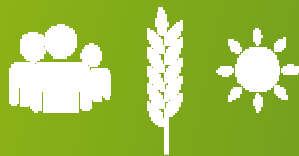


#DigitAg



Model assisted phenomics and phenotype modeling



Pierre Martre, Shouyang Liu



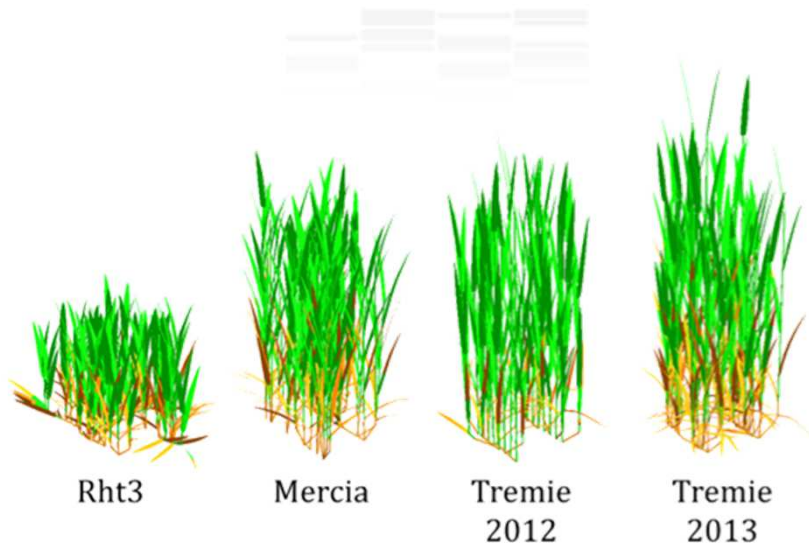
#DigitAg – Modelia Seminar on
Data assimilation

Montpellier, 21 June 2018

Outline

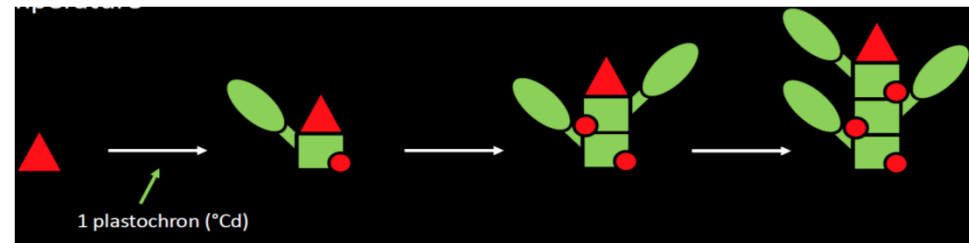
1. Dynamics models for phenotype modeling
 - Process-based models
 - Functional structural plant models
2. Role of crop modeling in genetics and breeding
3. Interpretation of high-throughput phenotyping data
4. Model assisted phenotyping
5. high-throughput phenotyping data assimilation(parameter estimation)
6. Conclusions

Functional Structural Plant Model (FSPM)



Robert et al., 2017

Garin et al., 2017



Fournier et al., 2003

- ❑ Individual based models (behavior of individual plants determine canopies properties)
- ❑ Plant architecture is represented in 3D
- ❑ Explicit feedback between plant growth and environmental drivers
- ❑ Only few functional processes considered

Process-Based Model (PBM)

Inputs

Crop Management

Sowing date
N fertilization
Irrigation
....

Soil

Hydraulic characteristics
% organic matter
pH
....

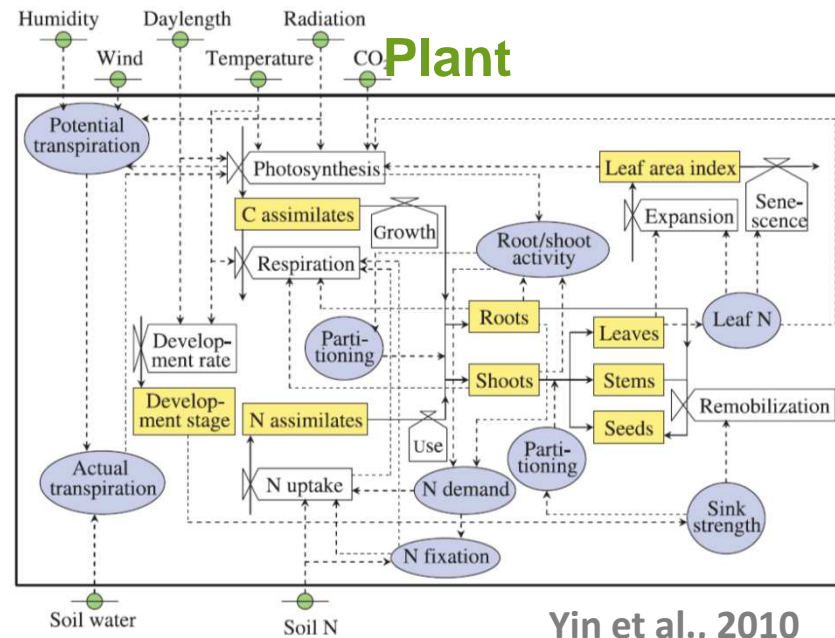
Weather

Temperature
Radiation
Precipitation
....

Cultivar

Radiation use efficiency
Phyllochron
Potential grain size
Fruiting efficiency
.....

