

# Data Management

## S4: Advanced data management (1)

### Variables & Measurements

Thursday 07/12/2023 - 14:00-16:00 (CET)

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General objectives: Overview of data management for plant phenotyping - focus on FAIR data

Session 4

Advanced data management (1):  
Variables & Measurements

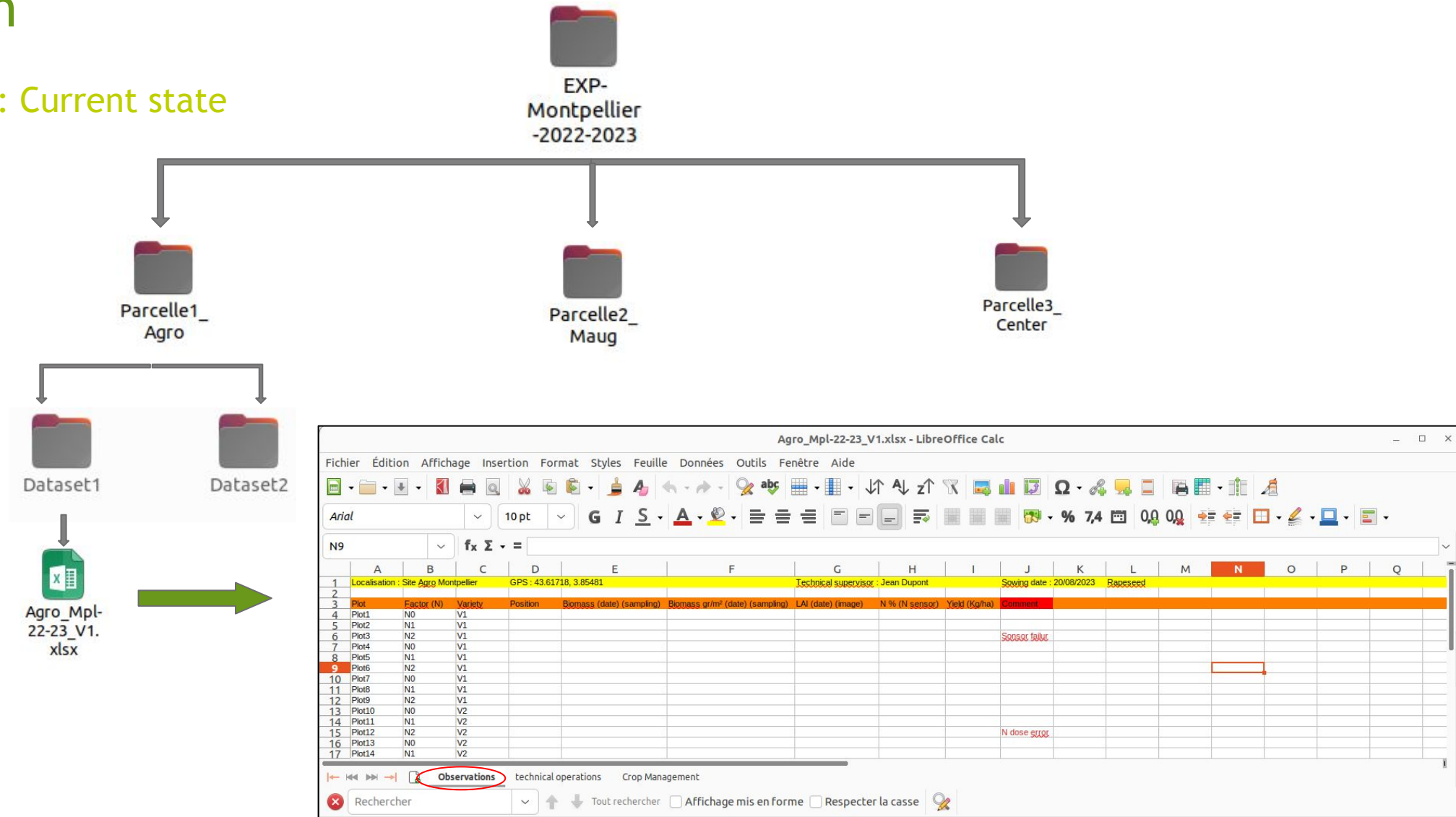
# Overview

## Advanced data management Variables & Measurements (1)

- Reminder of concepts
- Measured data, Variables & Provenance
- Data Pipeline, Automatized data insertion
- Use Case: Sending data to PHIS

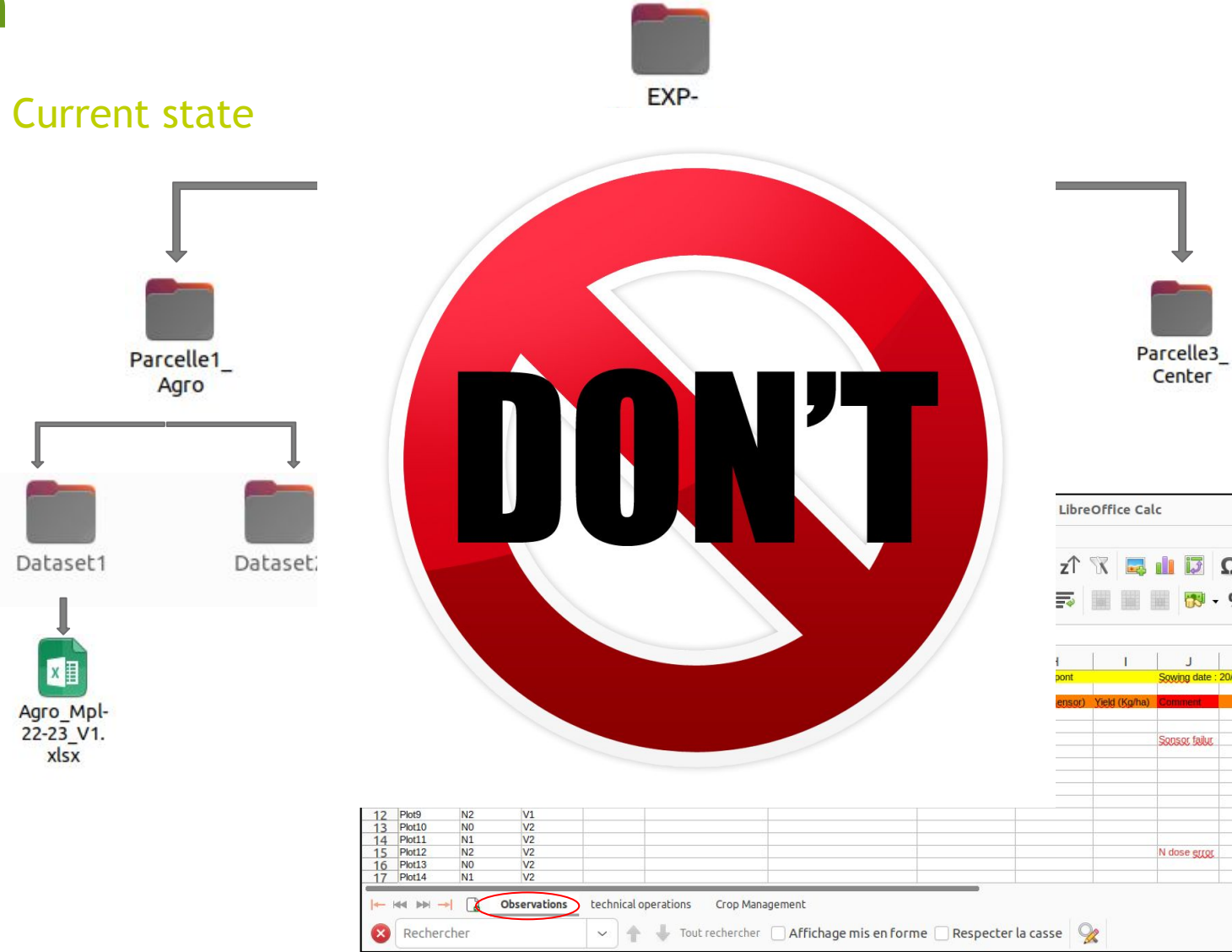
# Introduction

## Experimental Data: Current state



# Introduction

## Experimental Data: Current state



12	Plot9	N2	V1							
13	Plot10	N0	V2							
14	Plot11	N1	V2							
15	Plot12	N2	V2							
16	Plot13	N0	V2							
17	Plot14	N1	V2							

12						Sowing date : 20/08/2023	Rapeseed			
13						Yield (Kp/ha)	Comment			
14							Soosoc failure			
17							N dose g/ha			

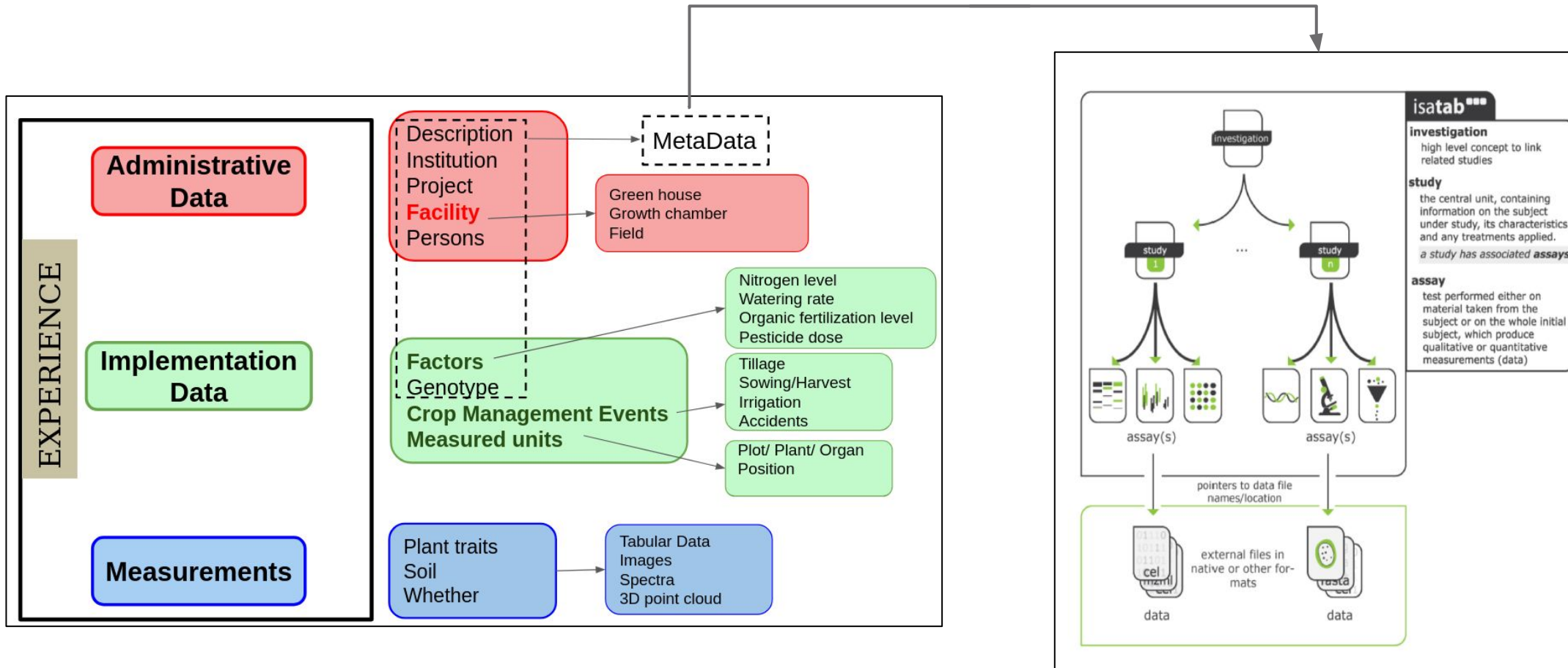
# Introduction

## Reproducible science levels

- **Re-Run:** same lab, same experiments, redo analytics (checking)
- **Repeat:** same lab, redo experiments and analytics
- **Transparency:** same experiments, other lab redo analytics (proof)
- **Replicate, Reproduce:** other lab redo experiments and analytics (robustness)
- **Reuse:** for different scientific results and/or other contexts

