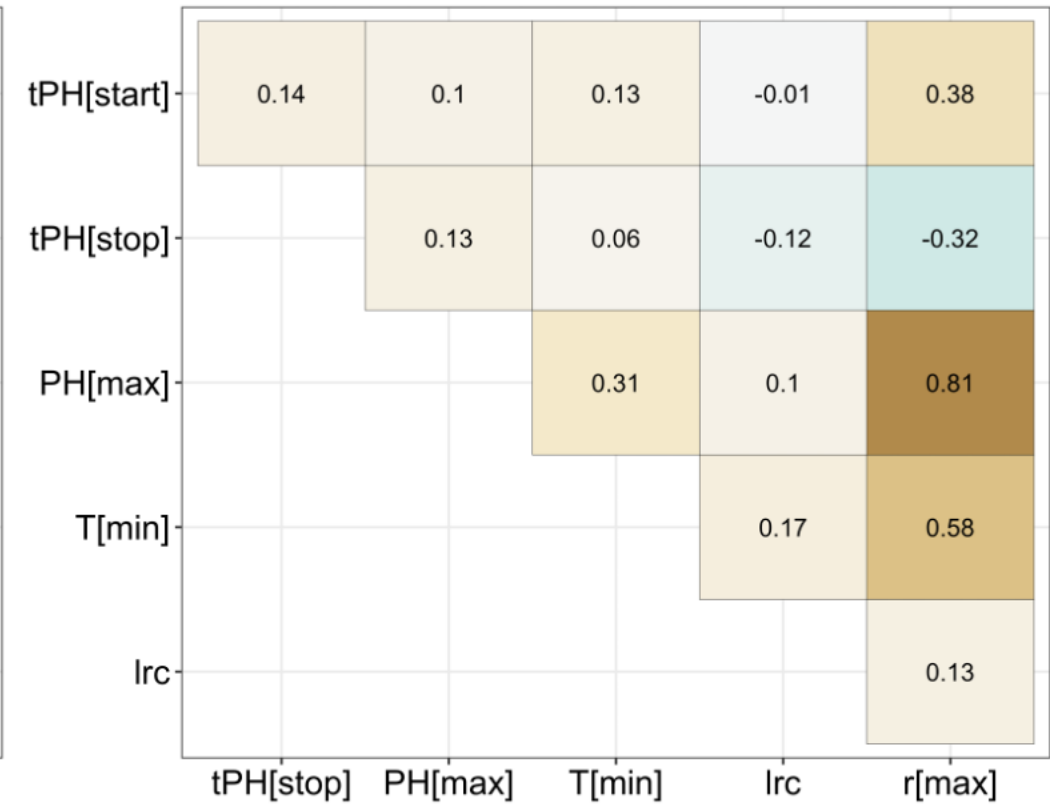
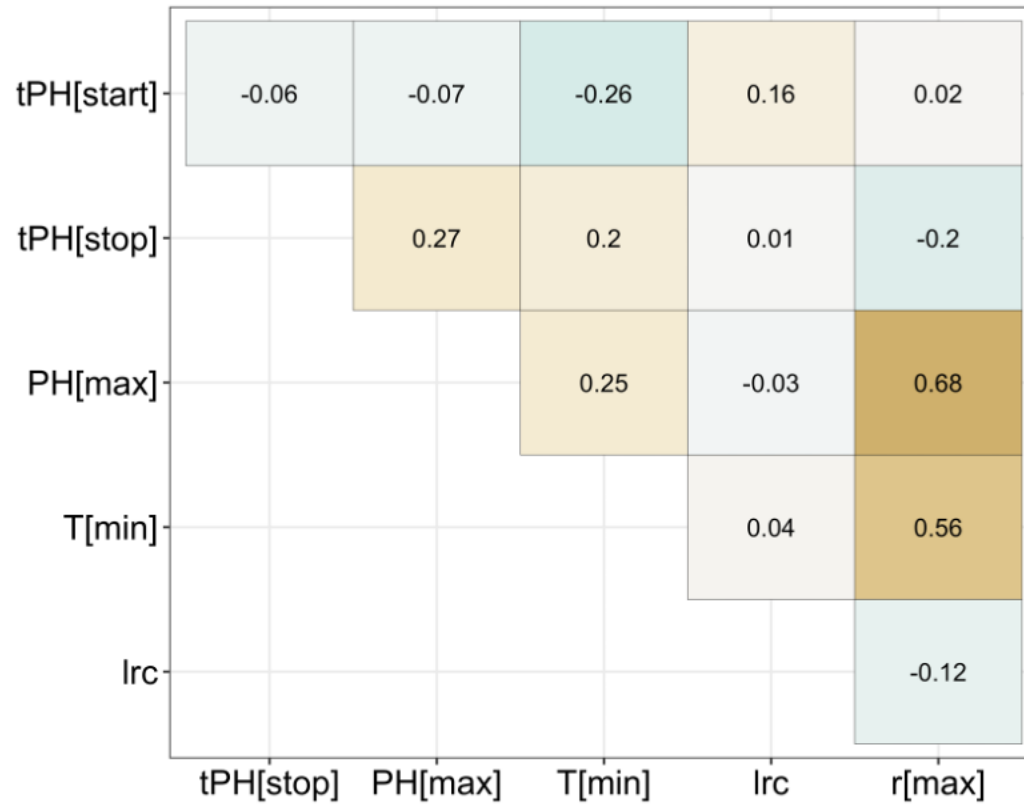
$$H^2 = \frac{\sigma_g^2}{\sigma_P^2}$$

$$R = \frac{i \sigma_G h}{L}$$

Height related traits		Temperature related traits	
Final canopy height	0.97	Maximal growth rate	0.43
End stem elongation	0.86	Minimal temperature	0.28
Start stem elongation	0.48	Steepness response	0.31