Crop model



Simulations

Grain yield

% Proteins

GPD

NUE

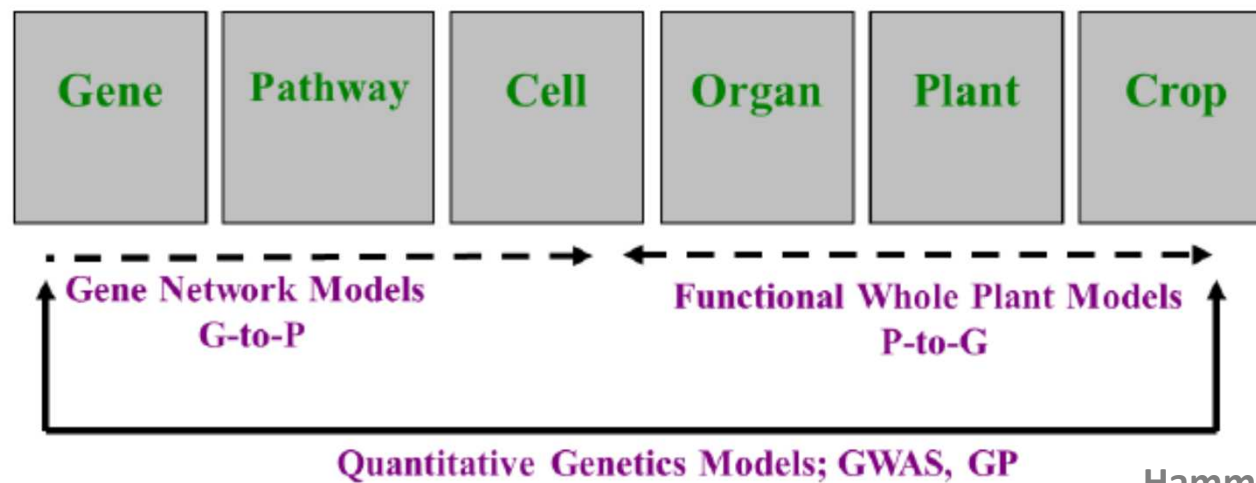
WUE

....

- ❑ Focus on functional processes and feedbacks (explicit or implicit {emerging properties])
- ❑ Canopy architecture not explicitly considered (canopy = 1D turbid medium)

Phenotyping distance and prediction

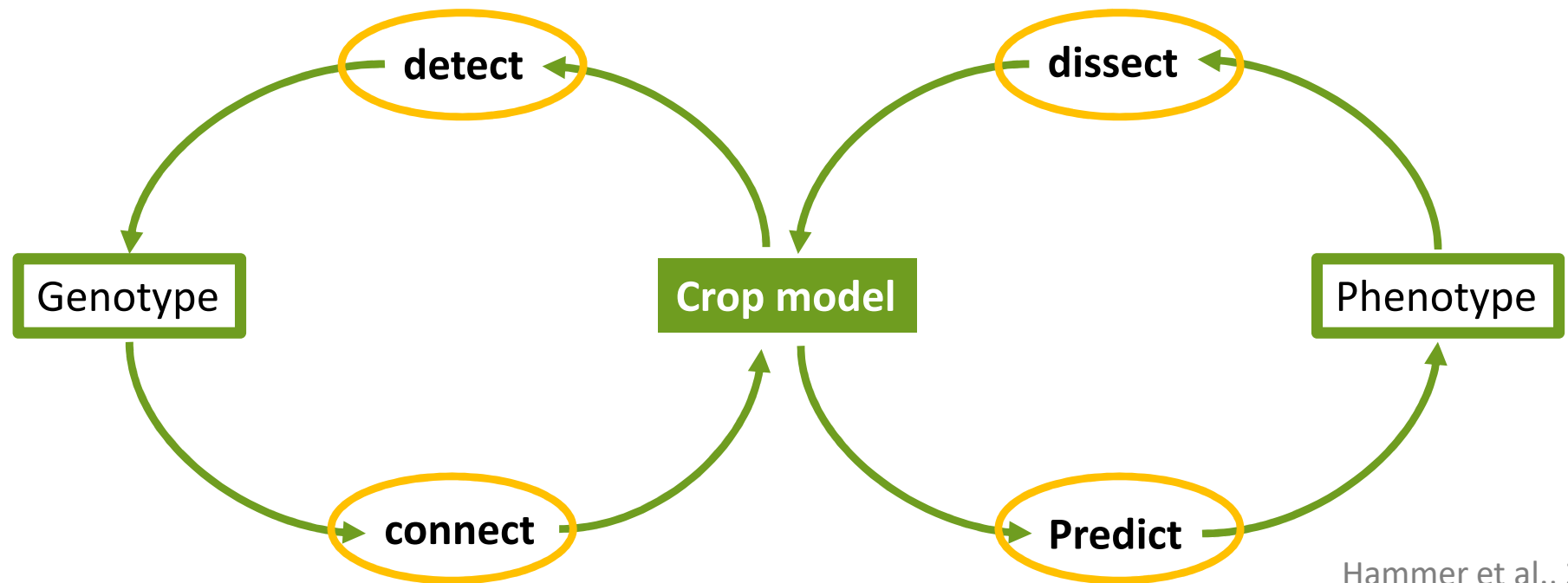
Crossing scales of biological organization confounds prediction



Hammer et al., 2016

- Crop model by integrating traits effect reduce the phenotypic distance
- Genetic analysis should be carried on simple traits and crop model allow scaling to the crop