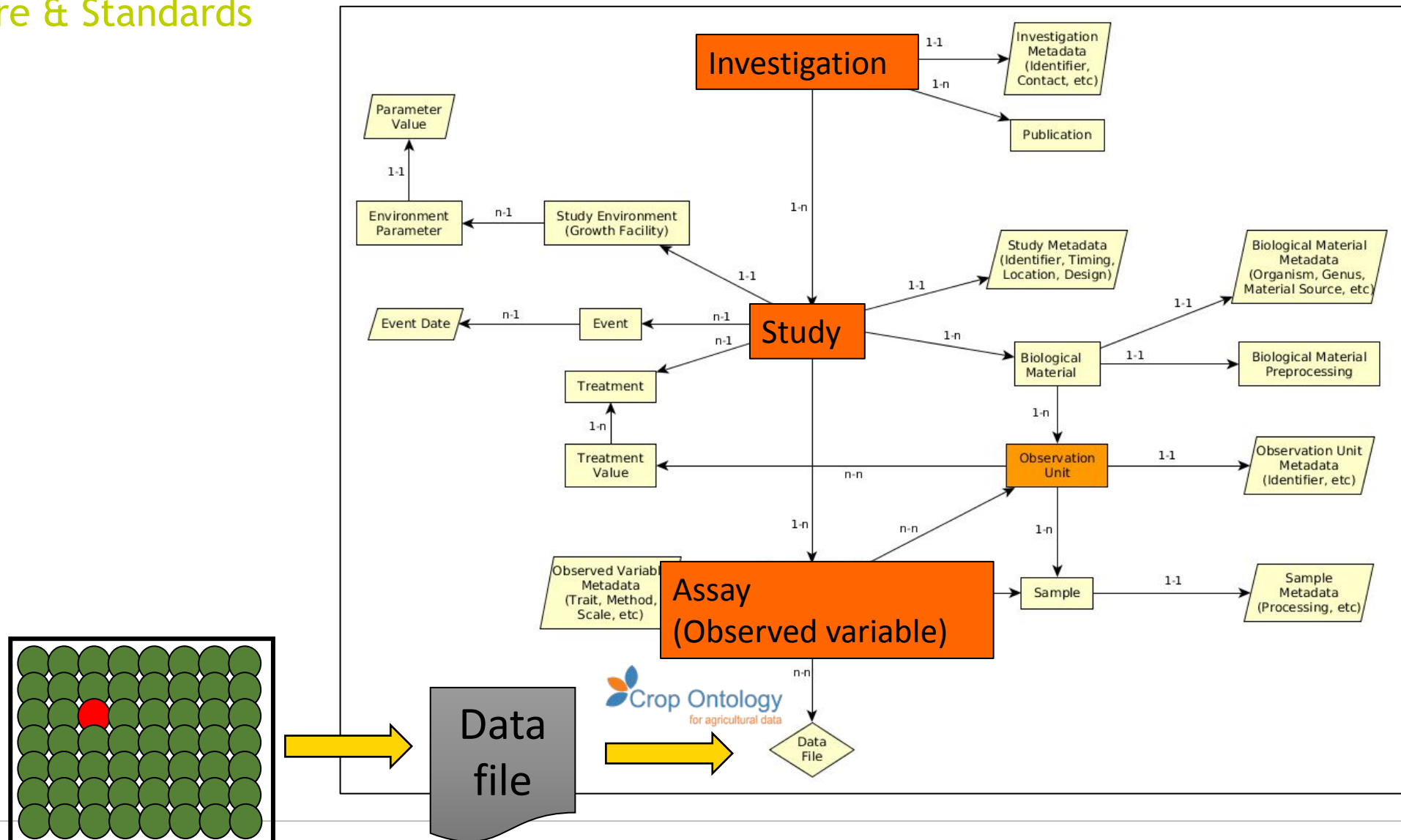
# Introduction

## Data structure



# Introduction

## Data structure & Standards

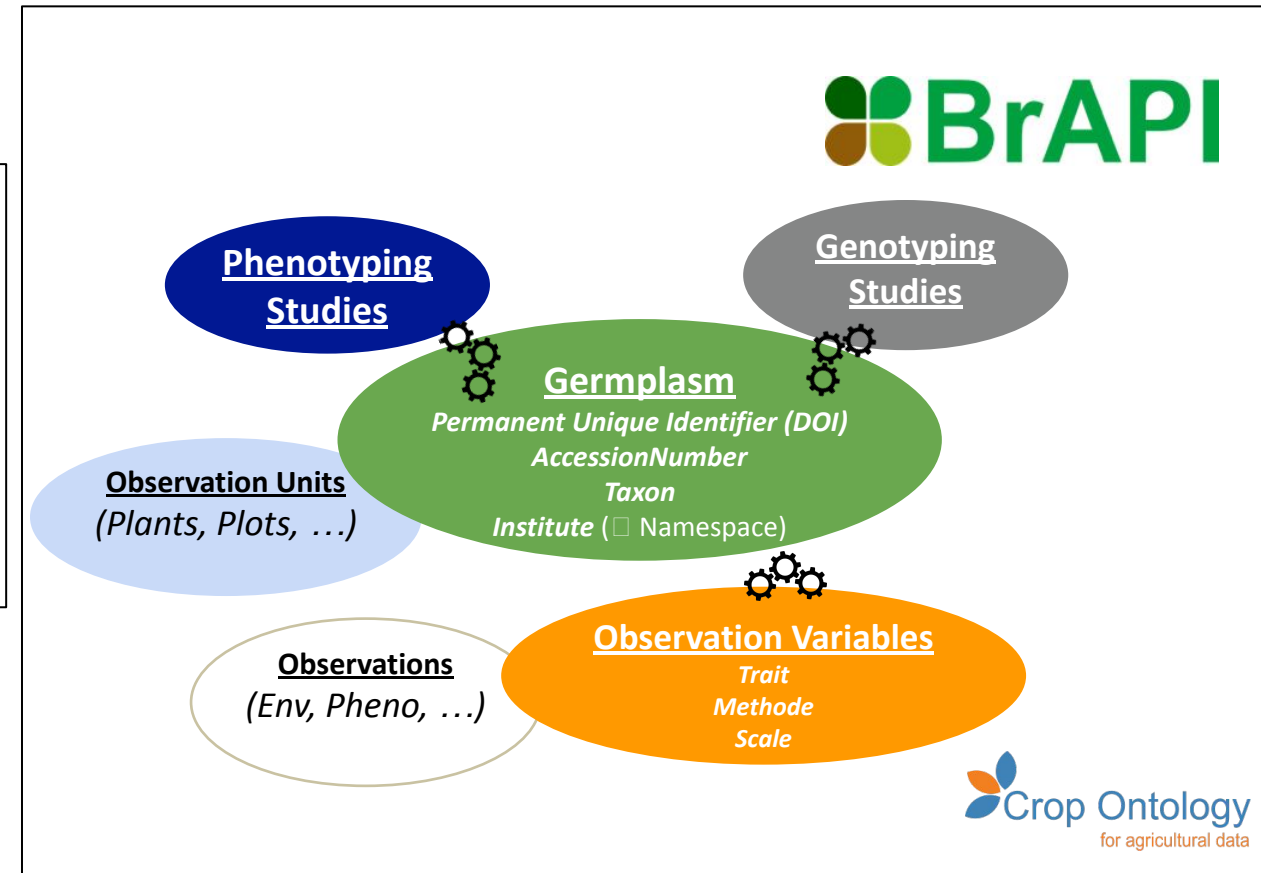
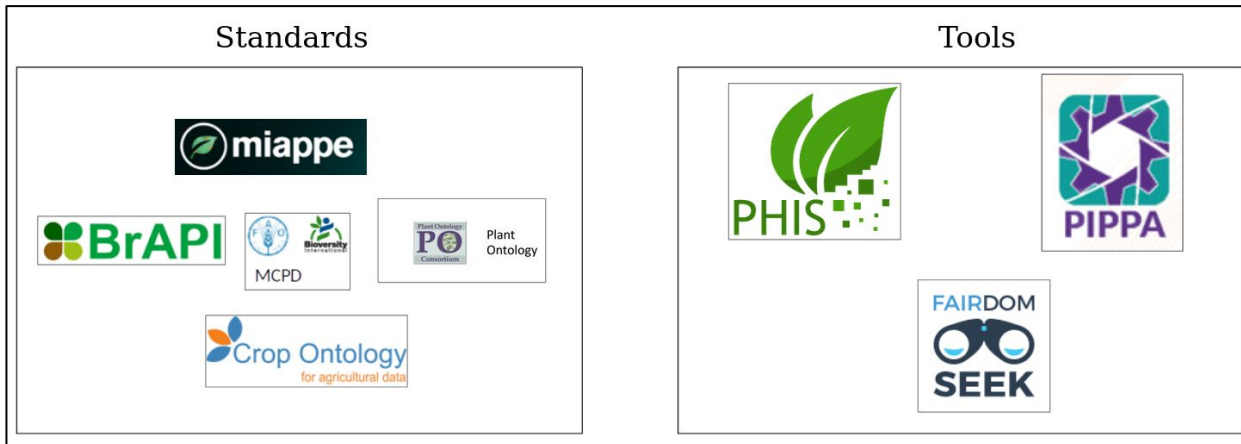




# Introduction

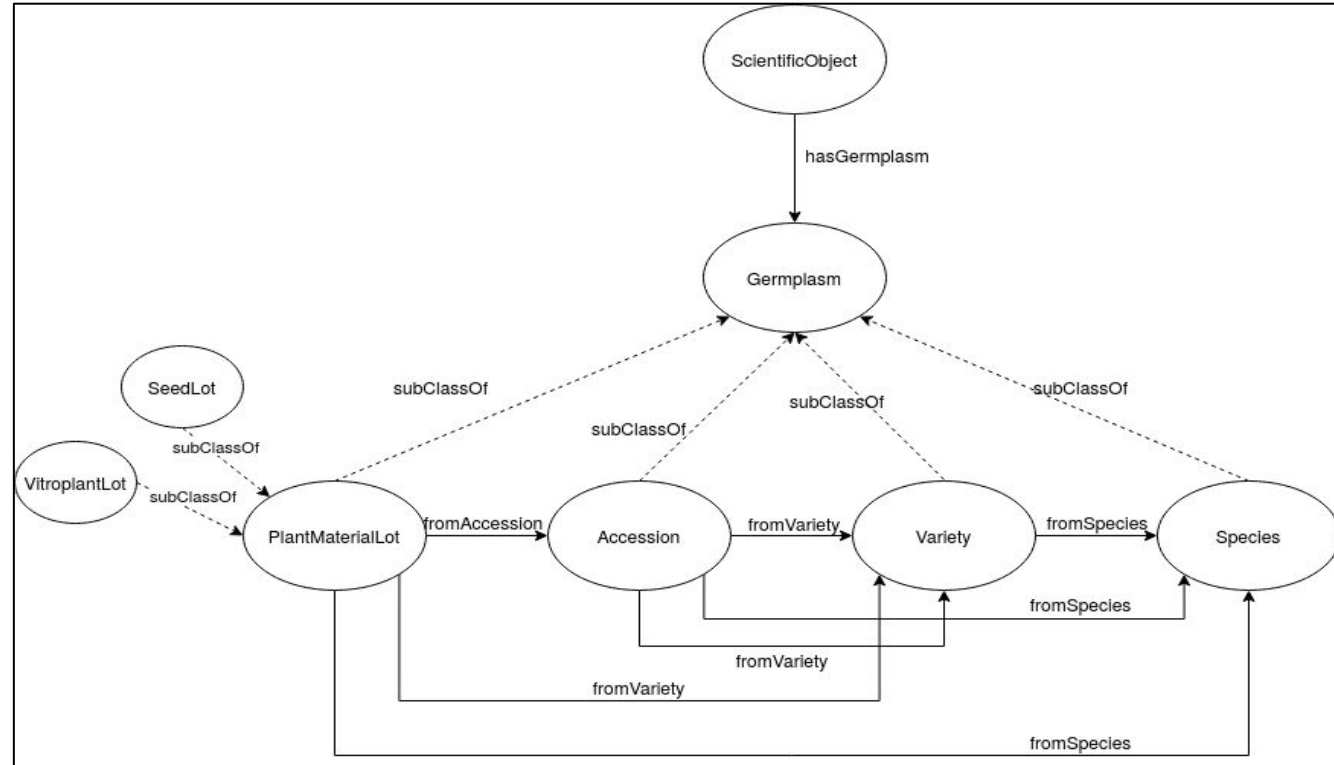
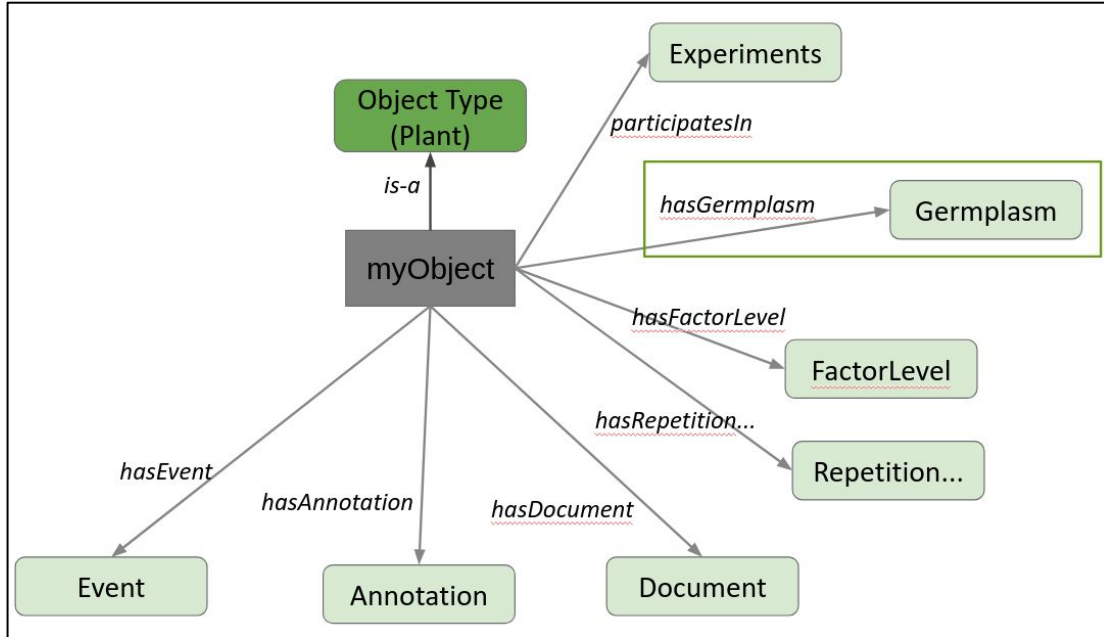
## Use Standards, Compatible tools and APIs

- Standard Open Web Service API
- Information Exchange, Main target: Breeding



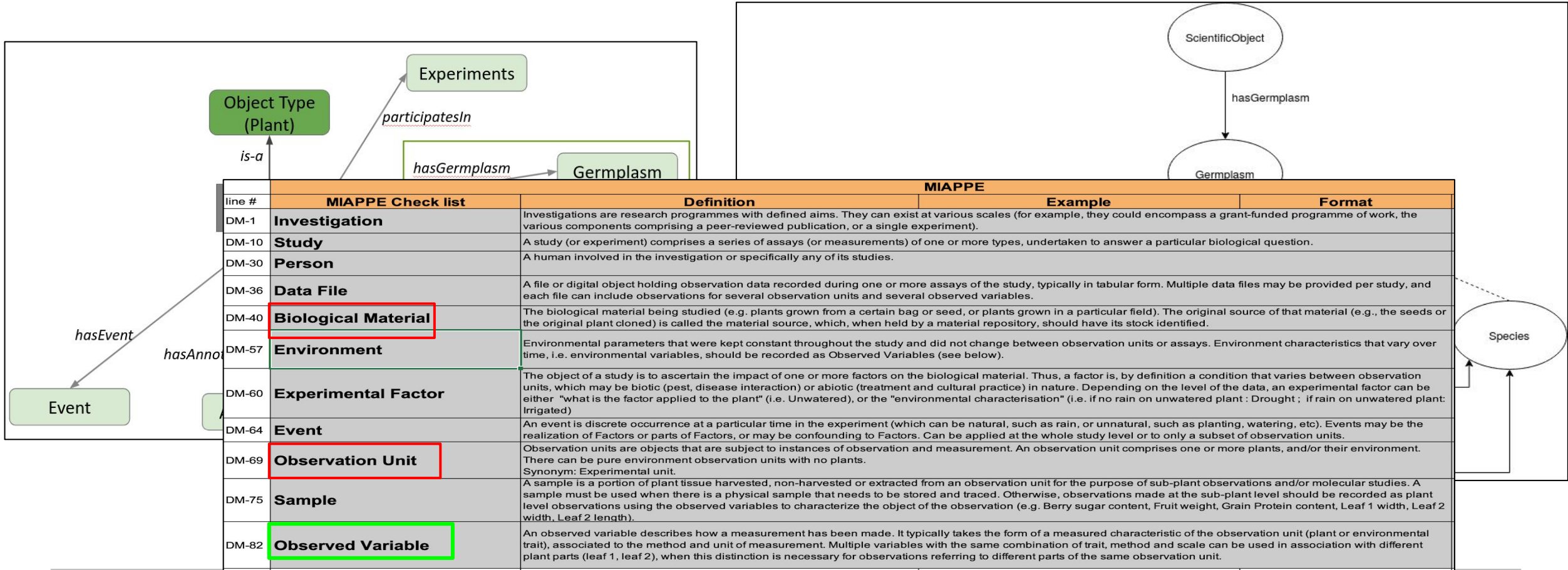
# Introduction

MIAPPE/BrAPI Compliant S.I.



# Introduction

## MIAPPE/BrAPI Compliant S.I.



# Context

Compared to 20 years ago...

- Number and diversity of data sources has increased
- Stronger complexity of data pipelines to be designed has increased

Increasing the difficulties for a reproducible and open science

Makes hard to understand how and under what conditions the data were produced

**We need to know about data provenance**

**We need to know about method of data production**

# Variables - Definition

Variable is the key element of data structuring and analytics

A variable can be represented by a combination of descriptive components giving meaning to the value derived from a data acquisition event be it an observation, a measurement, a simulation or a calculation.

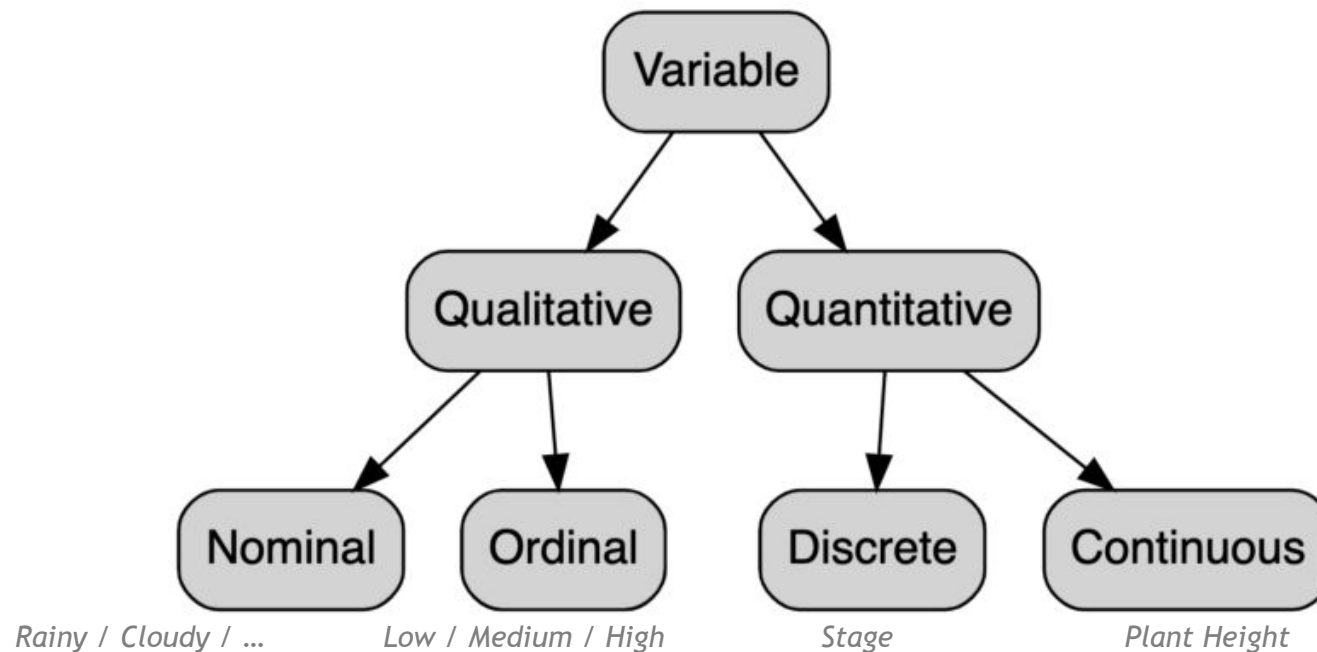


Image: Moira Sheehan

# Variables - Definition

Reality is that too often...

