

High-throughput phenotyping of fruit tree development, light interception and wateruse related traits: a case study on apple tree

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Tree architecture is genetically highly variable in fruit species



Monet (1983); Scorza (1984); Scorza 1984; Bassi et al., 1992) photos from Hollender and Dardick (2014)

Apple tree



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Lespinasse, (1992); Lauri and Lespinasse, (1993); Forshey and Elfving, (1989)

Both tree topology and geometry are varying depending on the variety

Tree architecture is a key feature for fruit production in the face of climate change



In silico modeling achieved optimal combinations of architectural and functional traits

> Resource use efficiencies are important targets, not accounted for in breeding programs

Their genetic control has rarely been investigated jointly with architecture
Major bottleneck : phenotyping these traits in field conditions on adult trees

Phenotyping: overall strategy



<u>T-LiDAR</u> : Architecture (geometry-topology) Nb of axes, tree volume, etc.





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Airborne imaging:

- Multispectral imagery (Vegetation indices)
- Thermal images (Stress indices) Response of the canopy water loss to drought





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Production and other variables:

- Yield (fruit number & harvest weight)
- Trunk cross sectional area

Experimental design

INRA Mauguio : Diascope Experimental Unit (43°36′N, 3°58′E, 10m a.s.l.)

- Apple tree French core collection (Lassois et al., 2015)
- 241 genotypes x 4 replicates (same M9 rootstock)



- 10 rows (ca. 200m)
- Plantation spring 2014
- Micro-irrigation, individual emitters
- Row weeding ; grassed inter-rows
- No pruning, fruitlet thinning,
- Conventional spraying
- 5x2m spacing, compatible with architecture characterization

Micrometeorology: Global & PAR radiation, Air temperature & RH, Wind speed & direction

Soil water status: Sentek capacitive probes (0-90cm) +Tensiometers (-30 and -60cm) For 14 tree locations



Mediterranean climate (dry summer) Soil : limited water holding capacity: irrigation withdrawal → water stress

- 2 different water regimes during summer
- WW = well irrigated
- WD = water deficit
- WW / WD trees, facing each other (2 pairs)

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Using T-LiDAR to assess tree architectural variability

1. Field acquisition

Two campaigns in summer and winter 2017

250 scans Scan resolution 0.06 (6mm at 10m) Acquisition duration: 30'/scan 5 days for 998 trees





2. Data post-processing (filtering, topological reconstruction, tree segmentation)



Tree segmentation

ComputeTree and RiscanPro



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PlantScan3D, Boudon et al.,⁶2014

Architectural traits estimated from T-Lidar scans (winter)

Topological variables:

Number of axes according to their types – T-Lidar vs digitizing data



Axis number - T-LiDAR data



© B. Pallas, INRAe Complete dataset

Variables	Mean value (Digitizing)	R ²
Total nb of axes	36.3	0.66
Nb of long + medium axes	22.7	0.98
Nb of short axes	13.6	0.14
Cumulative axis length	7.2m	0.99
Cumulative length of long & medium axes	6.6m	0.99
Cumulative length of short axes	0.6m	0.10

Pallas et al. 2018

Architectural traits estimated from T-Lidar scans (summer)

Geometrical variables

- Tree radius R
- Tree height H
- Convex volume (c_volume)
- Alpha volume (a_volume)
- Convexity (a_volume/c_volume)





Alpha hulls

Convex hull

Coupel-Ledru et al. 2019

Architectural traits estimated from T-Lidar scans *Computation of leaf area profile parameters (3 trees)* (summer)

Geometrical variables

- Leaf area profile
- STAR (Silhouette to total leaf Area Ratio)



High-resolution multispectral and thermal-IR imagery



Thermal-IR imaging



Some results on traits retrieved from T-LIDAR scanner and multispectral imaging

Coupel-Ledru et al., 2022

A strong genotypic variability on all traits related to architecture and light interception, and vegetation indices



Leaf area

Shoot development



Light interception efficiency (self-shading)





No effect of WD on these traits (or very light)

Likely due to the short duration (1 month) and period (July) of WD application

A strong genotypic variability on all traits related to **architecture** and **light interception**, and **vegetation indices**

Heritability of tree traits

Mixed effect model $Y = \mu + G + T + X + \varepsilon$

μ : intercept, G: genotypic effect (randon), T: treatment effect (fixed), X possible other fixed effect (row, date...)

Measurement type	Trait	Broad sense H ²
Multi-spectral imagery	NDVI	0.94
	GNDVI	0.94
	MCARI2	0.92
T-LiDAR data	c_volume a_volume	0.82 0.77
	C _i	0.86
	STAR	0.79
	height	0.81
	total_length	0.80
	nb_axes	0.83



Two different groups of traits within the collection



Genotype-dependent control of canopy temperature in WW conditions





Large genotypic variations in the response of canopy surface temperature to intensifying soil drying



Different genomic regions control canopy temperature, vegetative architecture or light interception efficiency





Multiple allelic combinations exist within the collection for the three groups of traits Possibilities of selecting independently for architecture, light interception efficiency and water loss

Take home messages

- The simultaneous assessment of canopy vegetative development and canopy functioning revealed to be efficient to
 - 1. Classify trees according to their global shape
 - 2. Assess the relevance of indices derived from scans and airborne imagery as proxies for tree architecture and functionning
 - 3. Capture the genetic variability of those traits in an apple tree core collection
 - **GWAS revealed that multiple allelic combinations** exist within the collection for the three groups of traits

It will be possible to select independently for architecture, light interception efficiency and water loss

Find adequate combinations depending on the climatic conditions?

 How to define / evaluate the water-use efficiency on a medium to long term for perennial crops ?

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Diascope : INRAe experimental unit

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Amélioration génétique et adaptation des plantes méditerranéennes et tropicales



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Thank you for your attention

Some related literature:

Coupel-Ledru et al. 2022. Tree architecture, light interception and water use related traits are controlled by different genomic regions in an apple tree core collection. New Phytologist 234: 209-226 <u>https://doi.org/10.1111/nph.17960</u>

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