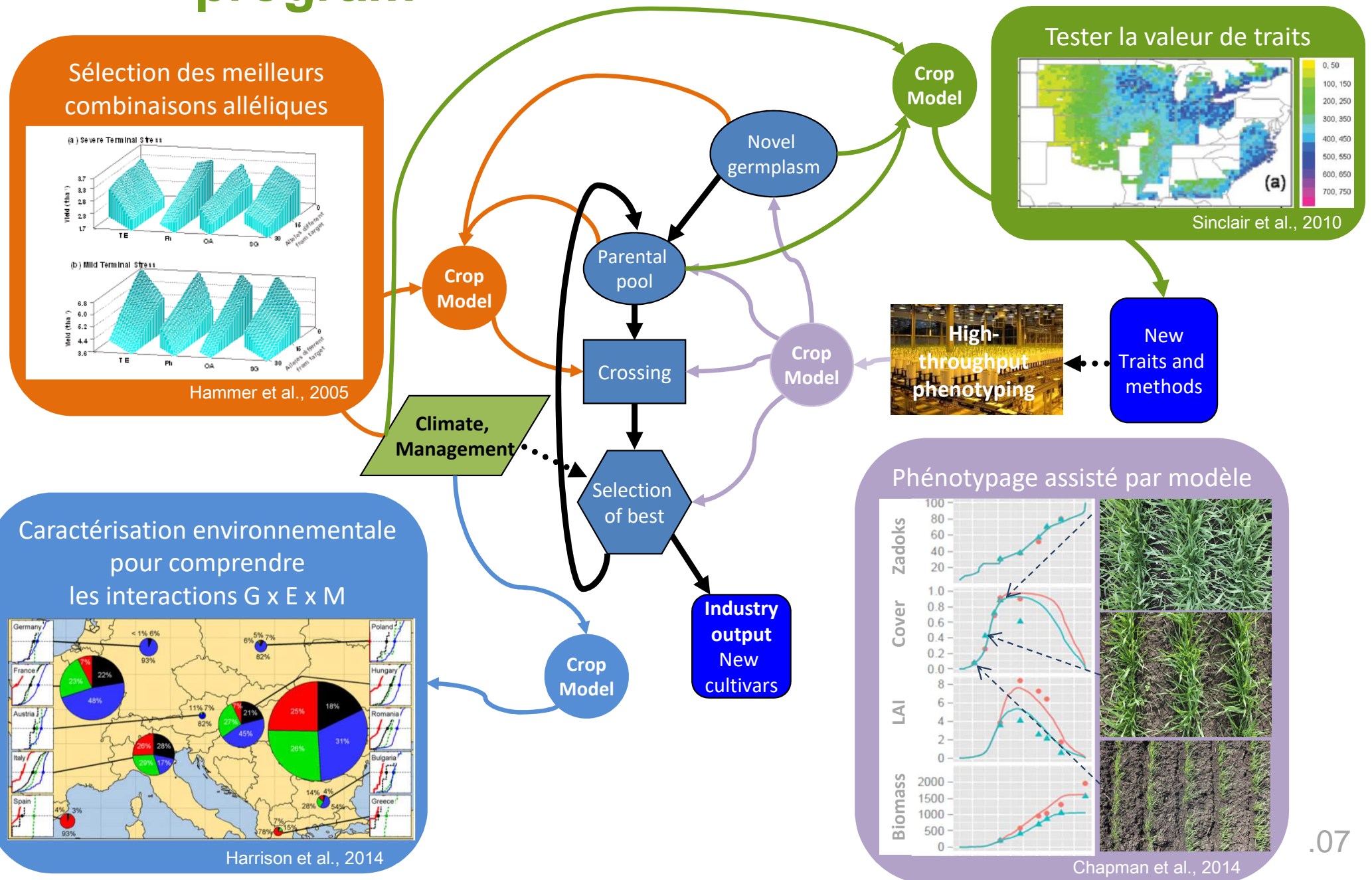
Role of crop modeling in genetics and breeding