- No variable ID
- Same name for different variables
- Different names for same variable
- No (or not machine readable) variable description
- Unstable variable

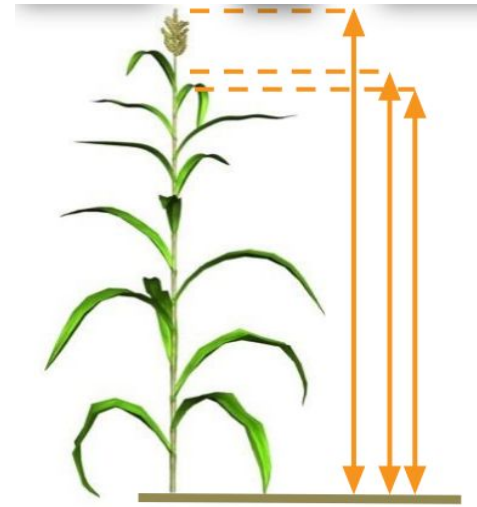


Figure From D. Pot, CIRAD

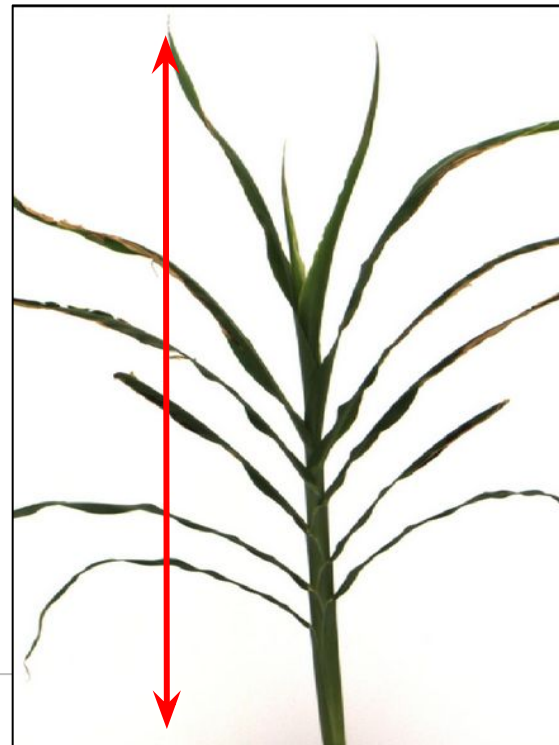


Image: Moira Sheehan

# Variables - Definition

## Formalized variable with ID...

...Aims FAIR data with a focus on interoperability and reuse

- Provides a **description**: simple, precise, unambiguous
- **Share/Reuse variables**: descriptions, structuring
- Make it easier to **aggregate and analyse** of harmonized data
- Be able to compare, merge, combine, etc.

**Covariables?**

Variables that affect a variable

Tair	size	BioMa	wt	pr
18.5	5.0	4.2	4.2	3.7
19.6	3.8	4.2	4.2	5.2
19.0	3.3	6.0	6.0	3.8
19.5	3.8	3.4	4.9	5.2
19.7	3.4	3.1	5.1	4.3
19.0	4.8	4.0	4.2	3.5
20.7	4.7	5.0	7.3	5.2

length	temp	biomass
3.79	18.55	2.93
3.62	19.16	3.32
3.83	19.79	3.41
4.19	19.87	5.38
5.14	19.12	5.33
3.66	19.16	2.35
3.84	19.03	4.74
5.74	20.27	4.55
2.90	18.66	3.75
4.65	19.52	2.53
3.50	18.66	2.27
4.06	17.56	2.98

same-as?

similar?

bias?

transform?

**Compatible individuals?**

# Variable Model

## How do you formalise the variables?

Requirements for formalising of variables:

- Focus on **WHAT, HOW, (WHERE, WHEN)**
- **In-depth understanding (area of expertise)**
- Step back from the variable, its use, its usefulness and its potential second use
- Provide a human-understandable definition



# Variable Model

We can create an infinite number of variables!

Need to formalise variables:

- that anyone can easily understand
- that can be manipulated with machine

## A Common Model in Plant Sciences

- **Trait** characteristic of an organism
- **Method** is the way we have to measure variable. Giving description of protocols and type of instruments used.
- **Unit** is an element from the Ontology of Units of Measurement

## Crop Ontology Trait Dictionary Standard

Variable = **trait** + method + scale

Study	Genotype	CIR	HT	SUR
Domaine de Valcros	6579	600	17	1
Domaine de Valcros	6580	482	14	1

What is the studied character?

Circumference of the tree

Total height of the tree

Survival state of the tree

Trait ID
Trait
Trait class
Trait description
Trait synonyms
Main trait abbreviation
Alternative trait abbreviations
Entity
Attribute
Trait status
Trait Xref

# Variable Model

We can create an infinite number of variables!

Need to formalise variables:

- that anyone can easily understand
- that can be manipulated with machine

## A Common Model in Plant Sciences

- **Entity** is the target of the variable, what we are measuring on **Trait**
- **Characteristic** is the observed property or physical quantity or quality.
- **Method** is the way we have to measure variable. Giving description of protocols and type of instruments used.
- **Unit** is an element from the Ontology of Units of Measurement

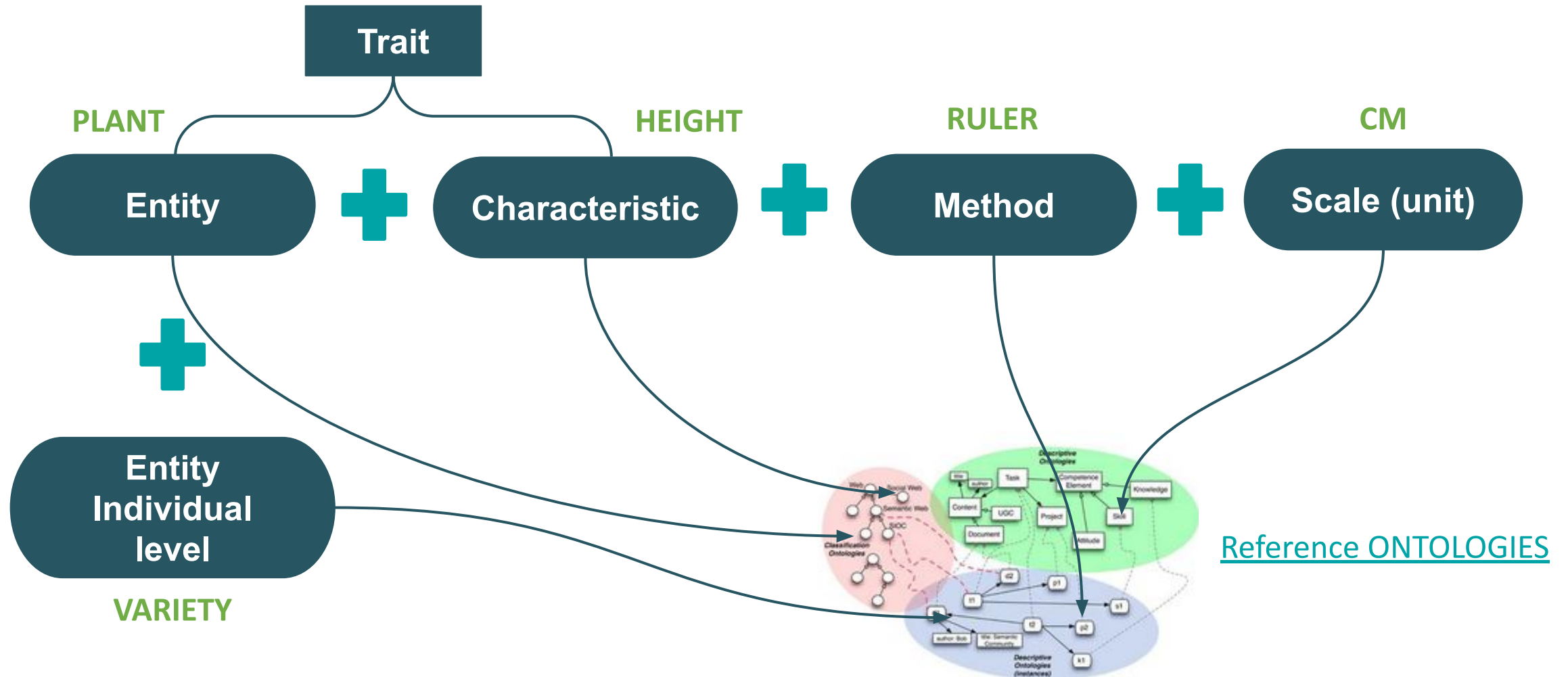
# Variable Model

## How to implement this model properly

- Enable semantically precise and **FAIR descriptions**
- Break down the description into **standardised atomic elements**
- **Link** these elements to **existing vocabularies/ontologies**
- Make the **description machine-readable**

# Variable Model

How to implement this model properly



# Variable Model

Enable semantically precise and FAIR descriptions - Machine readable

**PLANT**

**Entity**

**Entity Individual level**

**VARIETY**

### CanopyEar\_number\_counting\_PERm2

Variable

← Description Annotations Device associated Data Visualization Documents

**General informations**

**URI** [http://phenome.inrae.fr/id/variable/canopyEar\\_Number\\_c...](http://phenome.inrae.fr/id/variable/canopyEar_Number_c...)

**Name** CanopyEar\_number\_counting\_PERm2

**Alternative name** Ear\_nb\_m2

**Description**  
EN: mean ear number per m2 measured on a sample | FR: nb épis par m2 (échantillon)

**Structure**

**Entity** [canopyEar](#)

**Entity of interest** [number per area](#)

**Characteristic** [number per area](#)

**Method** [counting](#)

**Unit/Scale** [per square metre](#)

**Reference ontologies**

Relations	Reference URI
Narrower	<a href="https://croponology.org/term/CO_321:0001366">https://croponology.org/term/CO_321:0001366</a>
Exact match	<a href="https://agrovoc.fao.org/browse/agrovoc/en/page/c_8504">https://agrovoc.fao.org/browse/agrovoc/en/page/c_8504</a>

**Advanced information**

**Species** [barley](#), [bread wheat](#), [durum wheat](#)

**Data type** Decimal number

**Time interval**

**Sample interval**

**Trait uri** [https://croponology.org/term/CO\\_321:0000166](https://croponology.org/term/CO_321:0000166)

**Trait name** Ear number

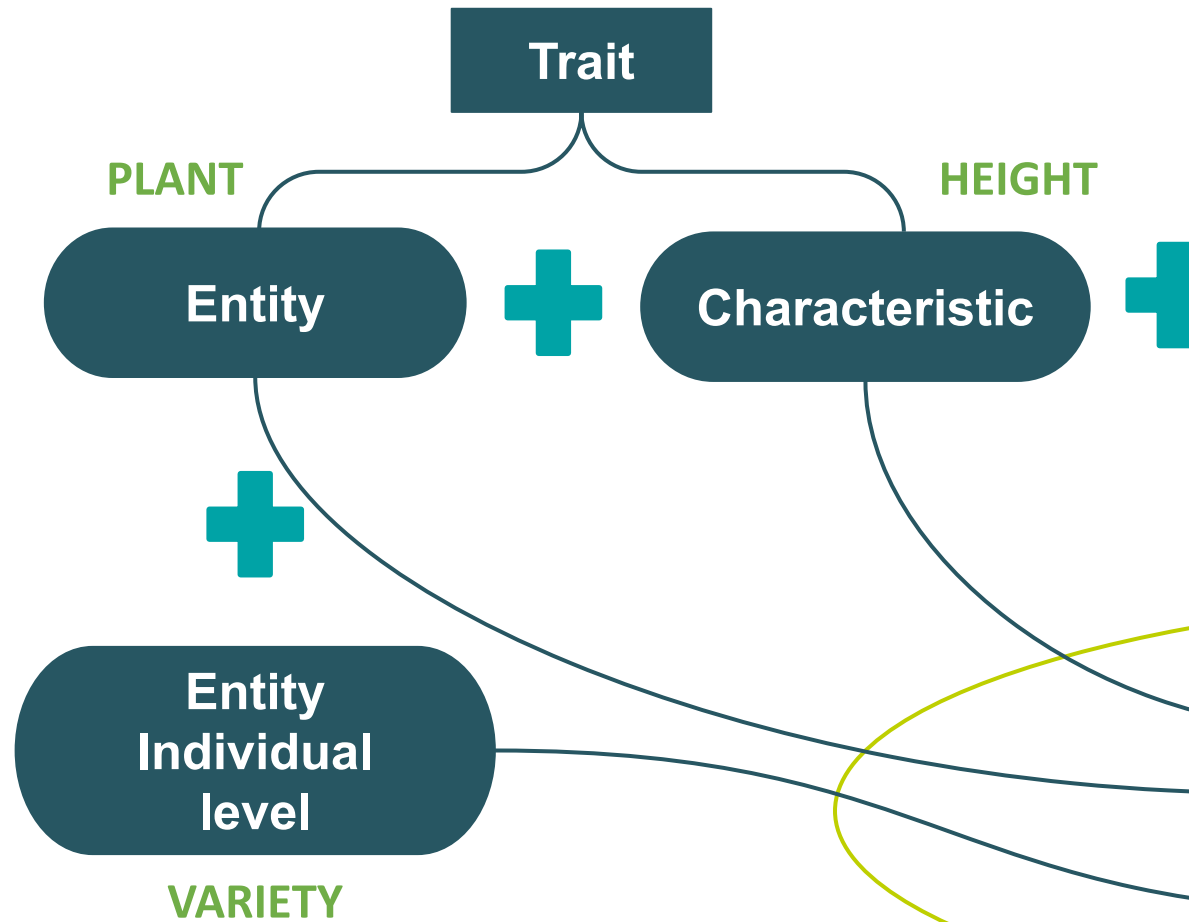
# Variable Model

Break down the description into standardised atomic elements



# Variable Model

Link these elements to existing vocabularies/ontologies



The screenshot shows the "Add trait" form in a web application. The form is titled "Add trait" and has a close button (X) in the top right corner. The form fields are:

- Trait uri** (required): A text input field containing "Height".
- Trait name** (required): A text input field containing "http://purl.obolibrary.org/obo/PATO\_0000119".
- Entity** (required): A dropdown menu with "Plant" selected.
- Characteristic** (required): A dropdown menu with "Height" selected.
- Method** (required): A dropdown menu with "Ruler" selected.
- Unit/Scale** (required): A dropdown menu with "Centimeter" selected.
- Name** (required): A text input field containing "Plant\_Height\_Ruler\_centimeter".
- Alternative name**: A text input field containing "Plant\_Height".
- Data type** (required): A dropdown menu with "Decimal number" selected.
- Time interval**: A dropdown menu with "Select an interval" selected.
- Sample interval**: A dropdown menu with "Select an interval" selected.