Genetic correlation F8



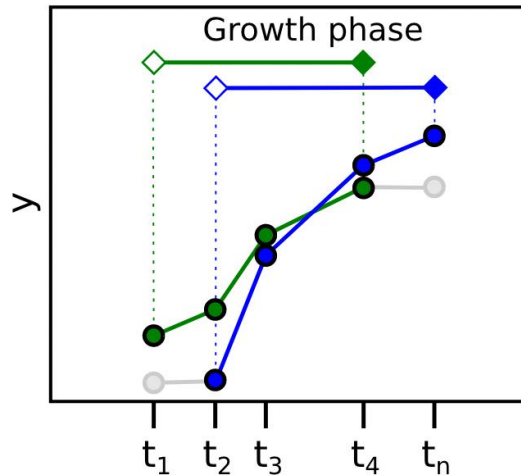
Modelling Temperature Response

$$r = r_{\max} * (1 - \exp(-\exp(s) * (T - T_{\min})))$$

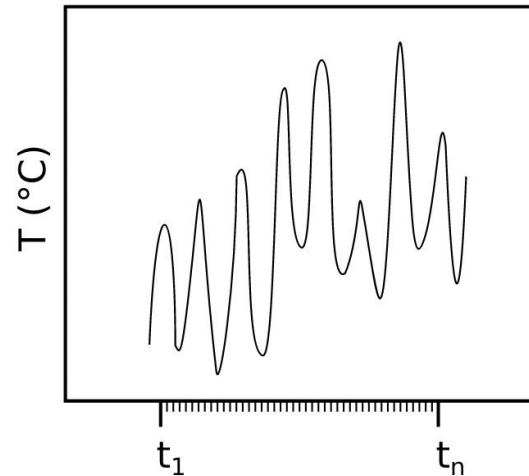
$$r_{\text{asym}}(T) = \begin{cases} r, & r > 0, \\ 0 & \text{otherwise} \end{cases}$$

$$y_t = \sum_{d=t_0}^t \left(\sum_{h=1}^{n_d} r(T_{dh}, \theta) * \Delta m_h \right),$$