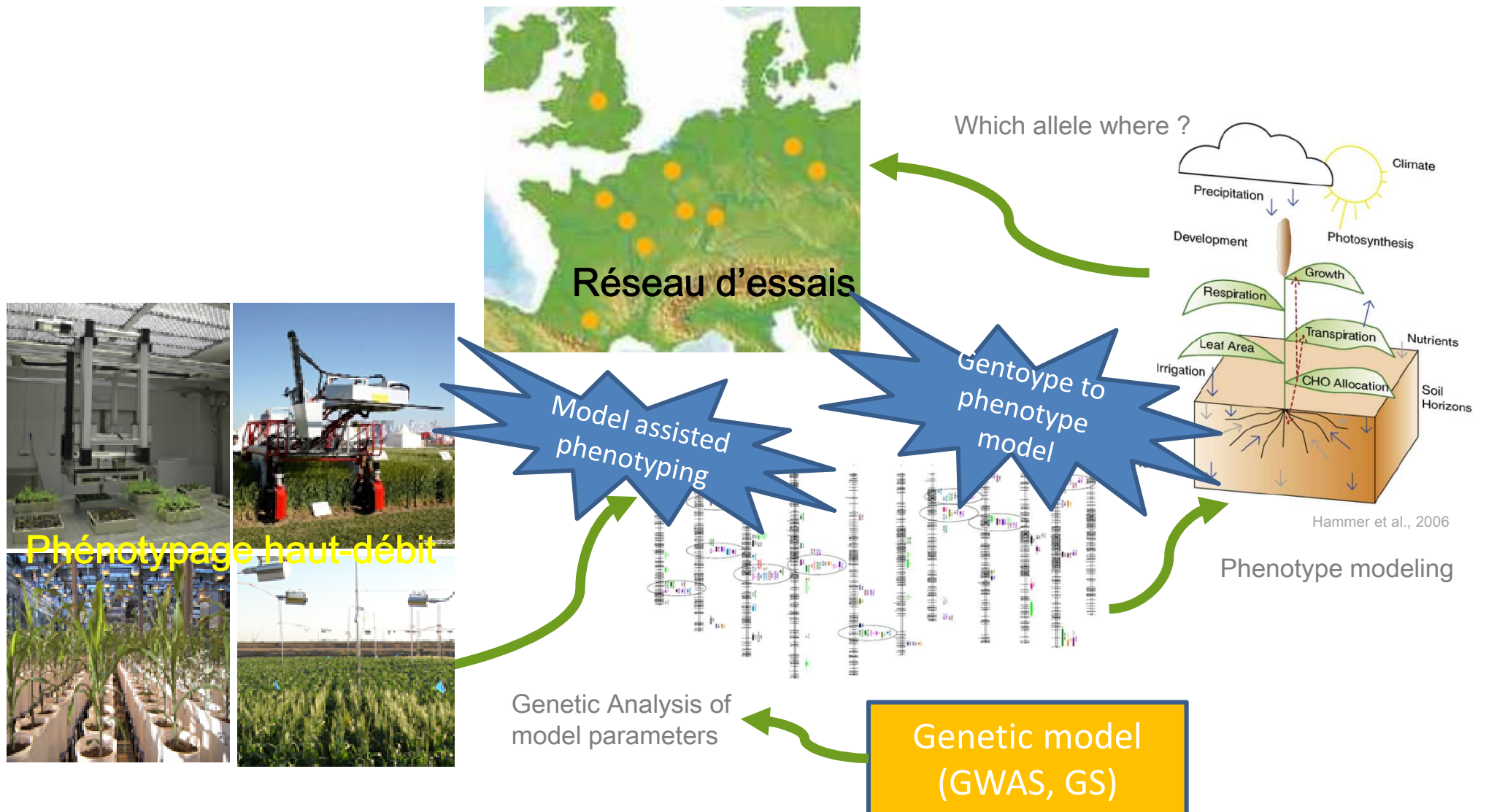
Hammer et al., 2016

- Predict** – Trait evaluation in target environments to unravel G x E interactions
- Dissect** – Understand and simplify complex traits (use of NILs, mutants)
- Detect** – Inform phenotyping for QTL detection
- Connect** – Link QTL/genes to crop attributes/processes

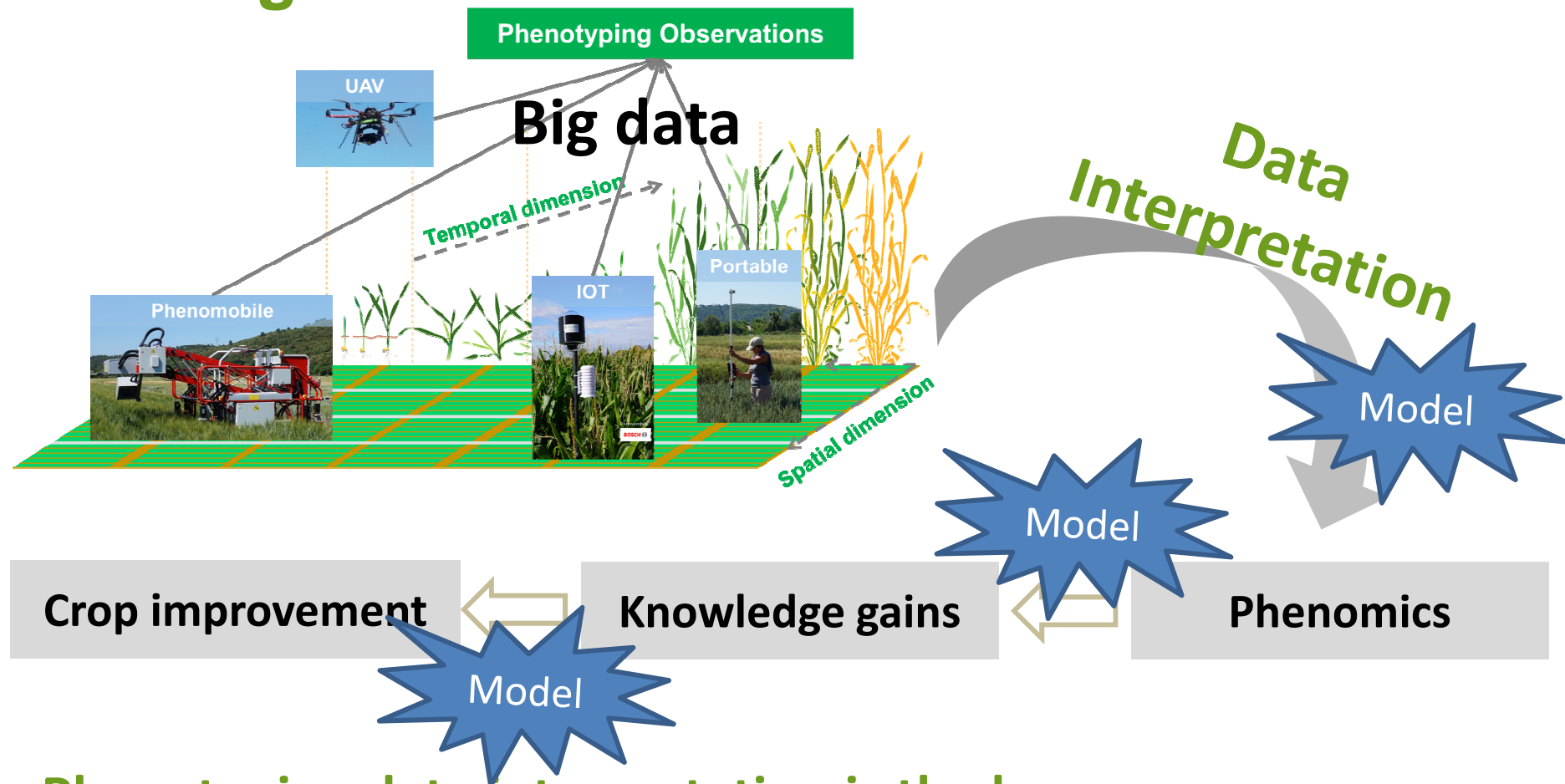
Integration of modeling in a breeding program