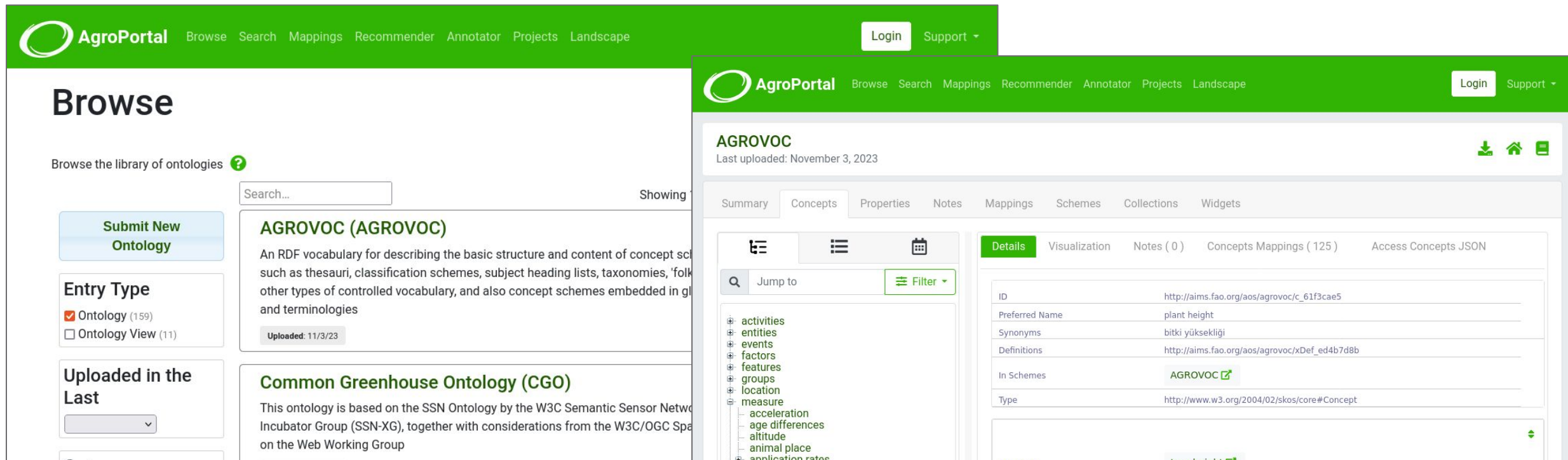
There are "Cancel" and "Save" buttons at the bottom of the form. A green button labeled "Trait already existing in an ontology" is also visible. A yellow box highlights the "Trait uri" and "Trait name" fields. A yellow question mark "How?" is overlaid on the form.

# Variable sharing

Link these elements to existing vocabularies/ontologies

Agroportal, a vocabulary and ontology repository for agronomy and related domains -

<http://agroportal.lirmm.fr/>



The screenshot displays the AgroPortal interface. The top navigation bar includes the AgroPortal logo, menu items (Browse, Search, Mappings, Recommender, Annotator, Projects, Landscape), and 'Login' and 'Support' buttons. The main content area is split into two panels. The left panel, titled 'Browse', shows a search bar and a list of ontologies. The right panel, titled 'AGROVOC', shows the details of the AGROVOC ontology, including a list of concepts and a 'Details' tab with a table of properties.

**Browse**

Browse the library of ontologies ?

Search... Showing

**Submit New Ontology**

**Entry Type**

- Ontology (159)
- Ontology View (11)

**Uploaded in the Last**

**AGROVOC (AGROVOC)**

An RDF vocabulary for describing the basic structure and content of concept schemes such as thesauri, classification schemes, subject heading lists, taxonomies, folksonomies, and other types of controlled vocabulary, and also concept schemes embedded in glossaries and terminologies

Uploaded: 11/3/23

**Common Greenhouse Ontology (CGO)**

This ontology is based on the SSN Ontology by the W3C Semantic Sensor Network Incubator Group (SSN-XG), together with considerations from the W3C/OGC Spatial Ontology Working Group on the Web Working Group

**AGROVOC**

Last updated: November 3, 2023

Summary Concepts Properties Notes Mappings Schemes Collections Widgets

Jump to Filter

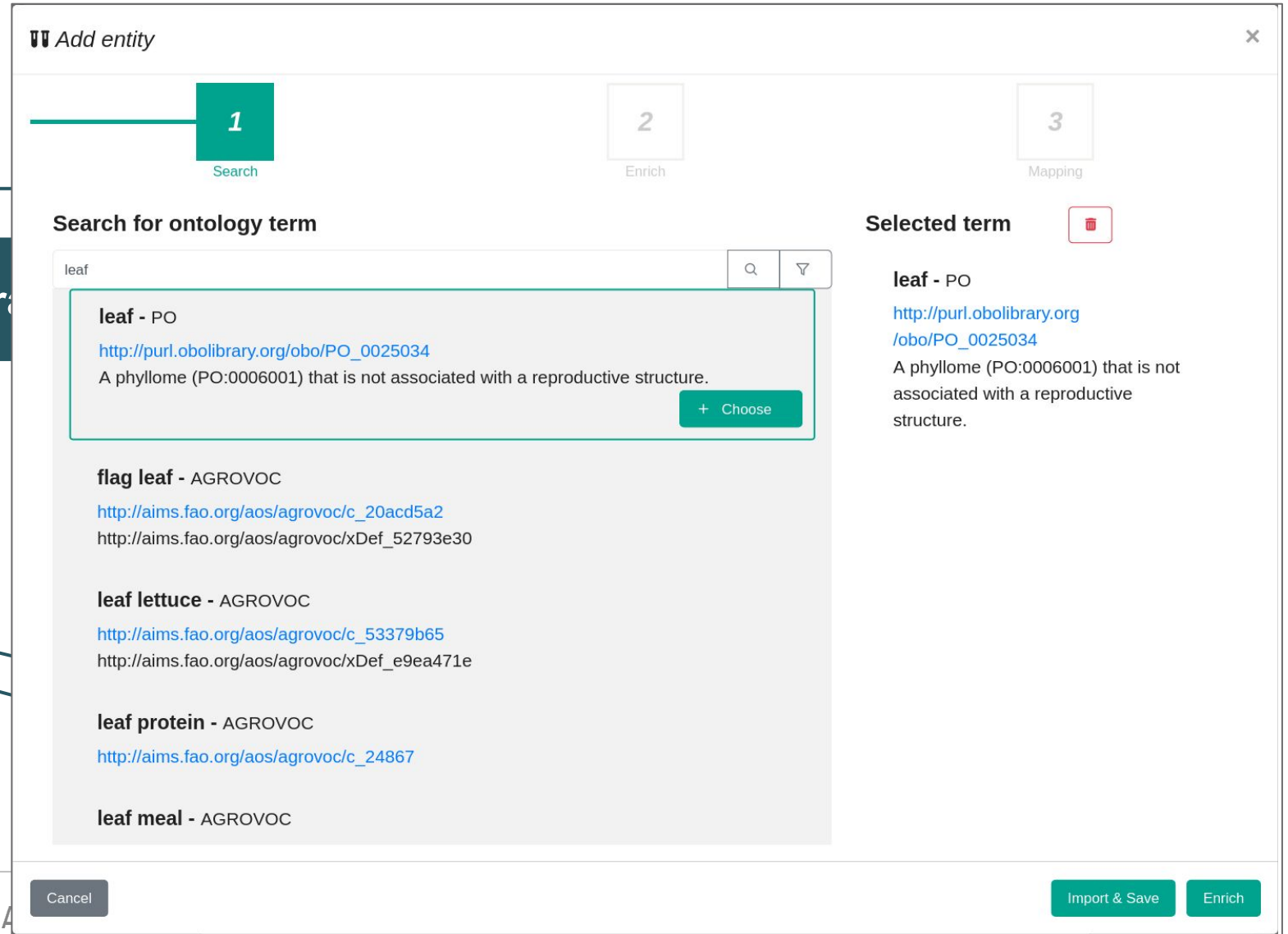
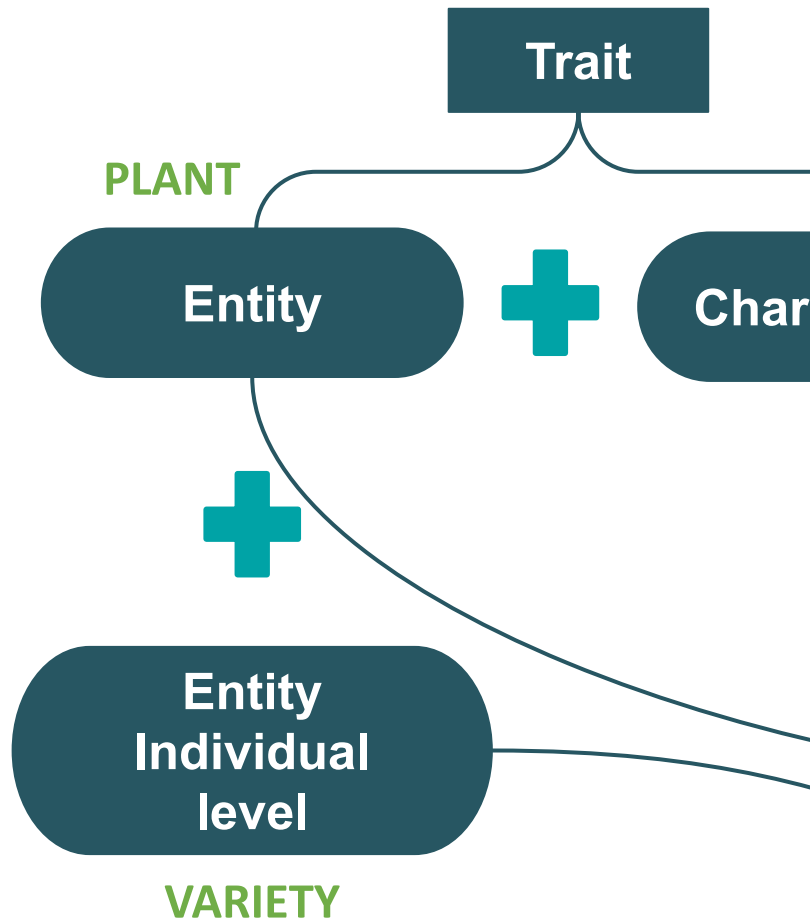
**Details** Visualization Notes (0) Concepts Mappings (125) Access Concepts JSON

ID	http://aims.fao.org/aos/agrovoc/c_61f3cae5
Preferred Name	plant height
Synonyms	bitki yüksekliği
Definitions	http://aims.fao.org/aos/agrovoc/xDef_ed4b7d8b
In Schemes	AGROVOC
Type	http://www.w3.org/2004/02/skos/core#Concept



# Describing Variables

Find an ontology concept



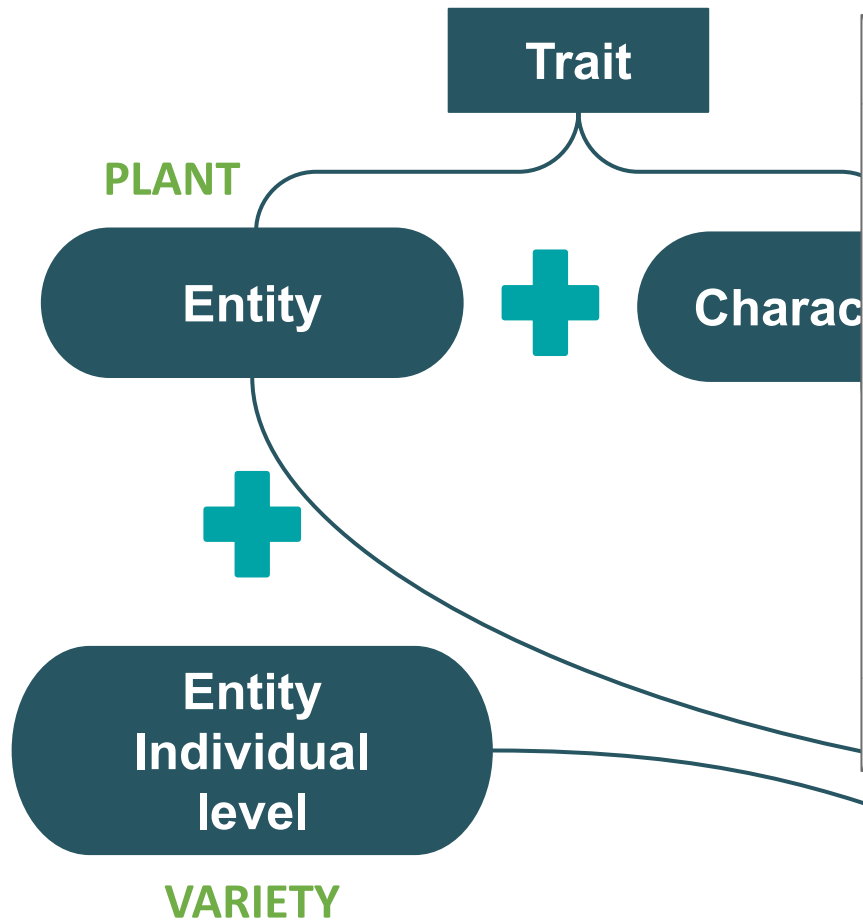
The screenshot shows the 'Add entity' interface with three steps: 1. Search, 2. Enrich, and 3. Mapping. Step 1 is active. The search input contains 'leaf'. The search results list several terms:

- leaf - PO**  
[http://purl.obolibrary.org/obo/PO\\_0025034](http://purl.obolibrary.org/obo/PO_0025034)  
A phyllome (PO:0006001) that is not associated with a reproductive structure.  
+ Choose
- flag leaf - AGROVOC**  
[http://aims.fao.org/aos/agrovoc/c\\_20acd5a2](http://aims.fao.org/aos/agrovoc/c_20acd5a2)  
[http://aims.fao.org/aos/agrovoc/xDef\\_52793e30](http://aims.fao.org/aos/agrovoc/xDef_52793e30)
- leaf lettuce - AGROVOC**  
[http://aims.fao.org/aos/agrovoc/c\\_53379b65](http://aims.fao.org/aos/agrovoc/c_53379b65)  
[http://aims.fao.org/aos/agrovoc/xDef\\_e9ea471e](http://aims.fao.org/aos/agrovoc/xDef_e9ea471e)
- leaf protein - AGROVOC**  
[http://aims.fao.org/aos/agrovoc/c\\_24867](http://aims.fao.org/aos/agrovoc/c_24867)
- leaf meal - AGROVOC**

The 'Selected term' section on the right shows the selected term: **leaf - PO** with its URL and description. At the bottom, there are buttons for 'Cancel', 'Import & Save', and 'Enrich'.

# Describing Variables

Enrich the concept



**Add entity** ✕

---

1  
Search

2  
Enrich

3  
Mapping

---

URI ?

autogenerated URI

Name \*

leaf

Description

A phyllome (PO:0006001) that is not associated with a reproductive structure.  
A phyllome vascular system that includes the totality of the portions of vascular tissue in their specific arrangement in a vascular leaf.