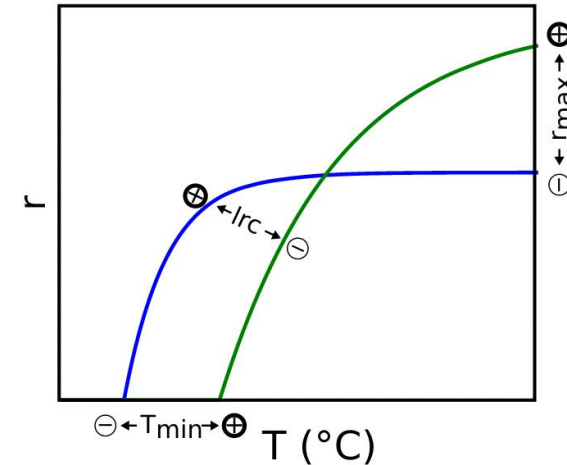
(a) Trait measurement



(b) Covariate measurement

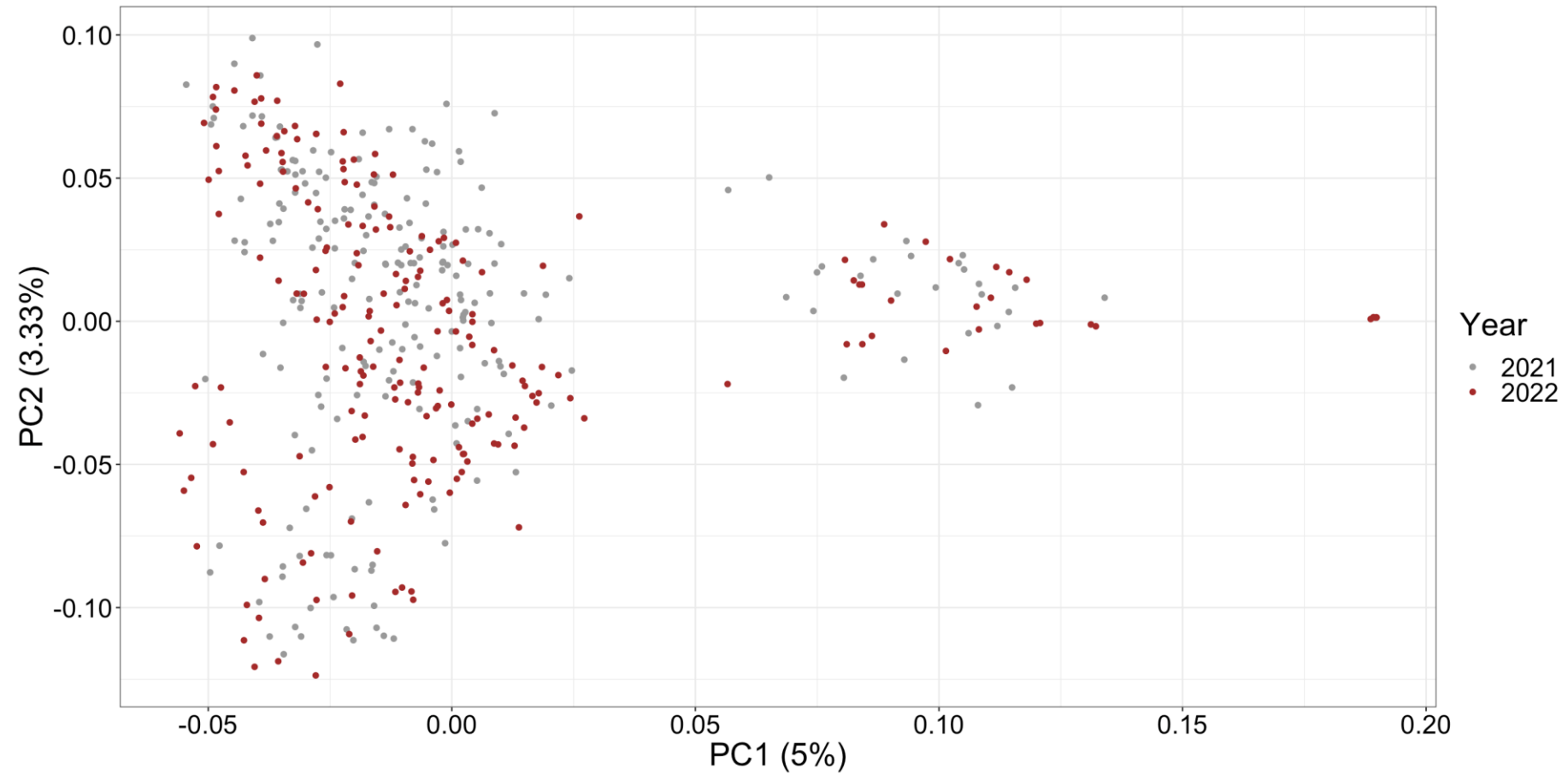


(c) Dose-response curve



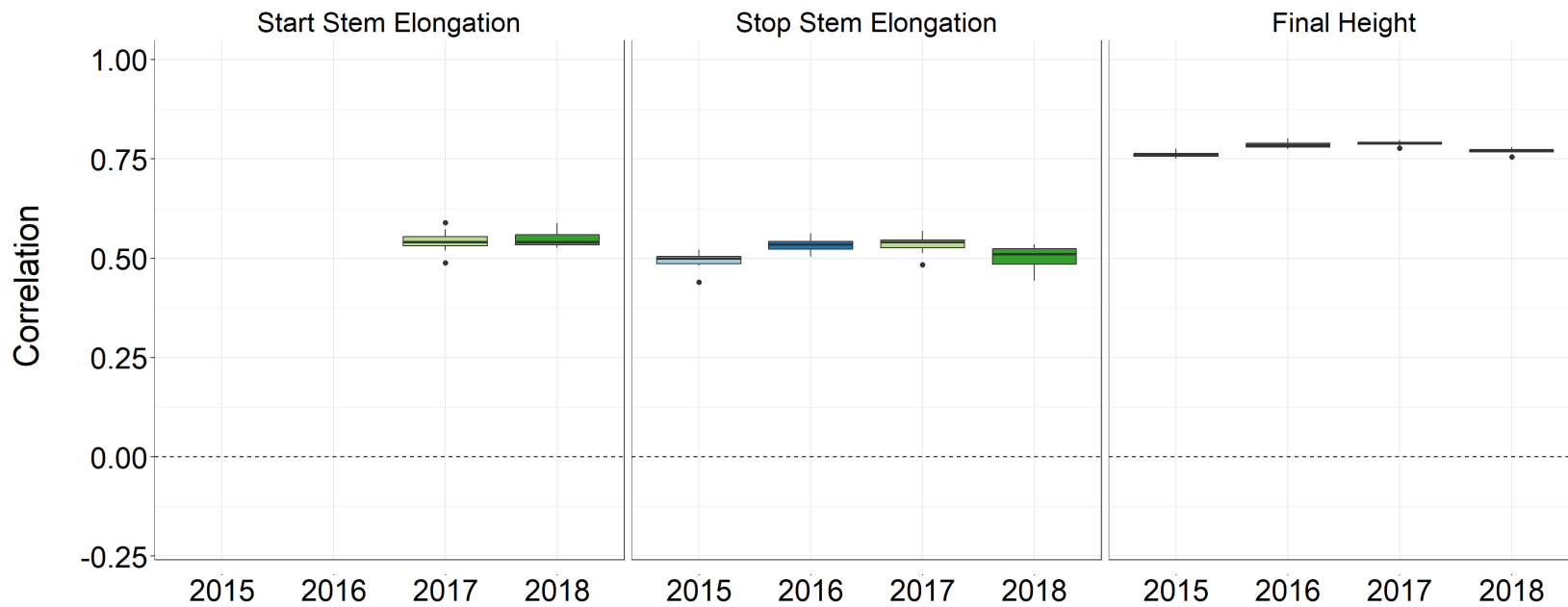
Lukas Roth, Hans-Peter Piepho, and Andreas Hund (2022). Phenomics data processing: extracting dose-response curve parameters from high-resolution temperature courses and repeated field-based wheat height measurements