QTL/gene-based modeling



High-throughput Phenotyping starts from big data

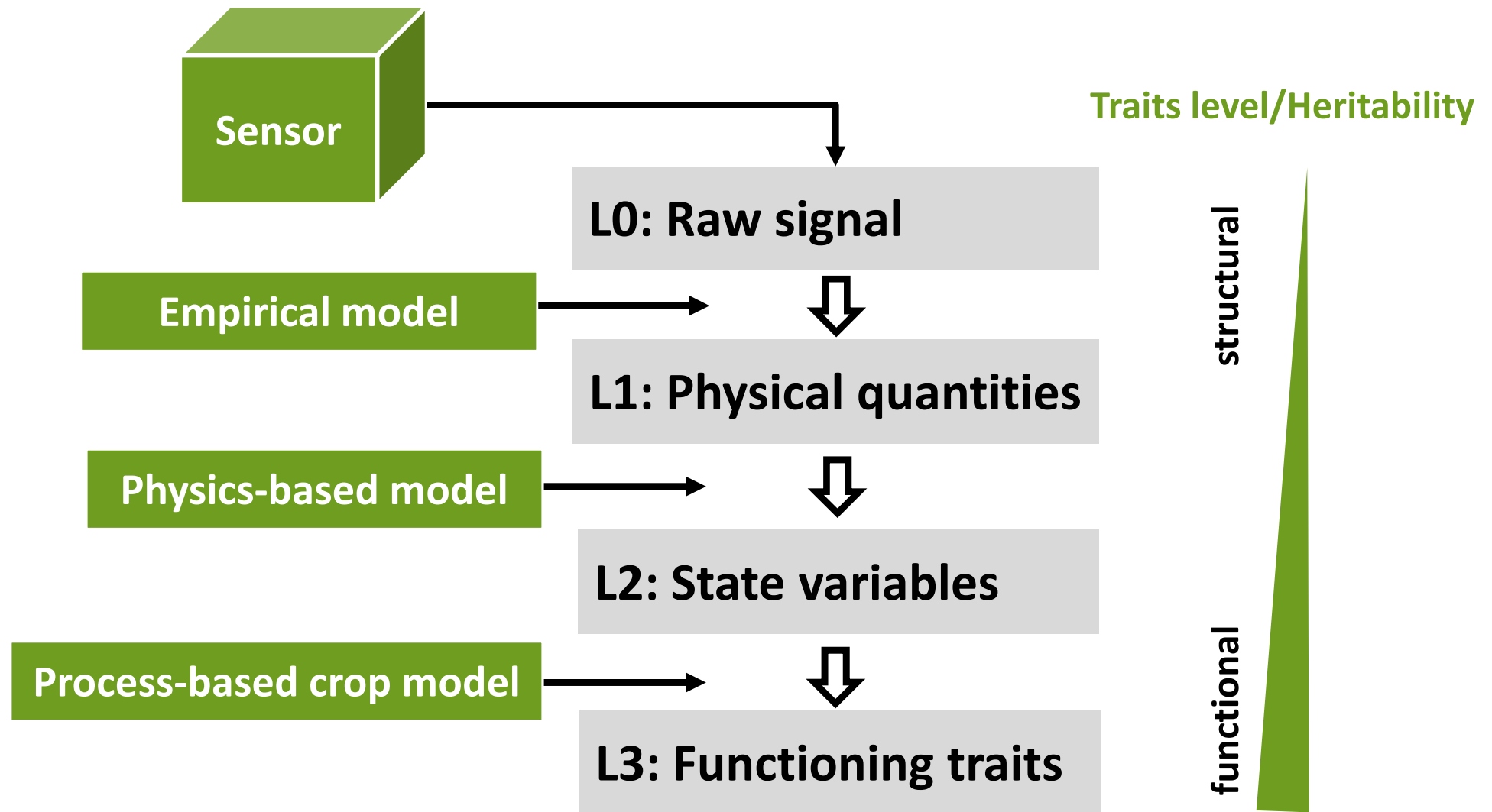


Phenotyping data interpretation is the key

- to transform big data to phenomics
- and finally contribute to the crop improvement