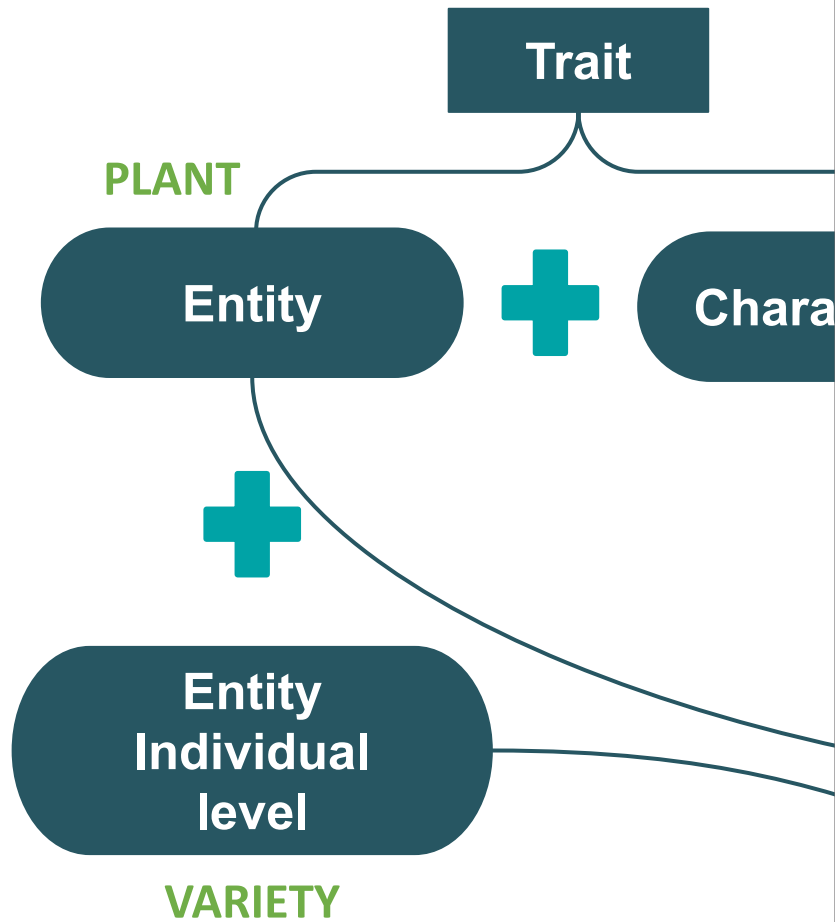
Cancel
Save
Previous
Next



Reference ONTOLOGIES

# Describing Variables

Map the concept



### Add entity

1 Search 2 Enrich 3 Mapping

**Search for mapping...**

leaf

Ontologies: OEPO and 2 more

**Selected term**

**leaf**  
A phyllome (PO:0006001) that is not associated with a reproductive structure.

Relations	Reference URI	Actions
Close match	<a href="http://purl.obolibrary.org/obo/PO_0025034">http://purl.obolibrary.org/obo/PO_0025034</a>	

**leaf - PO**  
[http://purl.obolibrary.org/obo/PO\\_0025034](http://purl.obolibrary.org/obo/PO_0025034)  
A phyllome (PO:0006001) that is not associated with a reproductive structure.

**Leaf - OEPO**  
<http://www.phenome-fppn.fr/vocabulary/2018/oeпо#Leaf>  
A phyllome vascular system (PO:0025206) that includes the totality of the portions of vascular tissue in their specific arrangement in a vascular leaf (PO:0009025).  
[database\_cross\_reference: POC:Ramona\_Walls]

**phyllode leaf - PO**  
[http://purl.obolibrary.org/obo/PO\\_0025335](http://purl.obolibrary.org/obo/PO_0025335)  
An adult ensiform leaf with a lamina that dev... median plane, rather a transverse plane, throughout the length of the leaf and is a result of increased activity of the

Map term as

- Narrower
- Broader
- Close match
- Exact match

**...Or map manually**

URI

[http://aims.fao.org/aos/agrovoc/c\\_8332](http://aims.fao.org/aos/agrovoc/c_8332) Map term as

Cancel Previous Save

# Variable sharing

## Resource Sharing Tool: Unified documentation and identification

- Centralise descriptions and identifiers
- Avoid duplicate entries and multiple sources
- Pool the work involved in describing variables and genetic resources
- Consistent management of identifiers
- Facilitate the adoption of standards

# Variable sharing

## Resource Sharing Tool: Unified

How?

- Shared File (Google drive, Sharepoint, etc.)
- List of variables published on a dataverse
- PHIS instance

<http://resources.plant-phenotyping.eu/emphasis/app/>

Variable uri	Variable ID	Variable name (long name in PHIS)	Variable description	Trait_Entity_name	Trait_Quality_name	Trait name	Trait class	Method nam
http://phenome.inrae.fr/4PId/variable/000001	4P-000001	Canopy_Height_Photogrammetry_Meter	Average Height of the canopy estimated from photogrammetry in m	Canopy	Height	Canopy_Height	morphological	Photogramme
http://phenome.inrae.fr/4PId/variable/000002	4P-000002	Canopy_HeightStd_Photogrammetry_Meter	Standard Deviation of canopy height estimated from photogrammetry in m	Canopy	HeightStd	Canopy_HeightStd	morphological	Photogramme
http://phenome.inrae.fr/4PId/variable/000003	4P-000003	Canopy_HeightFlag_Photogrammetry_Unlites	Canopy height quality flag estimated from multi or stereo vision techniques in m	Canopy	HeightFlag	Canopy_HeightFlag	morphological	Photogramme
http://phenome.inrae.fr/4PId/variable/000004	4P-000004	Soil_Height_Photogrammetry_Meter	Soil Abasse estimated from photogrammetry in m	Soil	Height	Soil_Height	morphological	Photogramme
http://phenome.inrae.fr/4PId/variable/000005	4P-000005	Canopy_SR950nm75nm_BandCombination_Unlites	Total Canopy vegetation index _ Simple Ratio (650nm,675nm), unlites	Canopy	SR	Canopy_SR	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000006	4P-000006	Canopy_CL1850nm570nm_BandCombination_Unlites	Total Canopy vegetation index _ Chlorophyll Index Green (850nm,570nm), unlites	Canopy	CLG	Canopy_CLG	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000007	4P-000007	Canopy_CL1850nm730nm_BandCombination_Unlites	Total Canopy vegetation index _ Chlorophyll Index Red Edge (850nm,730nm), unlites	Canopy	CLRE	Canopy_CLRE	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000008	4P-000008	Canopy_CL1850nm710nm_BandCombination_Unlites	Total Canopy vegetation index _ Chlorophyll Index Red Edge (850nm,710nm), unlites	Canopy	CLRE	Canopy_CLRE	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000009	4P-000009	Canopy_MCARI570nm730nm850nm_BandCombination_Unlites	Total Canopy vegetation index_Modified Chlorophyll Absorption Reflectance Index (570nm,730nm,850nm), unlites	Canopy	MCARI	Canopy_MCARI	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000010	4P-000010	Canopy_MND450nm530nm850nm_BandCombination_Unlites	Total Canopy vegetation index_Modified Normalized Difference (450nm,530nm,850nm), unlites	Canopy	mND	Canopy_mND	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000011	4P-000011	Canopy_MND450nm570nm850nm_BandCombination_Unlites	Total Canopy vegetation index_Modified Normalized Difference (450nm,570nm,850nm), unlites	Canopy	mND	Canopy_mND	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000012	4P-000012	Canopy_MND450nm675nm850nm_BandCombination_Unlites	Total Canopy vegetation index_Modified Normalized Difference (450nm,675nm,850nm), unlites	Canopy	mND	Canopy_mND	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000013	4P-000013	Canopy_MND450nm730nm850nm_BandCombination_Unlites	Total Canopy vegetation index_Modified Normalized Difference (450nm,730nm,850nm), unlites	Canopy	mND	Canopy_mND	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000014	4P-000014	Canopy_MND850nm730nm450nm_BandCombination_Unlites	Total Canopy vegetation index_Modified Normalized Difference (850nm,730nm,450nm), unlites	Canopy	mND	Canopy_mND	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000015	4P-000015	Canopy_MSR450nm850nm730nm_BandCombination_Unlites	Total Canopy vegetation index _ Modified Simple Ratio (450nm,850nm,730nm), unlites	Canopy	MSR	Canopy_MSR	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000016	4P-000016	Canopy_MTCI730nm850nm675nm_BandCombination_Unlites	Total Canopy vegetation index _ MERIS terrestrial Chlorophyll Index (730nm,850nm,675nm), unlites	Canopy	MTCI	Canopy_MTCI	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000017	4P-000017	Canopy_ND850nm70nm_BandCombination_Unlites	Total Canopy vegetation index_Normalized Difference Vegetation Index (850nm,570nm), unlites	Canopy	ND	Canopy_ND	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000018	4P-000018	Canopy_ND850nm730nm_BandCombination_Unlites	Total Canopy vegetation index_Normalized Difference Vegetation Index (850nm,730nm), unlites	Canopy	ND	Canopy_ND	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000019	4P-000019	Canopy_ND850nm675nm_BandCombination_Unlites	Total Canopy vegetation index_Normalized Difference Vegetation Index (850nm,675nm), unlites	Canopy	ND	Canopy_ND	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000020	4P-000020	Canopy_PRI570nm530nm_BandCombination_Unlites	Total Canopy vegetation index_Photchemical Reflectance Index (570nm,530nm), unlites	Canopy	PRI	Canopy_PRI	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000021	4P-000021	Canopy_InclinationAngle_PhysicalModel_Degree	Orientation of canopy elements retrieved from the inversion of radiative transfer model in decimal degrees	Canopy	InclinationAngle	Canopy_InclinationAngle	Optical	BandCombin
http://phenome.inrae.fr/4PId/variable/000022	4P-000022	Canopy_ChlorophyllContent_PhysicalModel_GramPerSquareMeter	Chlorophyll content of the canopy, in micrograms per square meters (µg/m²)	Canopy	ChlorophyllContent	Canopy_ChlorophyllContent	biochemical	PhysicalModel
http://phenome.inrae.fr/4PId/variable/000023	4P-000023	LeafGreen_ChlorophyllContent_PhysicalModel_MicroGramPerSquareCentimetre	Chlorophyll content of the green leaves, in micrograms per square centimetres (µg/cm²)	GreenLeaf	ChlorophyllContent	GreenLeaf_ChlorophyllContent	biochemical	PhysicalModel
http://phenome.inrae.fr/4PId/variable/000024	4P-000024	CanopyGreen_CoverFraction0deg_PhysicalModel_Unlites	Canopy Green Cover fraction in the nadir direction retrieved from the inversion of radiative transfer model (unlites)	GreenCanopy	CoverFraction	GreenCanopy_CoverFraction	morphological	PhysicalModel
http://phenome.inrae.fr/4PId/variable/000025	4P-000025	CanopyGreen_CoverFraction45deg_ImageSegmentation_Unlites	Canopy Green Cover fraction in the nadir direction retrieved from the segmentation of a RGB image (unlites)	GreenCanopy	CoverFraction	GreenCanopy_CoverFraction	morphological	ImageProces
http://phenome.inrae.fr/4PId/variable/000026	4P-000026	CanopyGreen_CoverFraction45deg_ImageSegmentation_Unlites	Canopy Green Cover fraction at 45° retrieved from the segmentation of a RGB image (unlites)	GreenCanopy	CoverFraction	GreenCanopy_CoverFraction	morphological	ImageProces
http://phenome.inrae.fr/4PId/variable/000027	4P-000027	CanopyGreen_FIPARWhiteSky_PhysicalModel_Unlites	Canopy fraction of intercepted diffuse (white sky) PAR (Photosynthetically Active Radiation - 400nm-700nm) absorption retrieved from the inversion of radiative transfer model (unlites)	GreenCanopy	FIPAR	GreenCanopy_FIPAR	morphological	PhysicalModel
http://phenome.inrae.fr/4PId/variable/000028	4P-000028	CanopyGreen_FIPARBlackSky_PhysicalModel_Unlites	Canopy fraction of intercepted PAR absorption in a given direction (black sky) retrieved from the inversion of radiative transfer model (unlites)	GreenCanopy	FIPAR	GreenCanopy_FIPAR	morphological	PhysicalModel
http://phenome.inrae.fr/4PId/variable/000029	4P-000029	CanopyGreen_FIPARDaily_PhysicalModel_Unlites	Daily integrated value of the canopy fraction of intercepted PAR absorption retrieved from the inversion of radiative transfer model (unlites)	GreenCanopy	FIPAR	GreenCanopy_FIPAR	morphological	PhysicalModel
http://phenome.inrae.fr/4PId/variable/000030	4P-000030	CanopyGreen_EffectiveAreaIndex_PhysicalModel_Unlites	Canopy Green Area Index retrieved from the inversion of radiative transfer model (unlites)	GreenCanopy	Area	GreenCanopy_Area	morphological	PhysicalModel
http://phenome.inrae.fr/4PId/variable/000031	4P-000031	Canopy_Reflectance450nm_Calibration_Unlites	Mean canopy reflectance in band 450nm (unlites)	Canopy	Reflectance	Canopy_Reflectance	Optical	ImageProces
http://phenome.inrae.fr/4PId/variable/000032	4P-000032	Canopy_Reflectance450nmStd_Calibration_Unlites	Standard Deviation of canopy reflectance in band 450nm (unlites)	Canopy	Reflectance	Canopy_Reflectance	Optical	ImageProces
http://phenome.inrae.fr/4PId/variable/000033	4P-000033	Canopy_Reflectance530nm_Calibration_Unlites	Mean canopy reflectance in band 530nm (unlites)	Canopy	Reflectance	Canopy_Reflectance	Optical	ImageProces

recherche.data.gov.fr

Submit / Publish a dataset EN

About The Ecosystem members Data Online help News

Home > A multidisciplinary repository

### A multidisciplinary repository

Updated at: 05/12/2023

The multidisciplinary repository is a sovereign publishing solution for sharing and opening up data for communities which are yet to set up their own recognised thematic repository.



EMPHAS

Variables

Germplasm

Tools

Web API

Name	Entity	Entity of interest	Characteristic	Method	Unit/Scale	Actions
awn_type_estimation_1to3 AwnType_1to3	awn		type	estimation	Score[1-3]	[Eye icon]
awn_type_estimation_presORabs AwnType_presAbs	awn		type	estimation	Presence-absence	[Eye icon]
canopy_FCover0_estimation_1to5	canopy		CoverFraction0deg	estimation	Score[1-5]	[Eye icon]
canopy_FCover0_estimation_1to9	canopy		CoverFraction0deg	estimation	score[1-9]	[Eye icon]
canopy_FCover0_estimation_percent	canopy		CoverFraction0deg	estimation	percent	[Eye icon]
canopy_height_estimation_cm Plant_height	canopy		height	estimation	centimetre	[Eye icon]
canopy_standEstablishment_estimation_1to5	canopy		stand establishment	estimation	Score[1-5]	[Eye icon]

# Variables & Provenance: Understanding Data

Building credibility and trust in data, analysis and reporting

- Method is associated to Variable
- Provenance is associated to Data
  
- Variable method: protocol and type of thermometer (not compatible values)
- Provenance: name of the nurse or the manufacturer of the thermometer



# Variable definition - Special focus on method

Building credibility and trust in data, analysis and reporting

Trait = Plant Height

- Method 1 : Manual measurement
  - Method 2 : Image processing
- => 2 different and non-comparable variables!



Image: Moira Sheehan

# Whats is Provenance ?

That is metadata to know...