Relationship

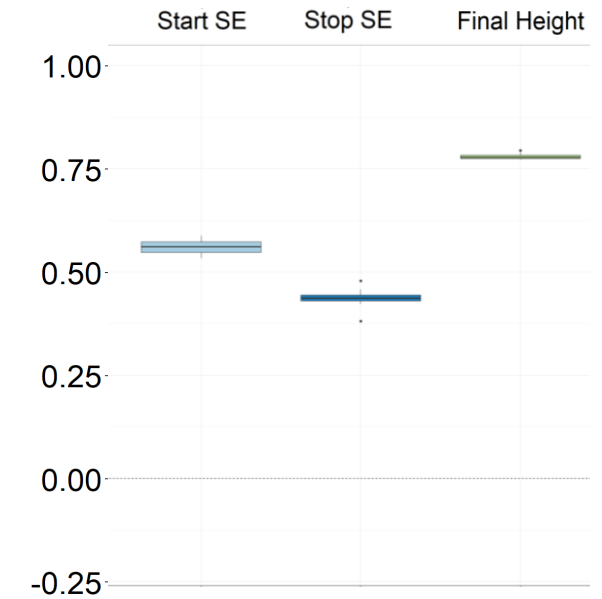


GABI: Year-wise and overall CV of Height-related Traits

Year-wise



Overall

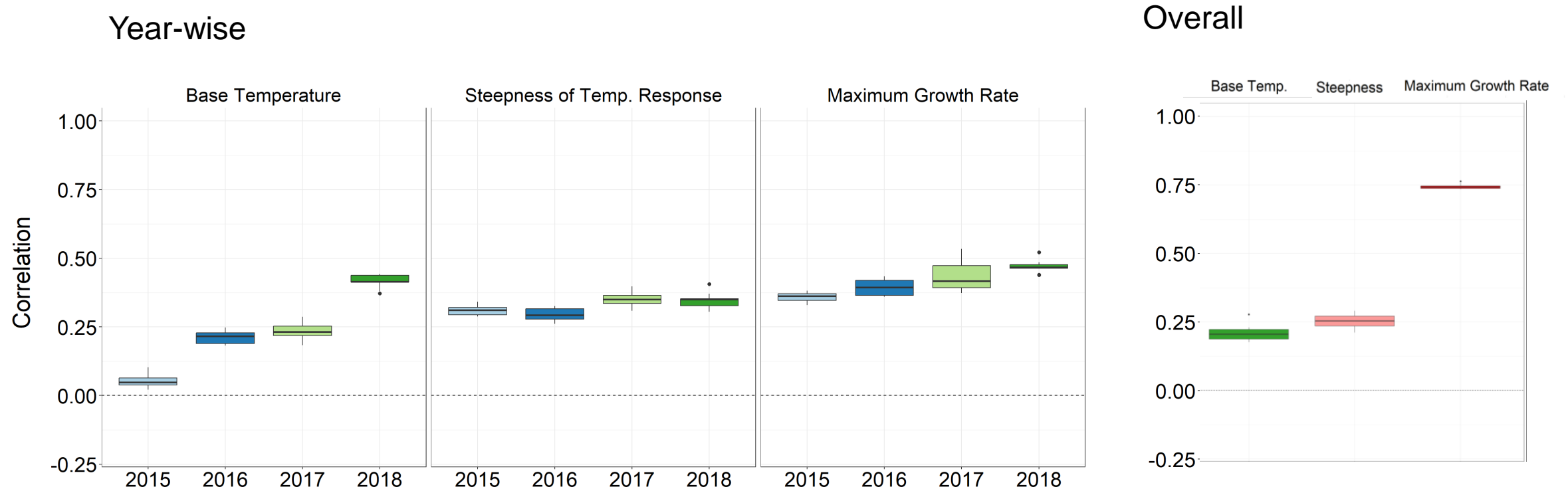


Overall genomic prediction accuracy didn't increase compared with year-wise prediction accuracy

Calculation of GDD

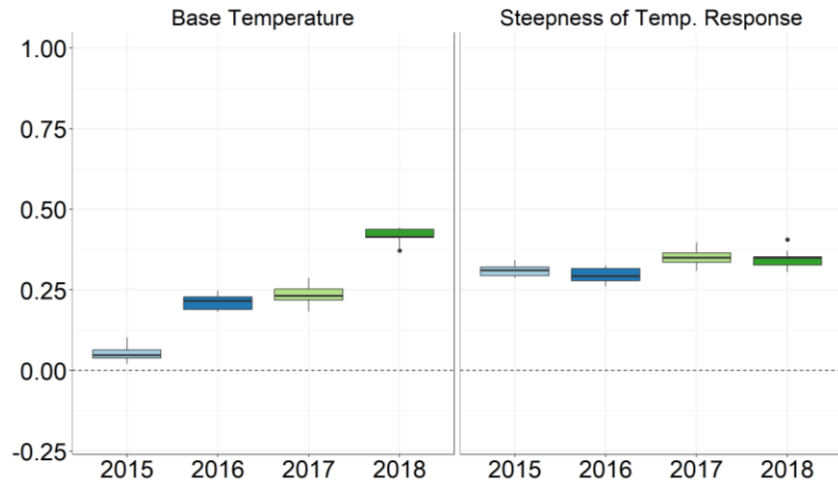
$$\text{GDD(DAS)} = \sum_{d=1}^{\text{DAS}} \sum_{h=1}^{24} \begin{cases} \frac{T_{d,h} - T_{\text{base}}}{24}, & T_{d,h} > T_{\text{base}}, \\ 0, & T_{d,h} \leq T_{\text{base}}, \end{cases}$$

GABI: Year-wise and overall CV of Temperature-related Traits

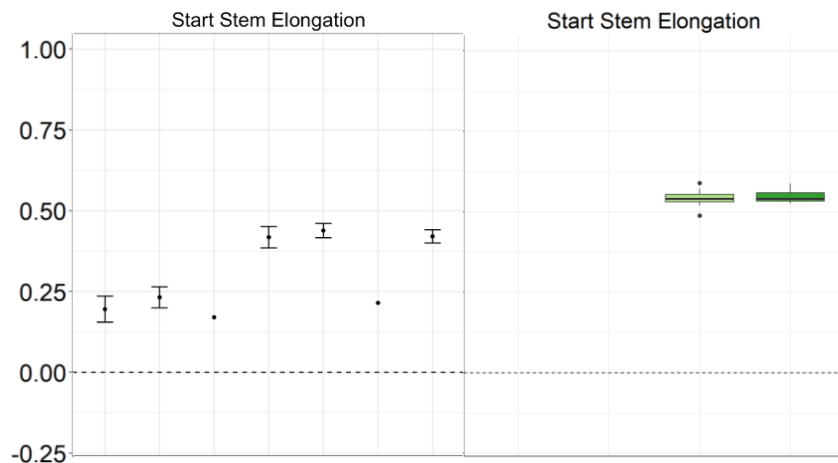


For maximum growth rate, overall genomic prediction accuracy increased compared with year-wise prediction accuracy

Discussion and Conclusion



Still quite low prediction accuracies for temperature related traits



Higher prediction accuracy of start of stem elongation with GABI panel

Crop Rotation

- Since 2012: winter wheat, soybean, buckwheat and maize

Example from RGB imaging at different scales



FIP dolly, 3m distance 0.6 mm GSD

UAS, 28 m altitude, 3 mm GSD

(a)