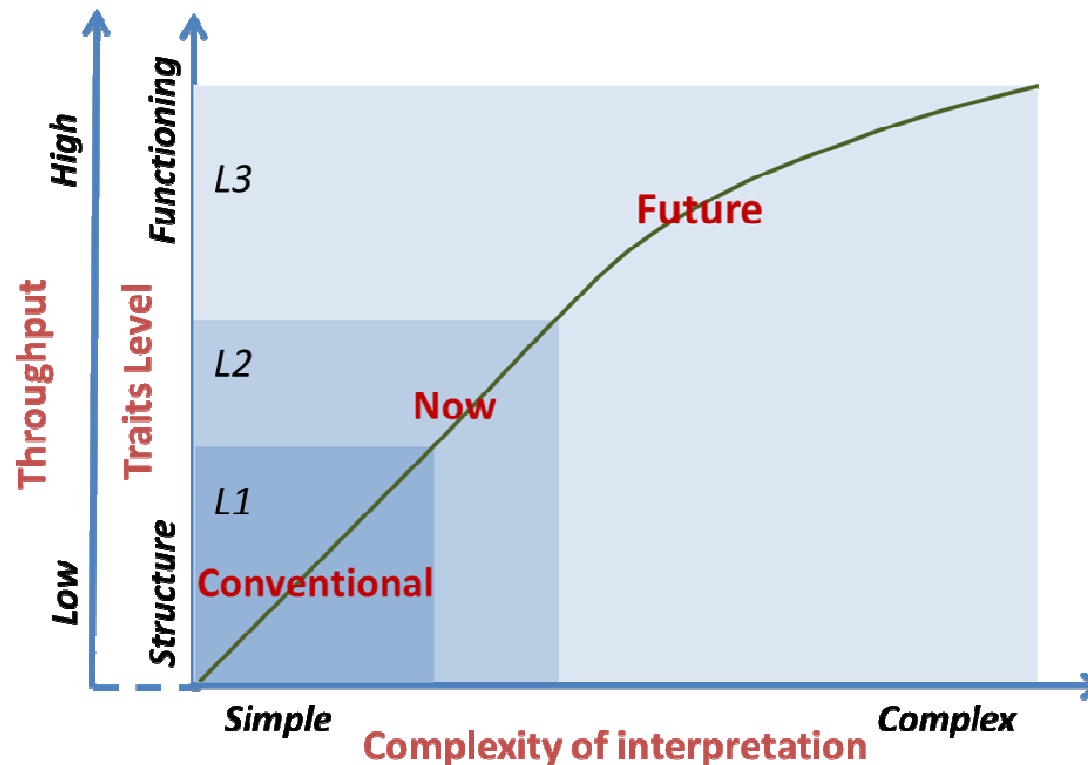
Slide from S. Liu

Levels of Phenotyping data interpretation



Slide from S. Liu

Phenotyping throughput is limited by interpretation



Phenotyping data interpretation:

Goal: maximize the traits' outcome with satisfactory accuracy

Strategy: Optimize the information use from measurements and prior knowledge

Slide from S. Liu

Levels in phenotyping data interpretation

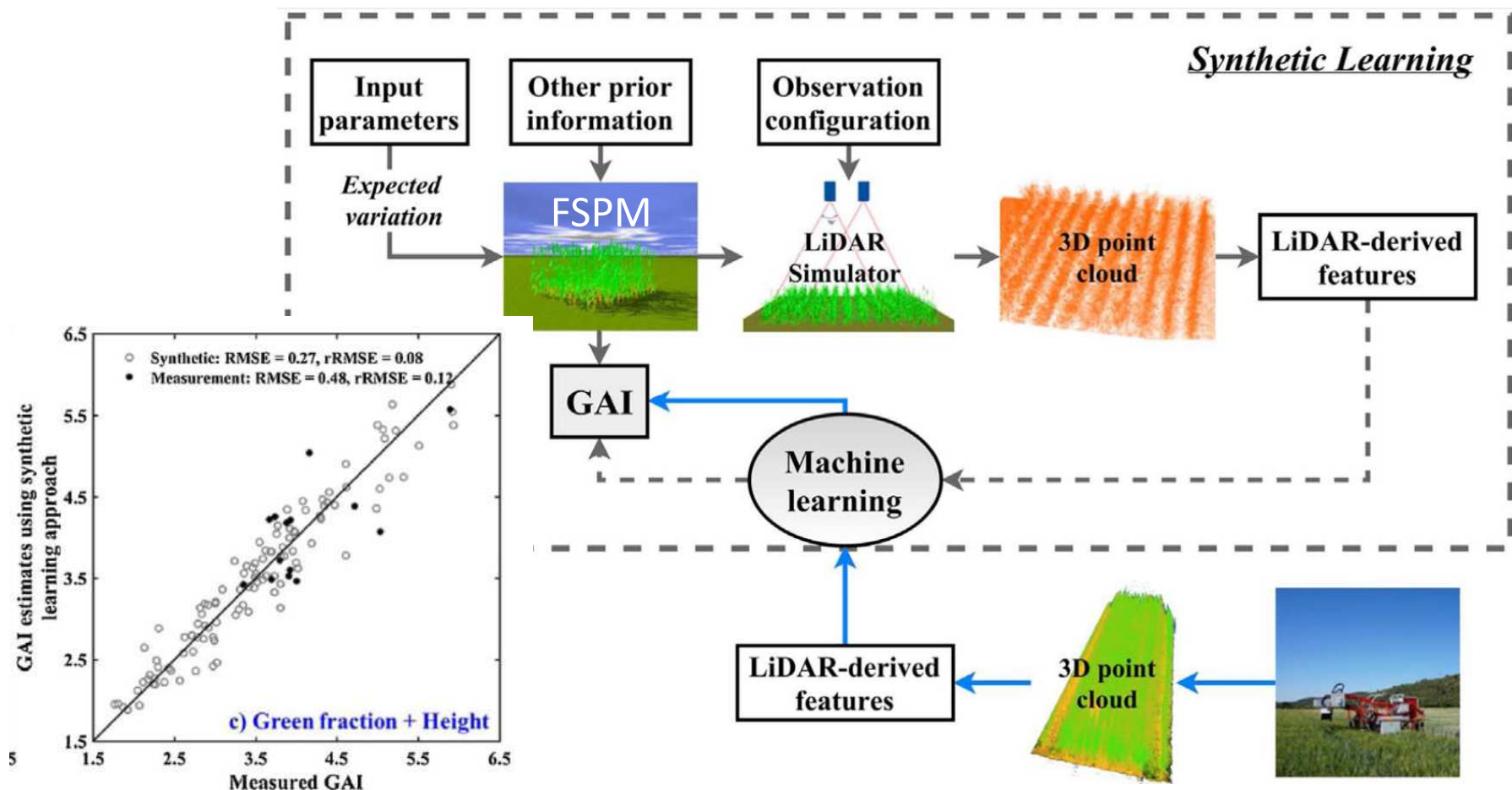
Level	Sensor	Traits	Method
L1	RGB	Plant distribution	Statistic model
		Plant density	Machine learning
		Ear density	Deep learning (Simon Madec)
L2	LiDAR	GAI	Digital Plant Phenotyping Platform (D3P)
	Multispectral	Chlorophyll content	
L3	RGB	3D canopy structure	
		Light interception	