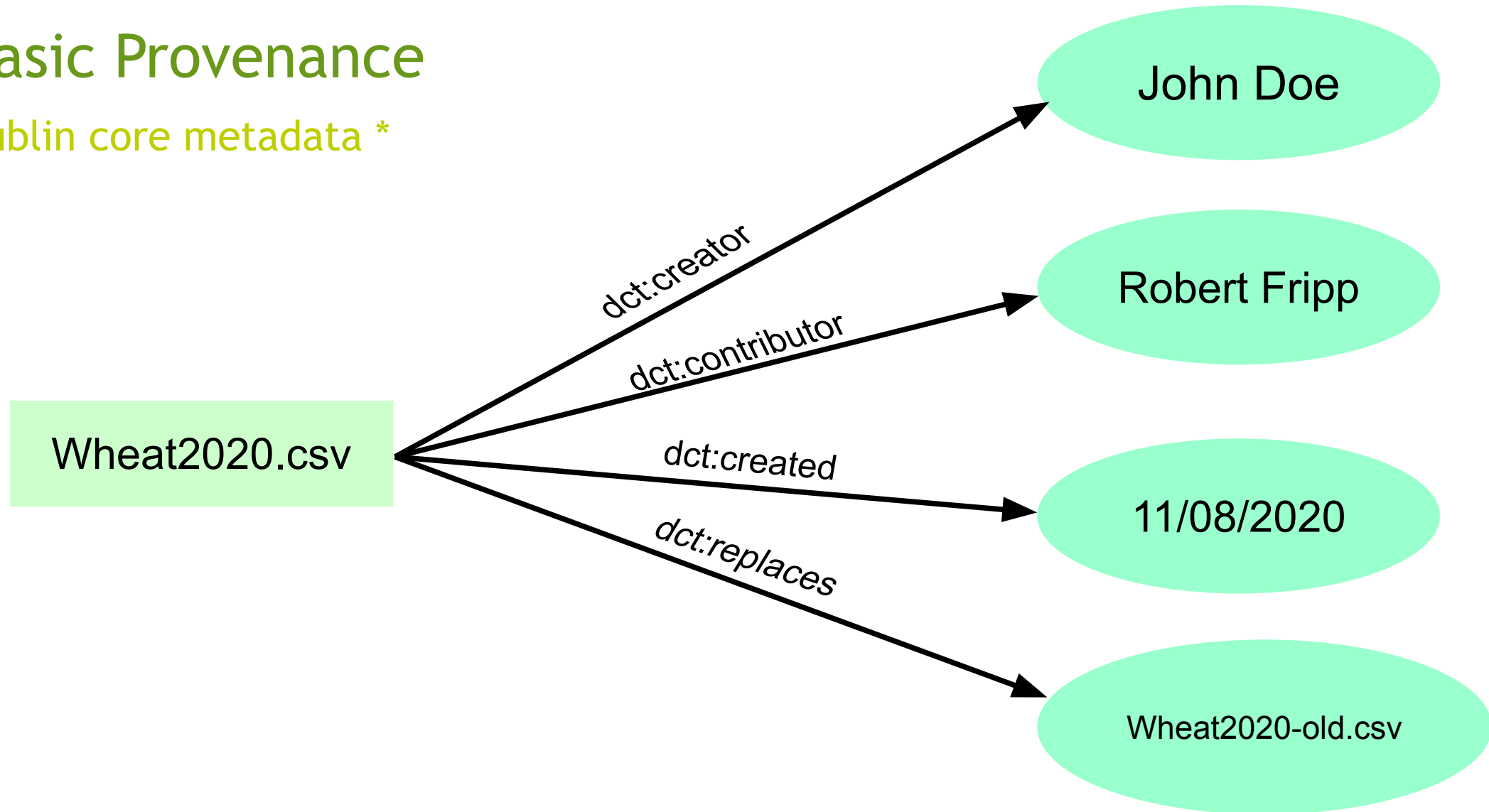
- Who played a role when creating the data?
- Who contributed to the data?
- How data was transformed?
- Which tools were used?
- When and how data were produced?
- etc.





# Basic Provenance

Dublin core metadata \*

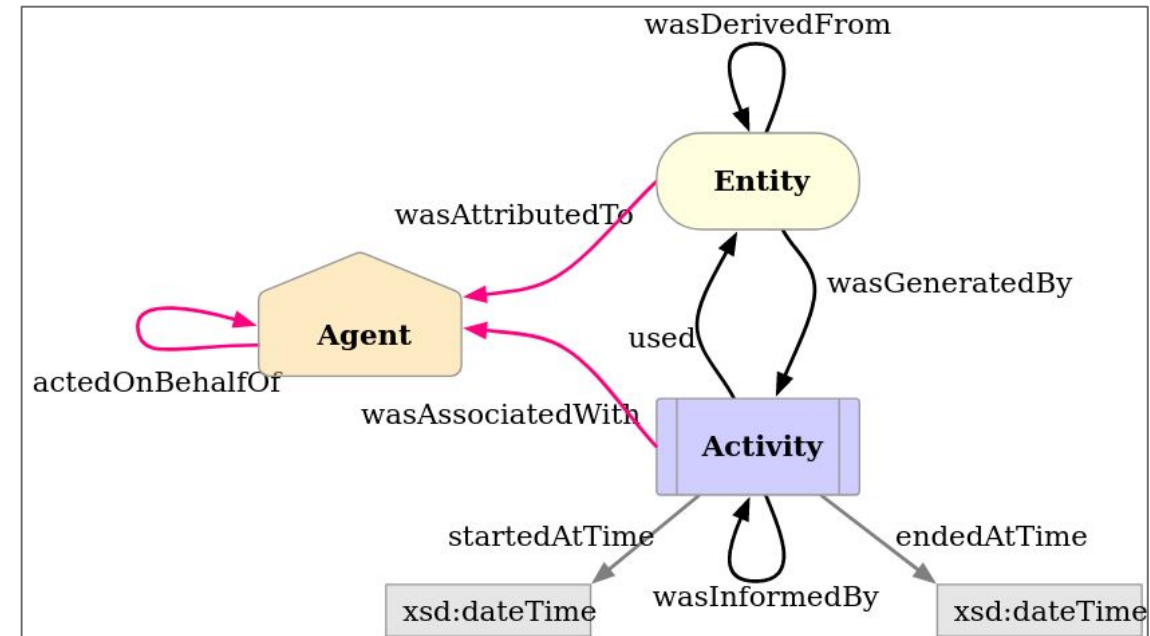


\* <https://www.dublincore.org/specifications/dublin-core/dcmi-terms>

# Advanced Provenance

## Representation model for provenance (PROV-O \*)

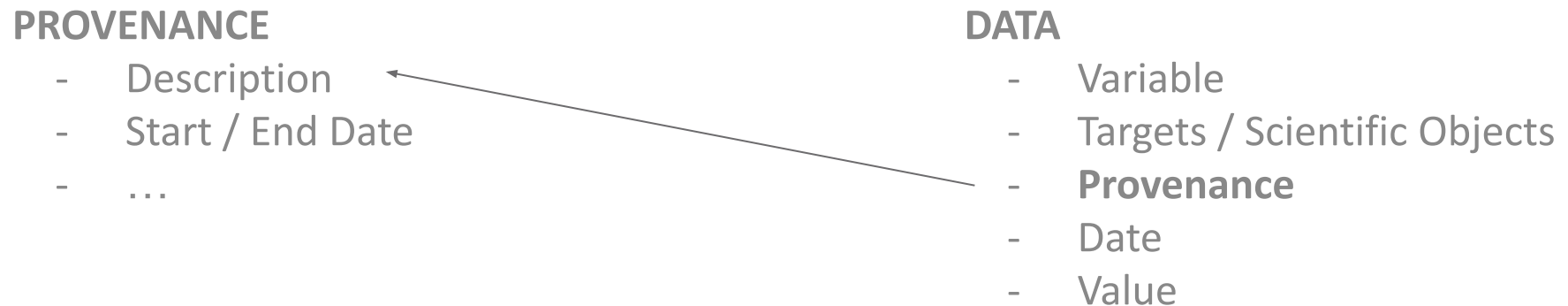
- **Entity:** data, data sources, documents, results, etc.  
Entity can be input or output
- **Agent:** person, software, Web services, institution, company, etc.  
Agent bears the responsibility
- **Activity:** generating, transforming, modifying, processing, etc.  
Activity occurs over a period of time and acts upon or with entities



\* <https://www.w3.org/TR/prov-o/>

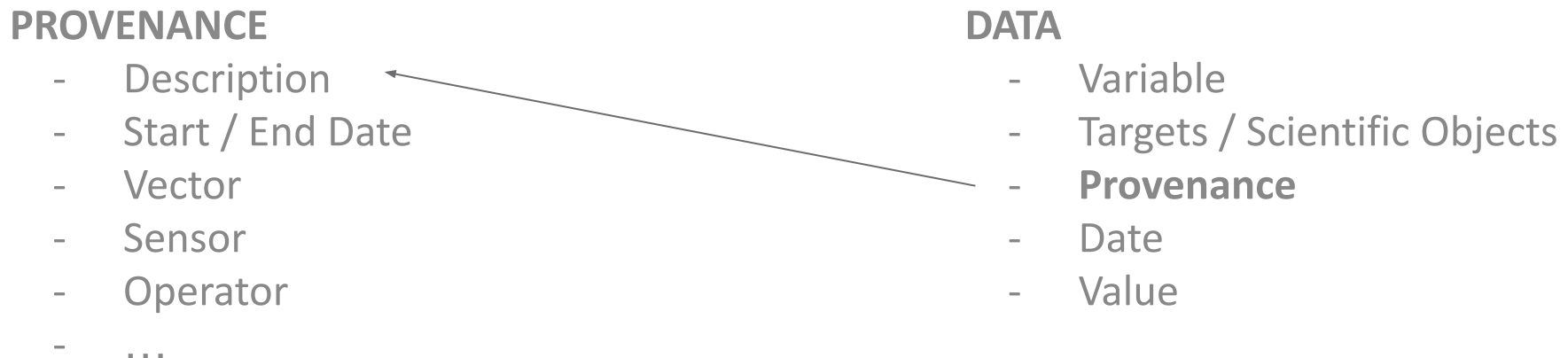
# Provenance: Data Acquisition description

- **Provenance (Activity)**
  - Formalisation of the transformation of raw data into processed data
  - Provenance will describe a process, a step in a data acquisition workflow
- **Datafiles and Data (Entity)**
  - Single item of factual information of variable complexity (e.g. a length or an image) resulting from measurement, observation or processing.



# Provenance: Data Acquisition description

- **Provenance (Activity)**
  - Formalisation of the transformation of raw data into processed data
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- **Datafiles and Data (Entity)**
  - Single item of factual information of variable complexity (e.g. a length or an image) resulting from measurement, observation or processing.
- **Devices (Agent)**
  - Sensors
  - Vectors
  - Softwares



# Devices

## Description

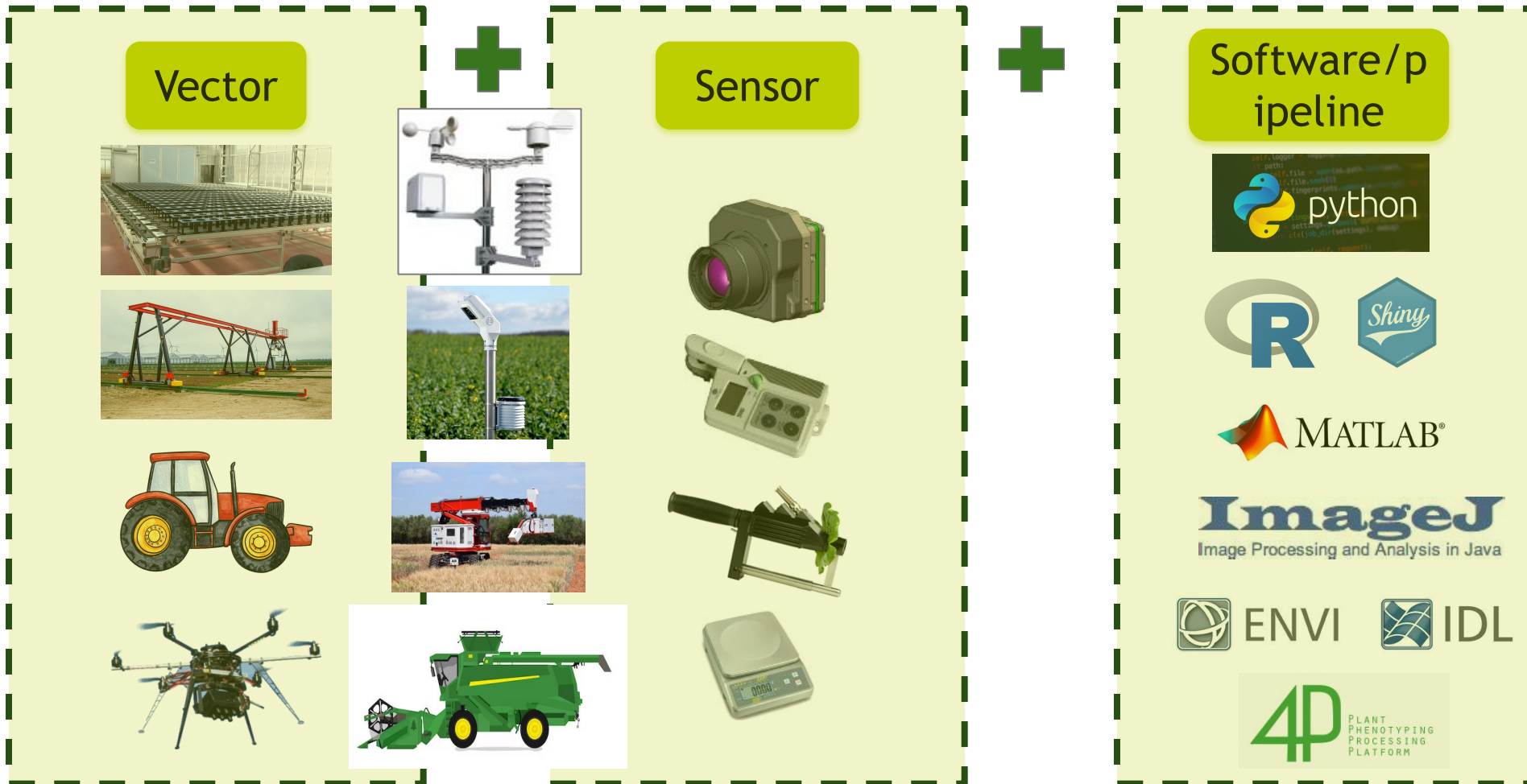
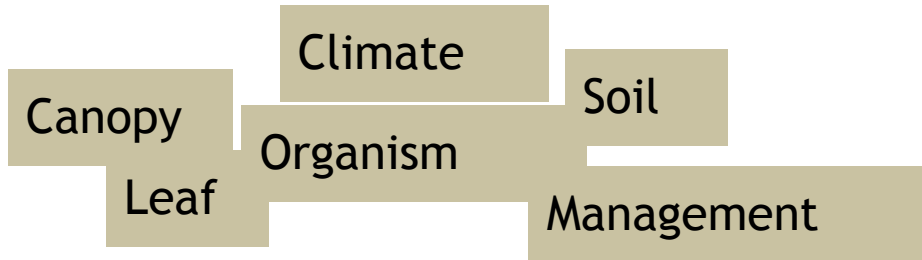
**Device** is any equipment to measure or control or calculate a property of an object, including also any equipment holding or carrying another one.

*Ex.:*

- *Camera, Thermometer, Software, Spectrometer*
- *UAV, Field Robots Conveyor*
- *Weather Station, Acquisition station*

# Devices Examples

Phenotyping & envirotyping



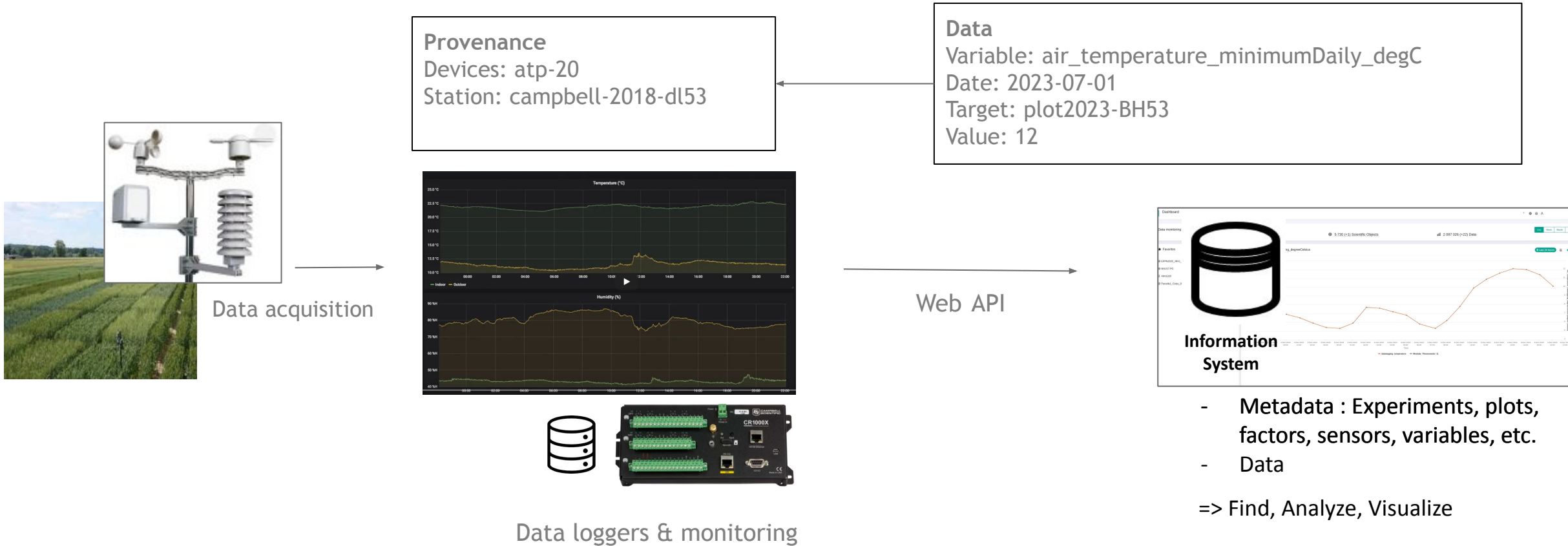
# Datafiles and data

## Description

- **Data:** Data is a single item of factual information of variable complexity (e.g. a length or an image) resulting from measurement, observation or processing. This element can be used as input or output by specialised IT tools.
- **Tabular Data:** : Scalar data.
- **Datafile / Complex Data:** Complex data often serialized in files (e.g. image file, spectrum output, orthomosaic, etc.)