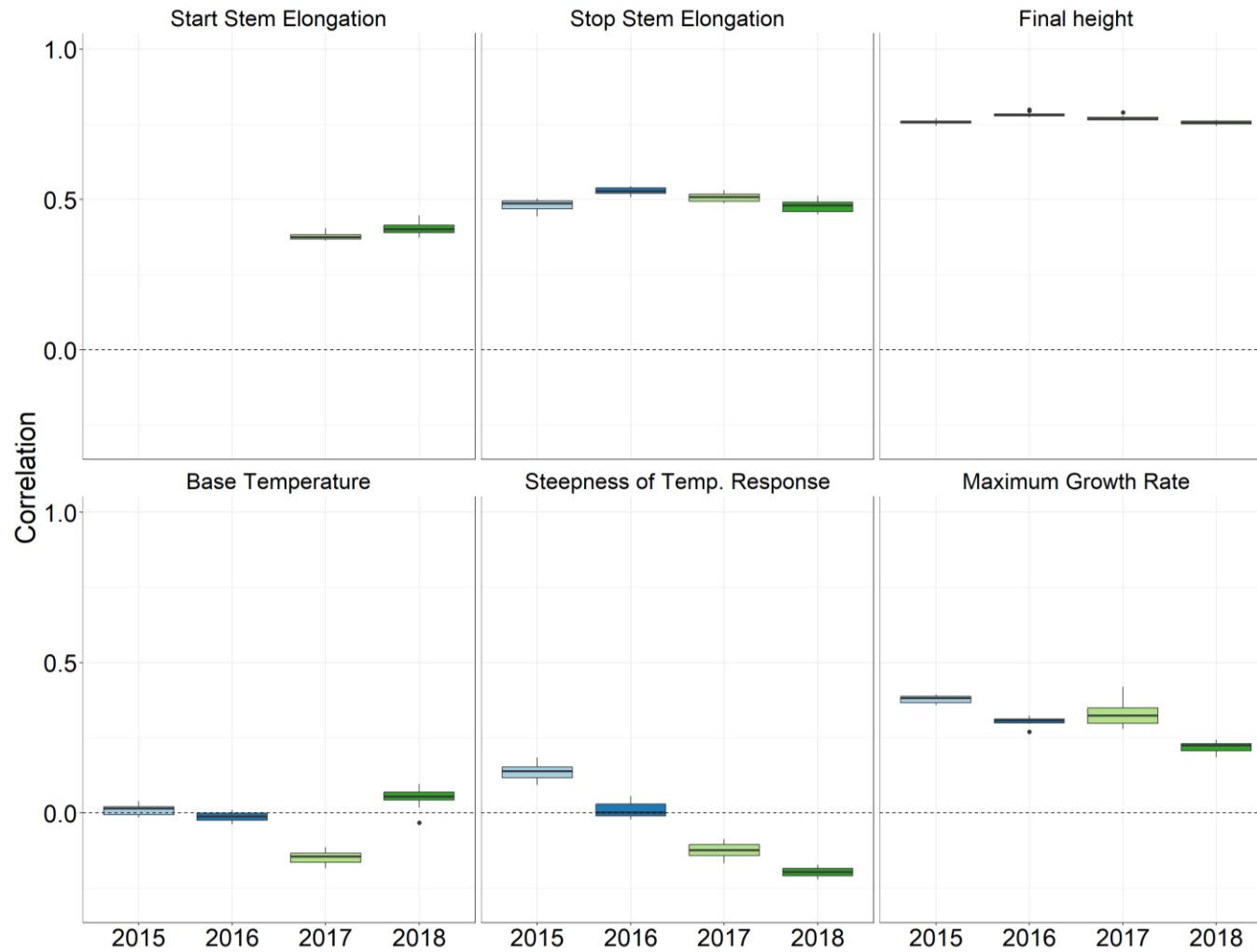
(b)



Ground control point

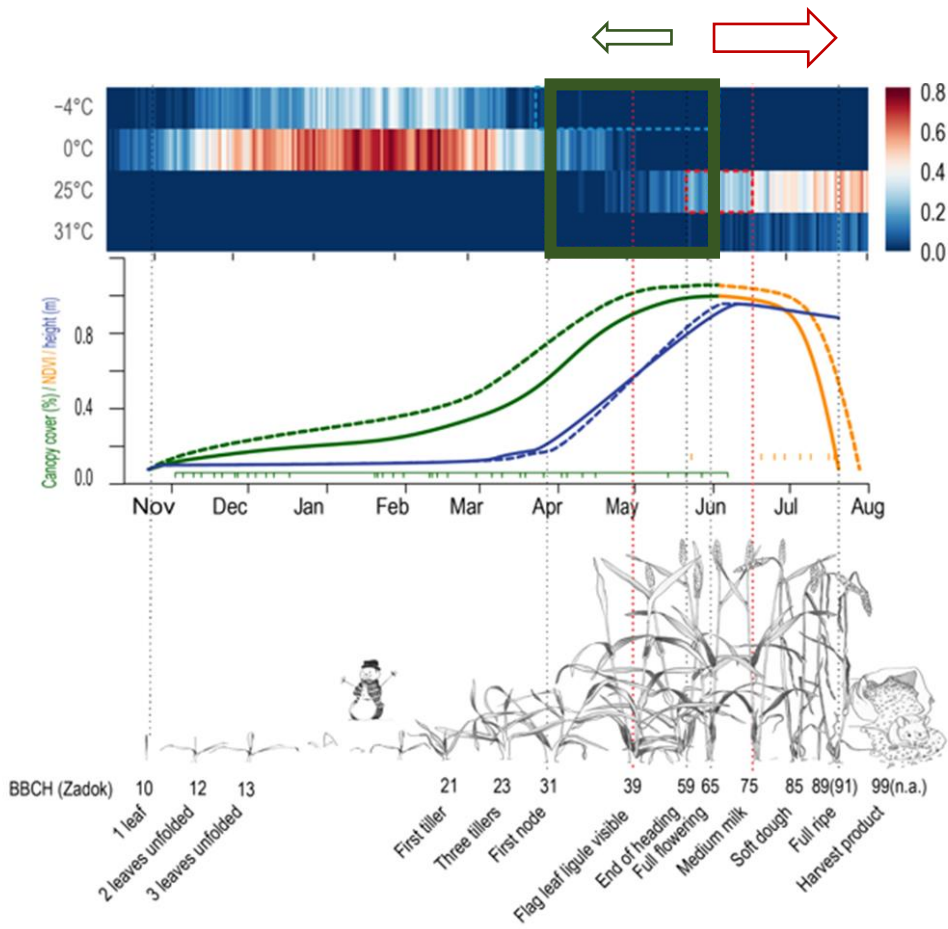
Roth et al. 2020, Plant Phenomics, DOI: 10.34133/2020/3729715