Slide from S. Liu

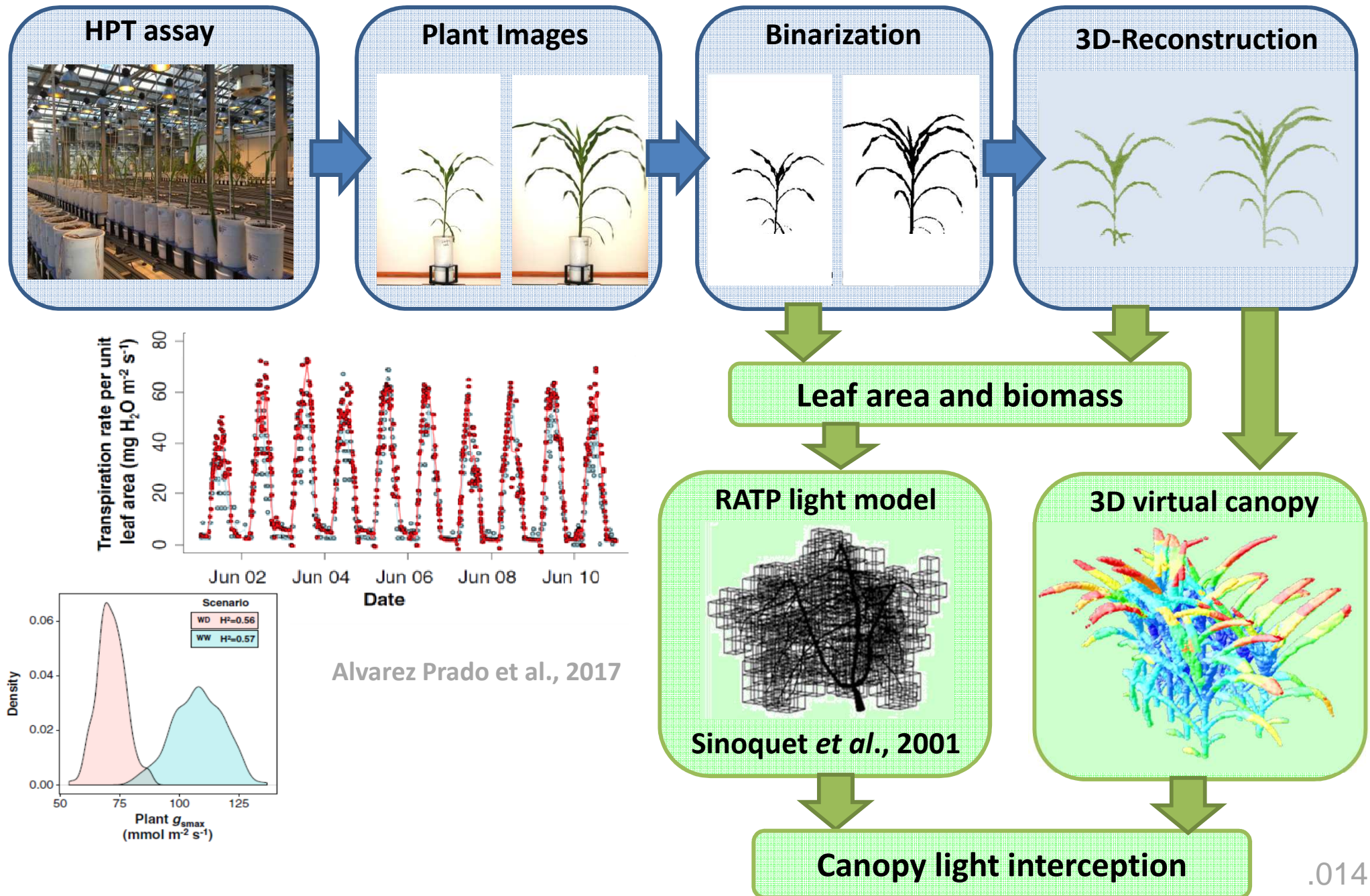
Model assisted field phenotyping

Estimating wheat green area index from ground-based LiDAR measurement using a 3D canopy structure model

Shouyang Liu^{a,*}, Fred Baret^a, Mariem Abichou^b, Fred Boudon^c, Samuel Thomas^d, Kaiguang Zhao^e, Christian Fournier^f, Bruno Andrieu^b, Kamran Irfan^a, Matthieu Hemmerlé^g, Benoit de Solan^d