# Environmental Data Acquisition

## Automatized data acquisition from sensors





# Phenotyping Data Acquisition

## Measuring Plant Height

Target: Observation unit / Scientific object

**URI of plant:**  
<<http://phenome.fr/arch/2017/c17000118>>

**URI of pot:**  
<<http://phenome.fr/arch/2013/pc13001542>>

**URI of cart:** 108  
<<http://phenome.fr/arch/2013/ct1300123>>

**URI of cabin:**  
<<http://phenome.fr/arch/2018/ac180015>>

Device

**URI of camera:**  
<<http://phenome.fr/arch/2018/ac180019>>

Datafile

**URI of image:** <<http://phenome.fr/arch/2017/ic17002295855>>



# Phenotyping Data Acquisition

## Measuring Plant Height

### Step 1: Image Acquisition

#### Provenance Datafile

- Image Acquisition
- Acquisition from 2017-06-01 to 2017-07-15
- Device: camera *ac180019*
  - Type: RGB Camera
  - Settings: Side View

### Step 2: Data Acquisition

#### Provenance Data

- Image Processing
- Acquisition from 2017-06-01 to 2017-07-15
- Device: camera *ac180019*
  - Settings: Side View
- Software: Image processing software
  - Settings: version 1.1.5
- Processing Datafile *ic17002295855*

Target: Observation unit / Scientific object

URI of plant:  
<<http://phenome.fr/arch/2017/c17000118>>

Software

URI of software:  
<<http://phenome.fr/arch/2018/sft019>>



108

Image processing software

Device

URI of camera:  
<<http://phenome.fr/arch/2018/ac180019>>

Datafile

URI of image: <<http://phenome.fr/arch/2017/ic17002295855>>



Trait = Plant Height

Variable =

- Plant
- Height
- Image Processing
- Centimetre

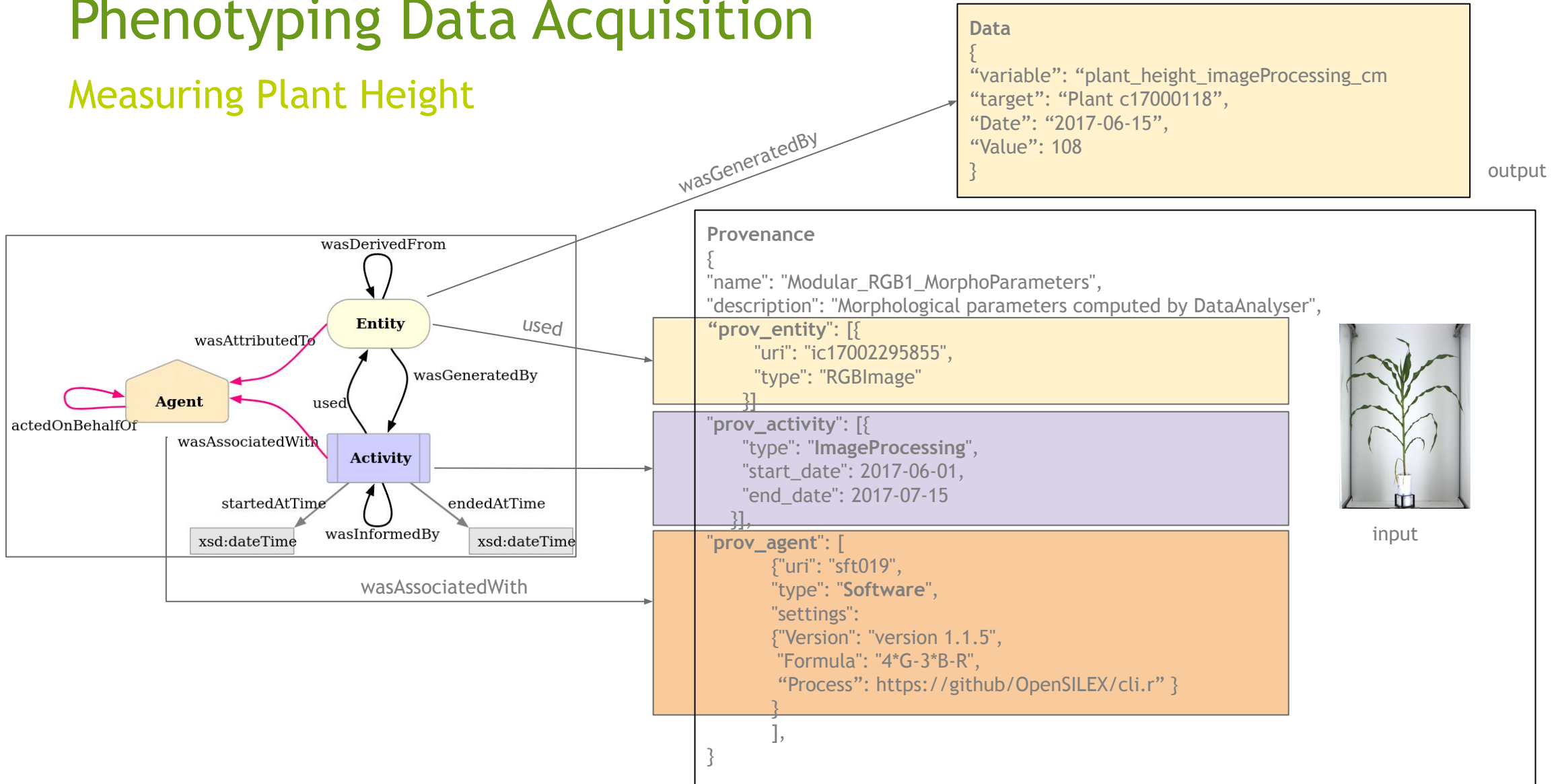
URI of Variable:  
<[https://emphasis/id/variable/plant\\_height\\_imageProcessing\\_cm](https://emphasis/id/variable/plant_height_imageProcessing_cm)>

Data

- Variable: *plant\_height\_imageProcessing\_cm*
- Target: Plant *c17000118*
- Date: 2017-06-15
- Value: 108

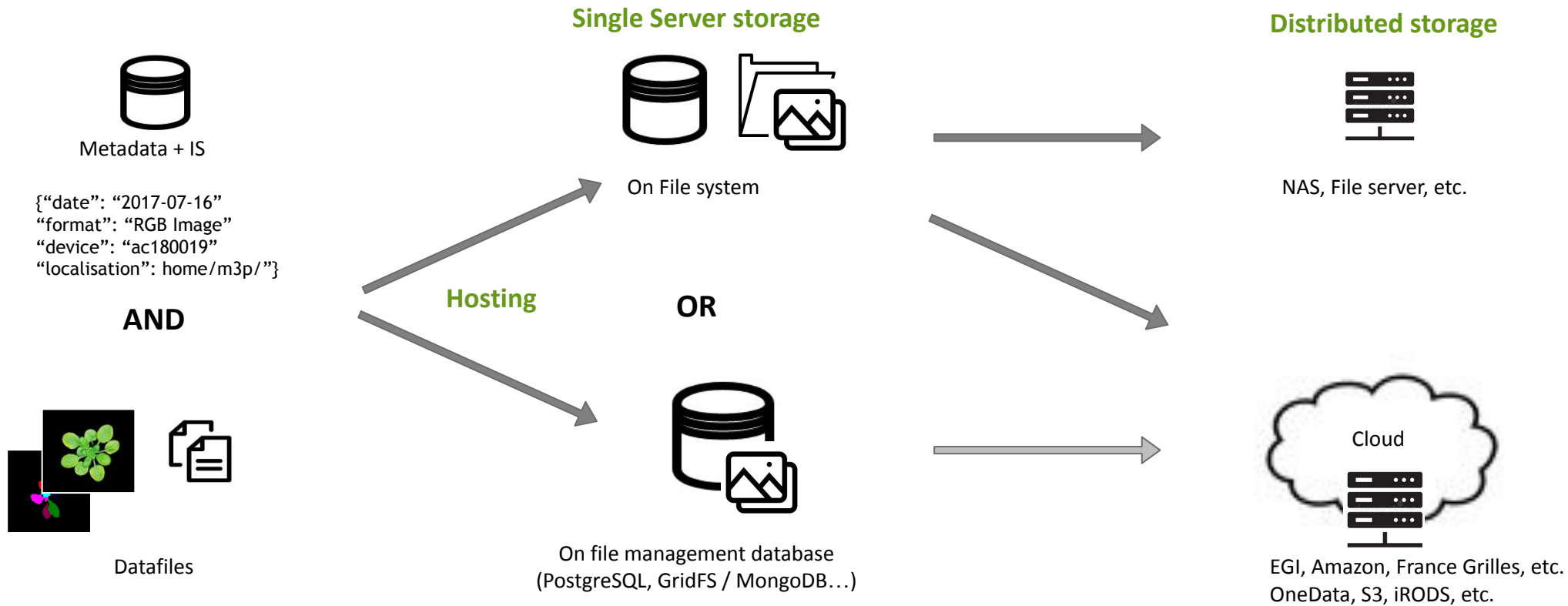
# Phenotyping Data Acquisition

## Measuring Plant Height



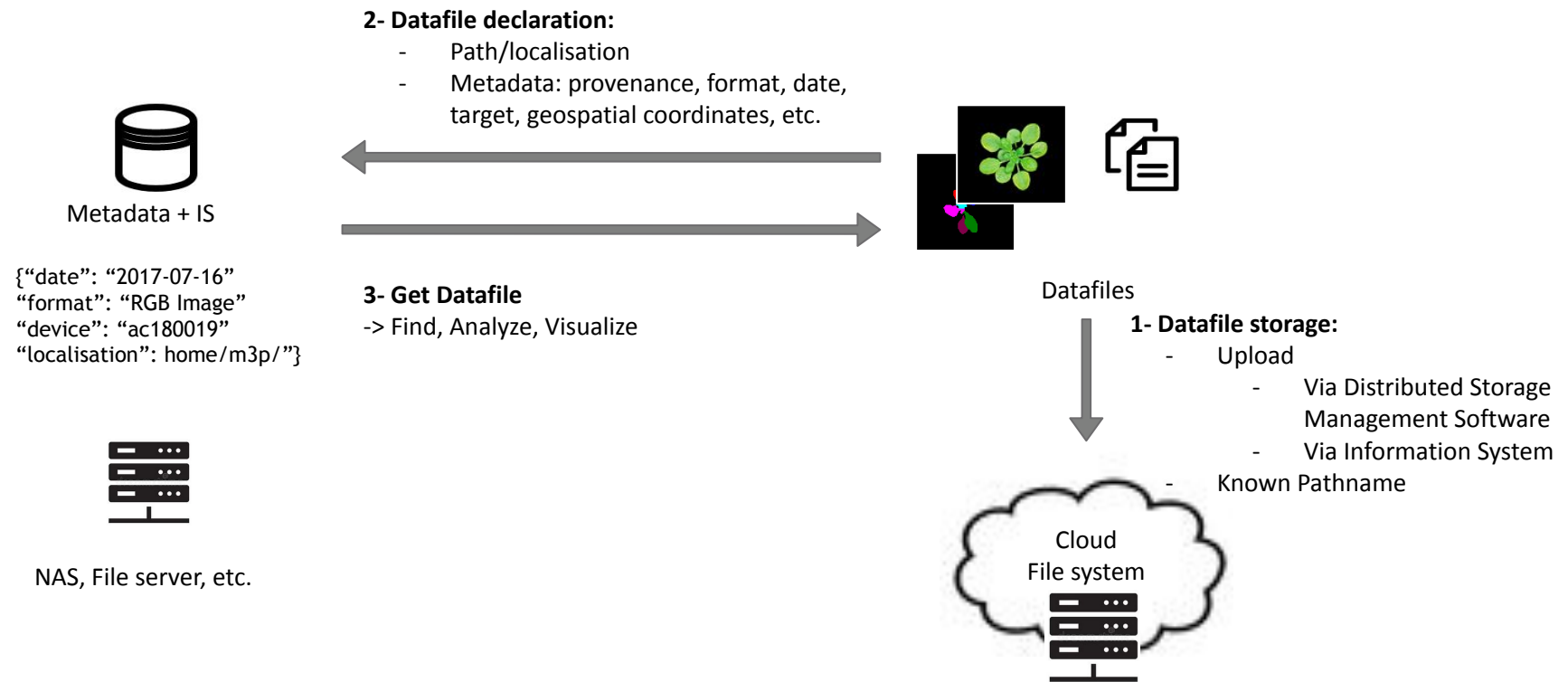
# Datafiles storage & link to Information System

## Different storage solutions



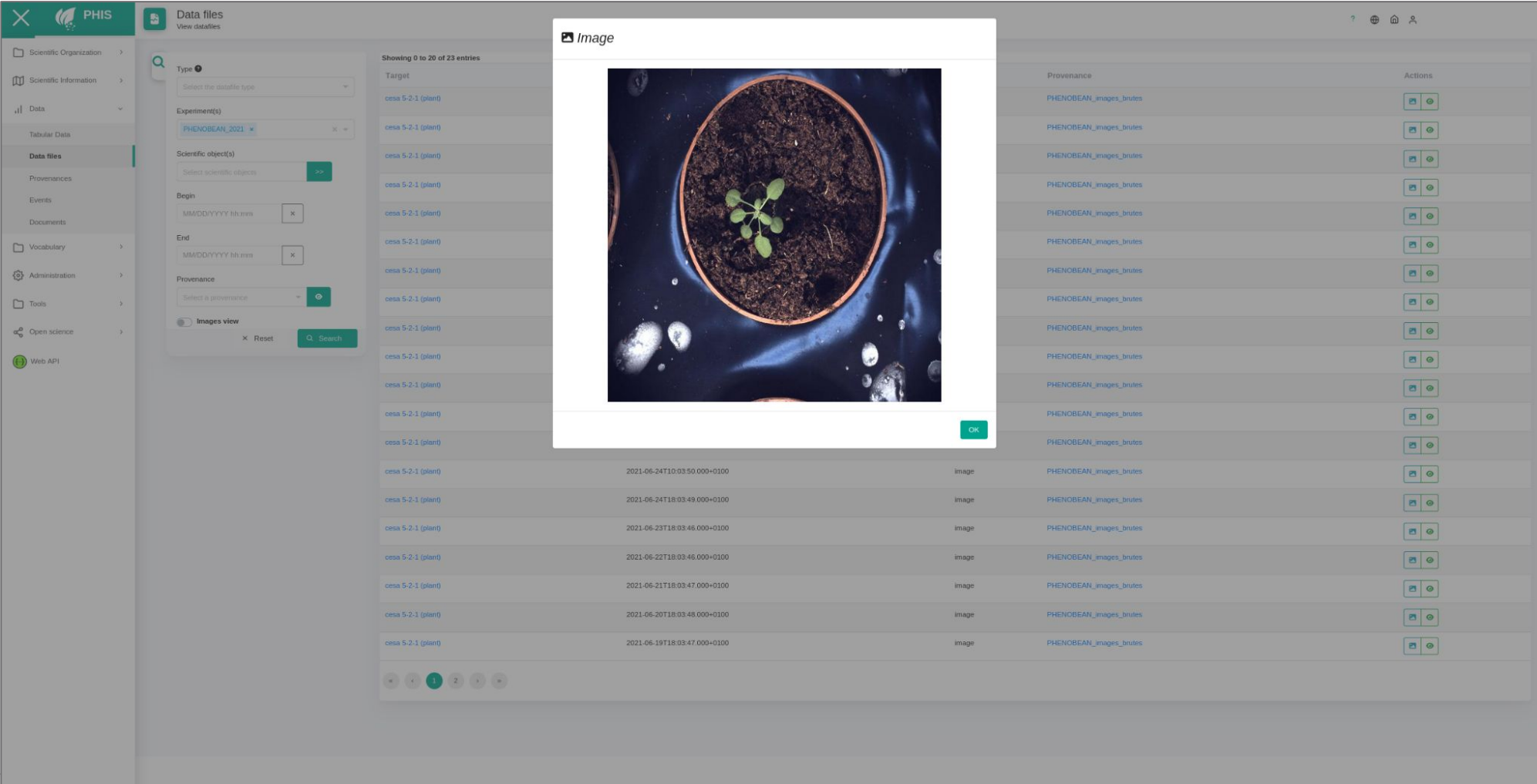
# Datafiles storage & link to Information System