GABI: Prediction of unseen genotypes in unseen years



Prediction of unseen genotypes in an unseen environment

Challenge



Focus: Effect of climate change on wheat yield

Breeders aim at keeping sensitive phase «stress free»

Higher temperatures due to climate change will result in altered phenology and earlier/shorter «stress free» phase

- **Escape:** stay in «stress free» zone: shorter vegetation period and yield reduction
- **Heat tolerance:** later varieties with heat tolerance

Where is the balance?

How can our phenotypic and genomic prediction methods contribute to answer this question?

2.2 GlobalWheat

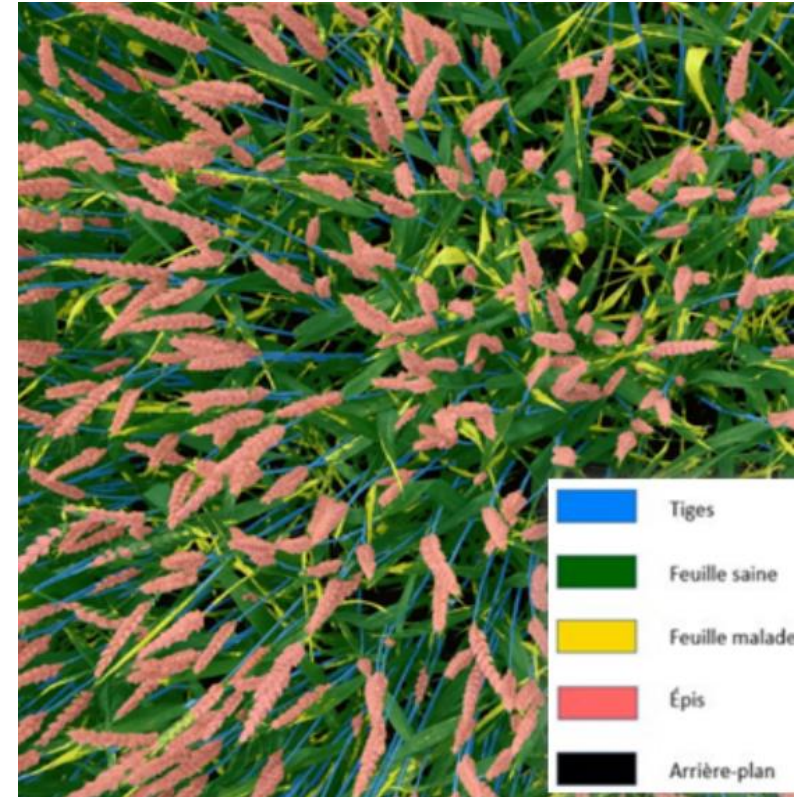


2.2 Global Wheat Dataset

Segmentation:

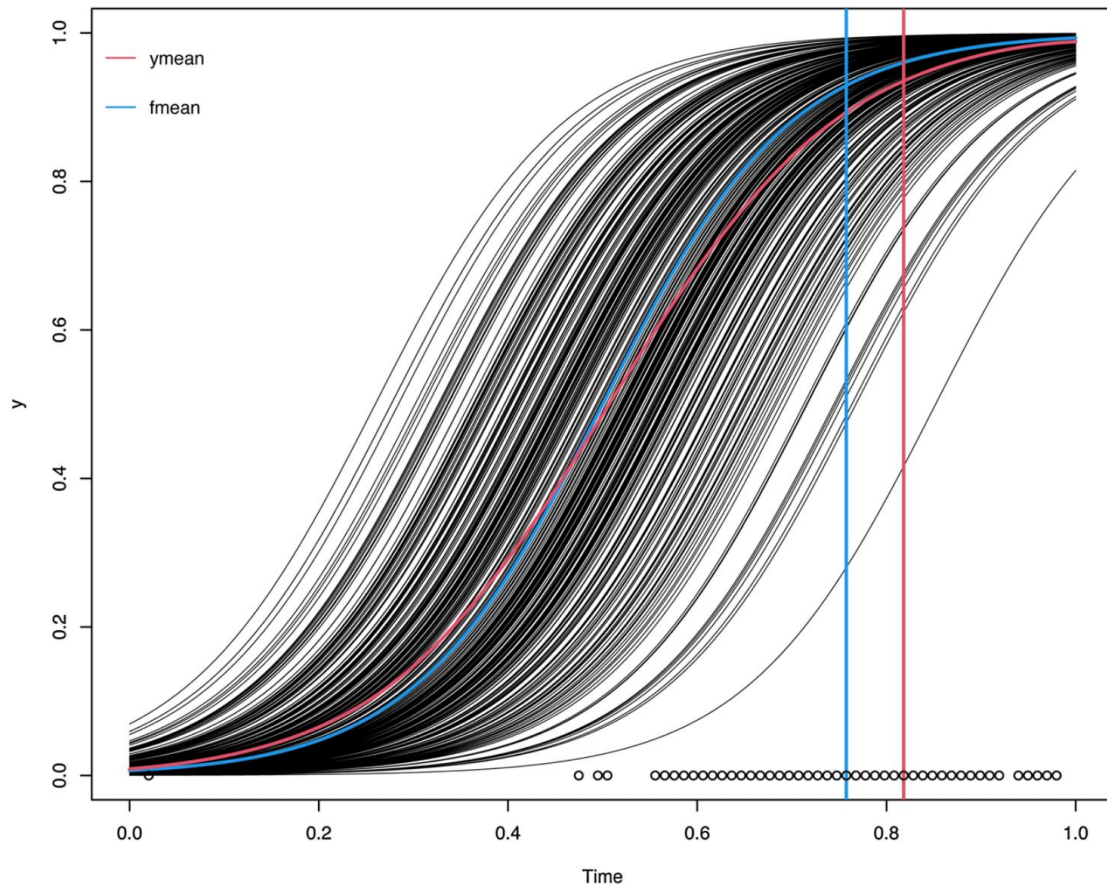
Define a dataset with most relevant features