Model assisted platform phenotyping



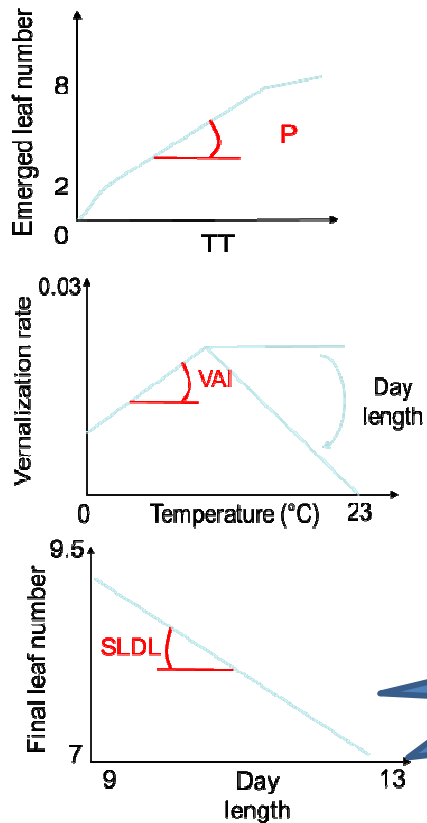
High-throughput data assimilation

Prediction of wheat flowering time

Phenotyping

Phenotype prediction
(in new environments)

Trait distinction



Genetic material:
100 RILs of Offanto x Cappelli

Vernalization at
4°C for 8 weeks

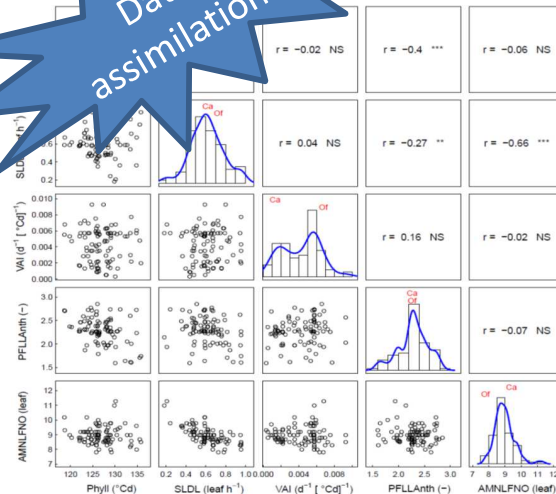
Plants grow until
anthesis in a
greenhouse or outside

Seeds germinated
in petri dishes

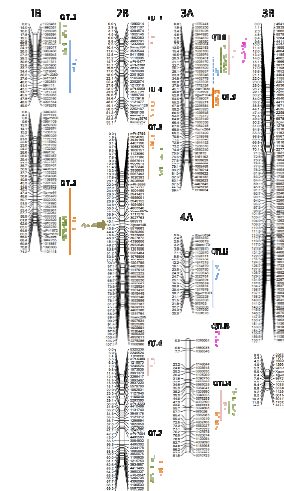
Plants transplanted
in pots

Measurements:
• Haun stage
• Final leaf number
• Anthesis date

**Data
assimilation**



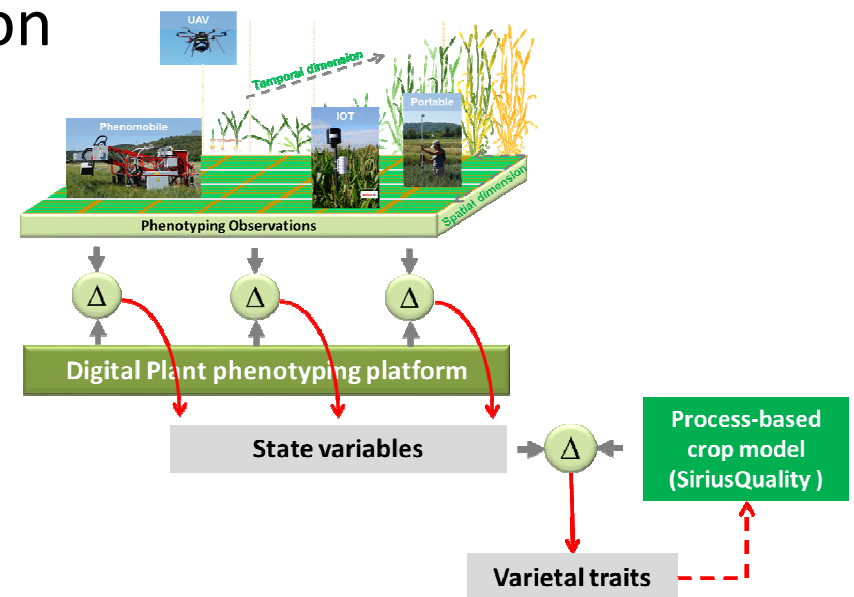
Genetic analysis



$$y_j = \mu + \sum_{i=1}^n a_i QTL_{ij}$$

Conclusions et perspectives

- ❑ Phenotyping-oriented crop model are required (biological/physical coherence).
- ❑ Phenomics is renewing model development, improvement, and testing.
- ❑ To retrieve functional traits Integration of D3P with process-based model is required.
- ❑ To develop real time data assimilation for precision agriculture emulators (meta-models) will probably be required (computation time).



Liu, 2016

Thank you for your time

pierre.Martre@inra.fr



M3P 
Montpellier Plant
Phenotyping Platforms

 **lepse**
Montpellier
Laboratoire d'Ecophysiologie des Plantes
Sous Stress Environnementaux

www6.montpellier.inra.fr/lepse
www1.clermont.inra.fr/siriusquality