- Spikes
- Stem
- Leaves
- Soil

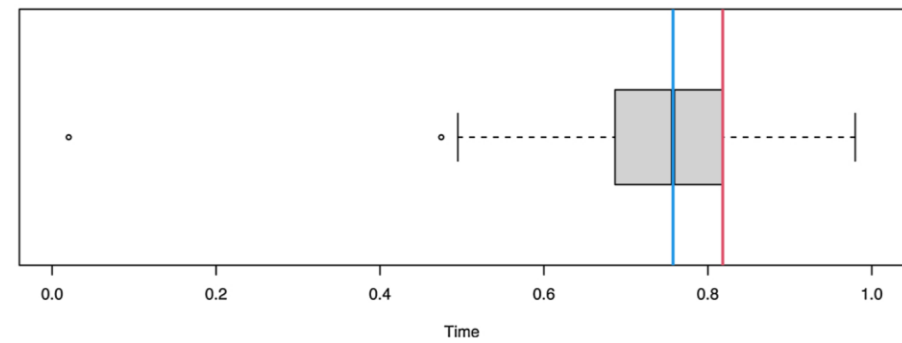


Spatial and temporal correction

a) Growth curves

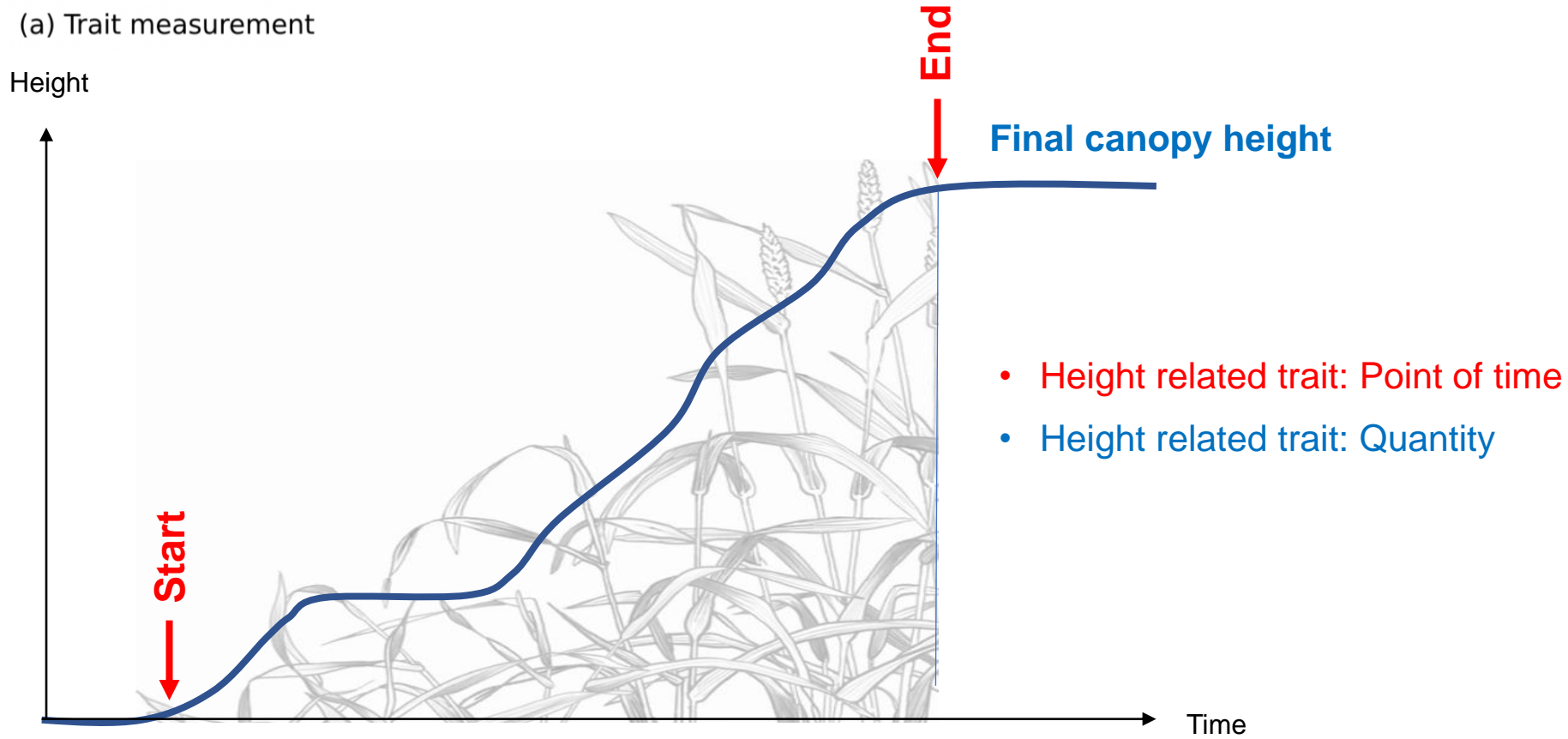


b) Extracted time points



Traits

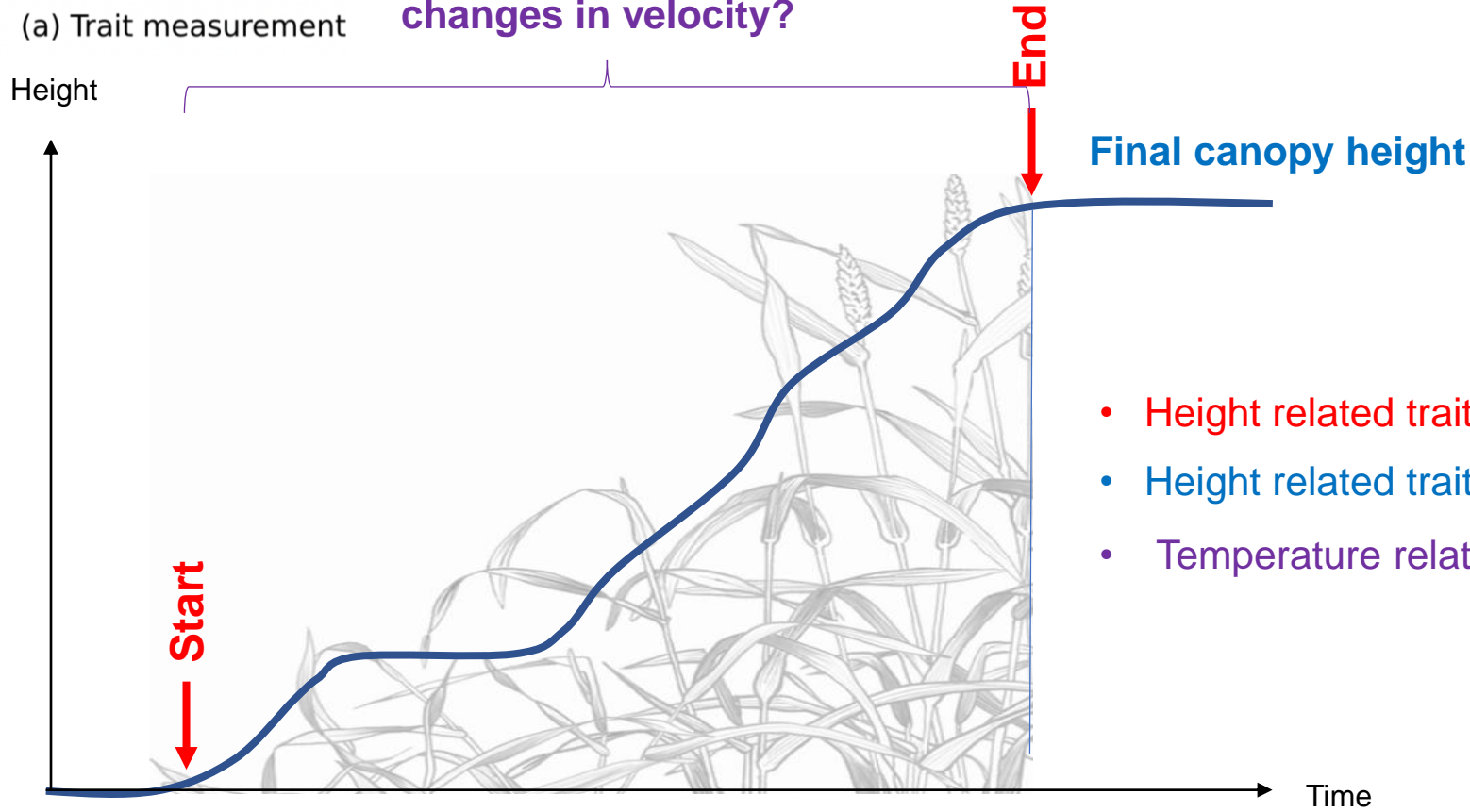
(a) Trait measurement



Lukas Roth et al. (2021) Phenomics data processing: A plot-level model for repeated measurements to extract the timing of key stages and quantities at defined time points, visualized by Andi Hund.

How well does temperature describe changes in velocity?

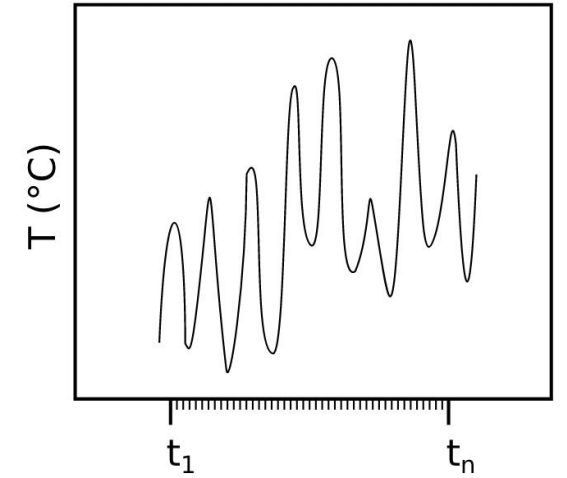
(a) Trait measurement



- Height related trait: Point of time
- Height related trait: Quantity
- Temperature related traits

Lukas Roth et al. (2021) Phenomics data processing: A plot-level model for repeated measurements to extract the timing of key stages and quantities at defined time points, visualized by Andi Hund.

(b) Covariate measurement



(c) Dose-response curve