## Implementation Example



# Datafiles storage & link to Information System

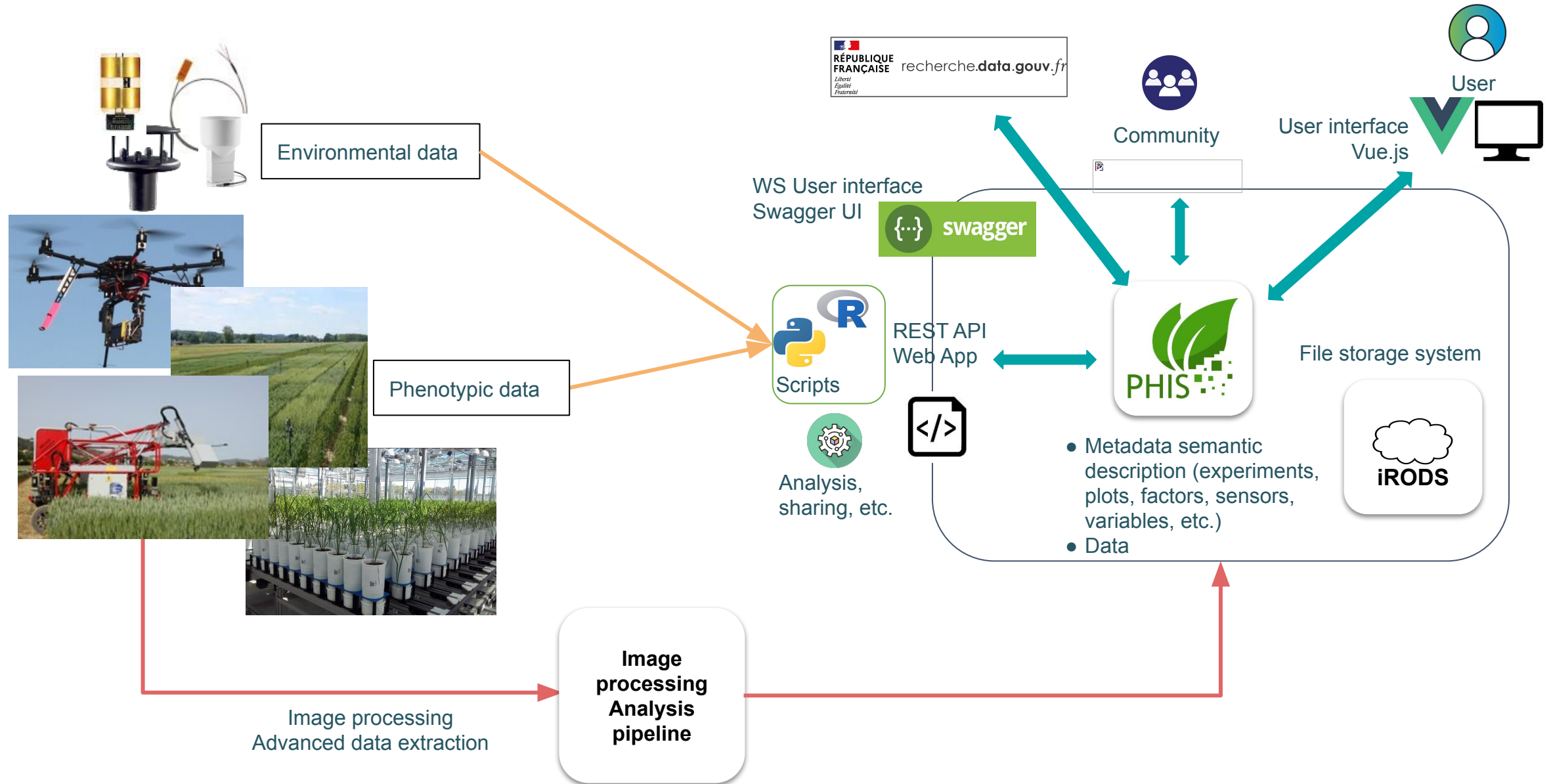
Implementation Example: view images directly from the web interface



The screenshot displays the PHIS web interface. On the left is a sidebar with a navigation menu. The main content area is titled 'Data files' and contains a search and filter panel. The search results show a list of 23 entries, with the first few visible in the background. A modal window titled 'Image' is centered, showing a photograph of a small green plant growing in a pot. The background list shows columns for Target, Date, Type, and Provenance.

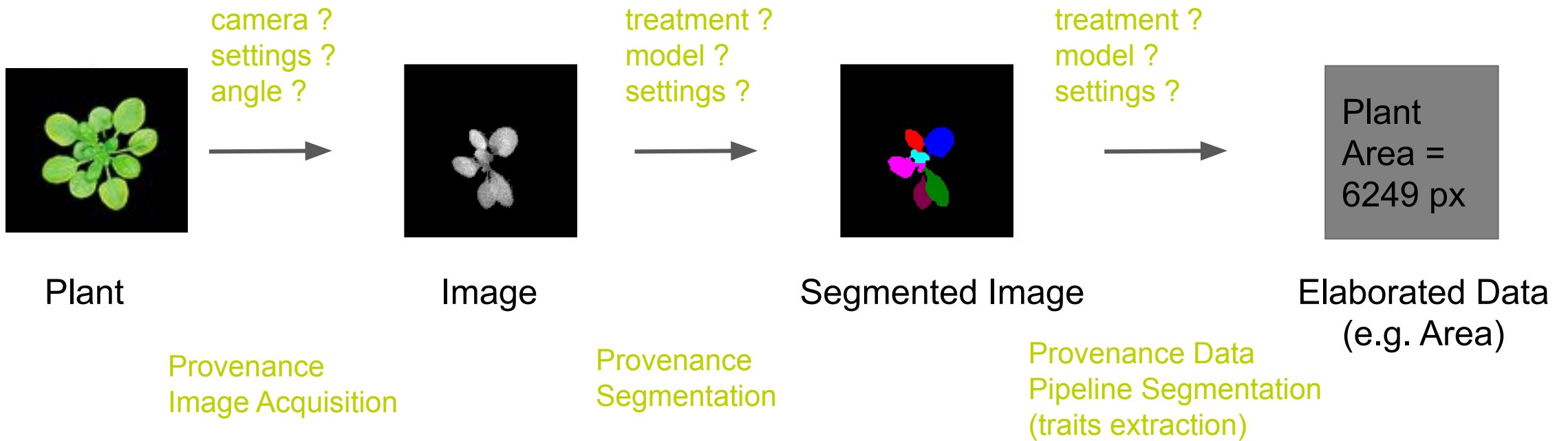
Target	Date	Type	Provenance
cesa 5-2-1 (plant)	2021-06-24T10:03:50.000+0100	image	PHENOBEAN_images_brutes
cesa 5-2-1 (plant)	2021-06-24T18:03:49.000+0100	image	PHENOBEAN_images_brutes
cesa 5-2-1 (plant)	2021-06-23T18:03:46.000+0100	image	PHENOBEAN_images_brutes
cesa 5-2-1 (plant)	2021-06-22T18:03:46.000+0100	image	PHENOBEAN_images_brutes
cesa 5-2-1 (plant)	2021-06-21T18:03:47.000+0100	image	PHENOBEAN_images_brutes
cesa 5-2-1 (plant)	2021-06-20T18:03:48.000+0100	image	PHENOBEAN_images_brutes
cesa 5-2-1 (plant)	2021-06-19T18:03:47.000+0100	image	PHENOBEAN_images_brutes

# Implementation Example: PHIS, an Information System for Plant Phenomics



# Data & Datafiles insertion

## Image processing - Analysis Pipeline





# Use Case - Sylvain Poque

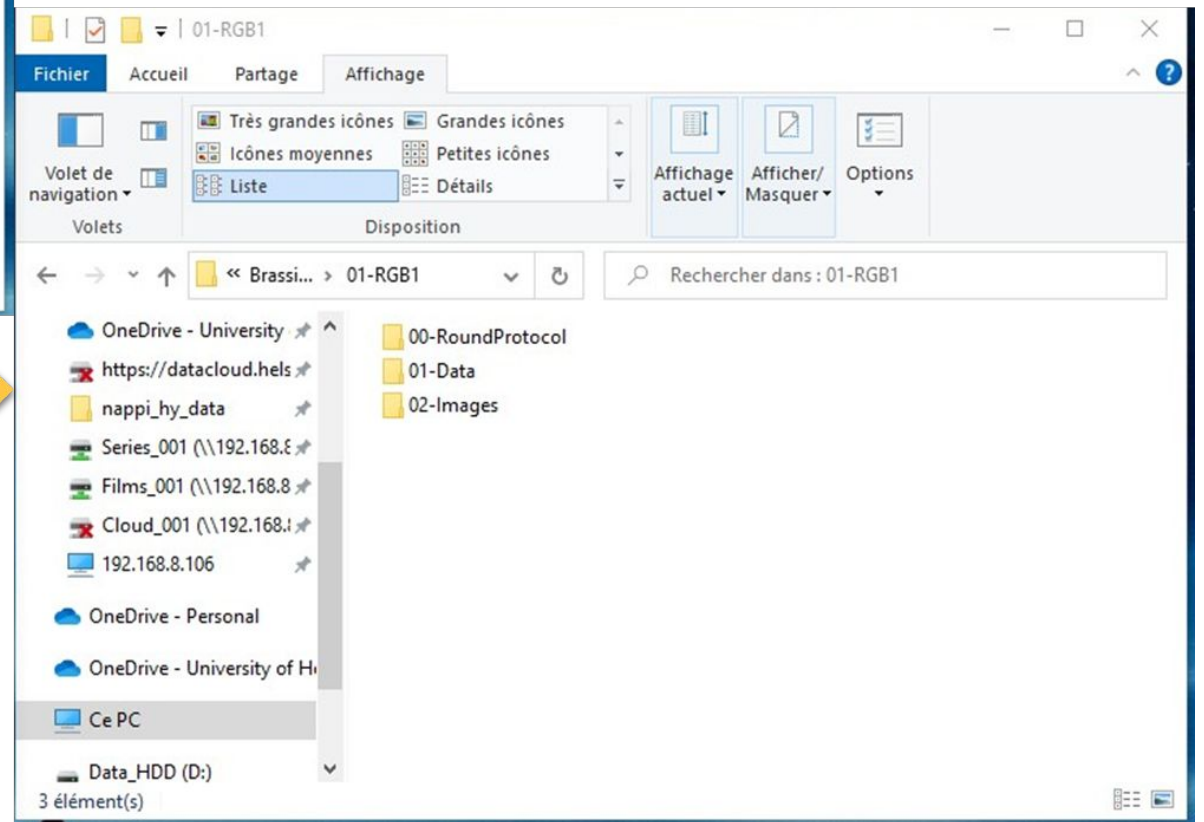
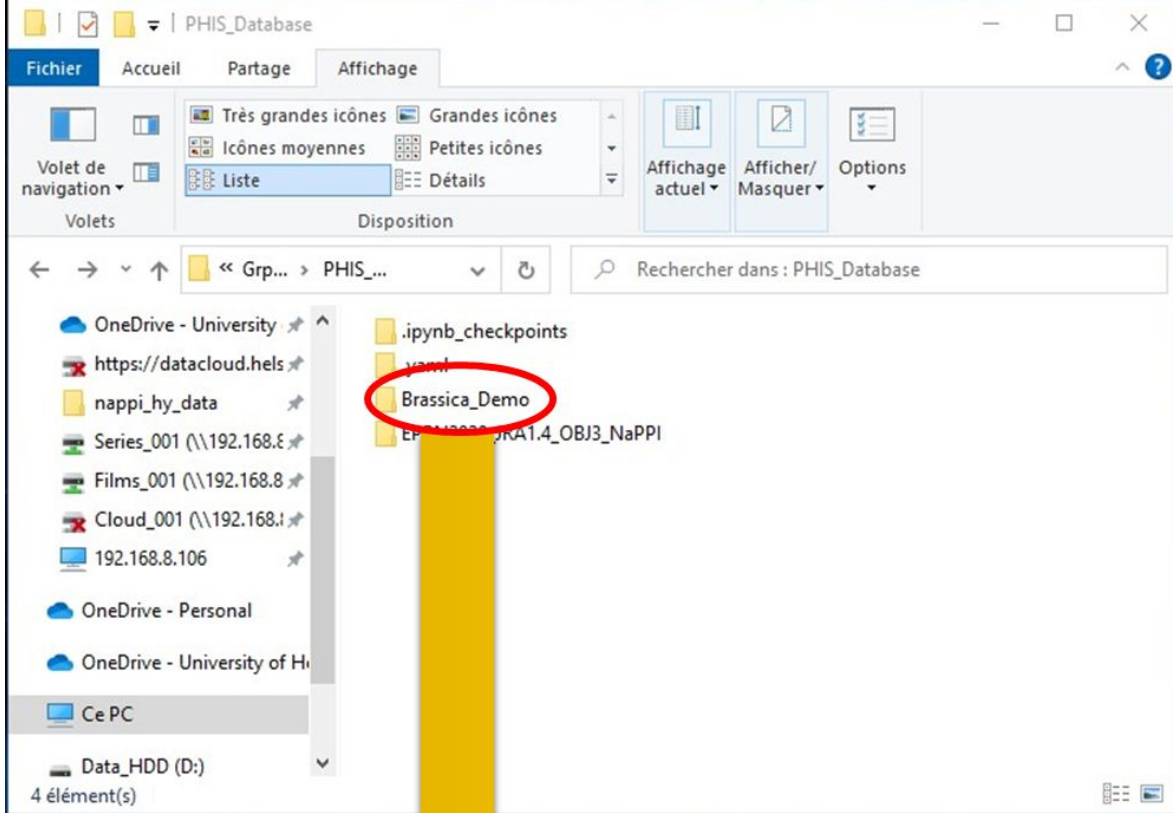
## Sending Data to PHIS Information System



Amazon S3

```

<Measure>
  <AdaptTime>00:00:00</AdaptTime>
  <Prescription id="1" name="Recipe New:1">
    <RGB1 height="Default">
      <Offset>0</Offset>
      <Frame angle="0" />
      <Frame angle="120" />
      <Frame angle="240" />
    </RGB1>
    <RGB2 height="Automatic">
      <Offset>0</Offset>
      <Frame angle="0" />
    </RGB2>
    <Analyse>
      <MaskErosionLevel>1</MaskErosionLevel>
      <RGB1>
        <PlantMask>
          <Formula>0 - ((2*R) + (3*B) - (5*G)) </Formula>
          <Threshold>0.1098039</Threshold>
          <MedianFilterSize>4</MedianFilterSize>
          <MinSize>100</MinSize>
          <MinHoleSize>10</MinHoleSize>
          <CropObjectsOnBorders>False</CropObjectsOnBorders>
          <UseReflectionReduction>False</UseReflectionReduction>
          <SkipBadExposedPoints>True</SkipBadExposedPoints>
        </PlantMask>
      </RGB1>
    </Analyse>
  </Prescription>
</Measure>
  
```



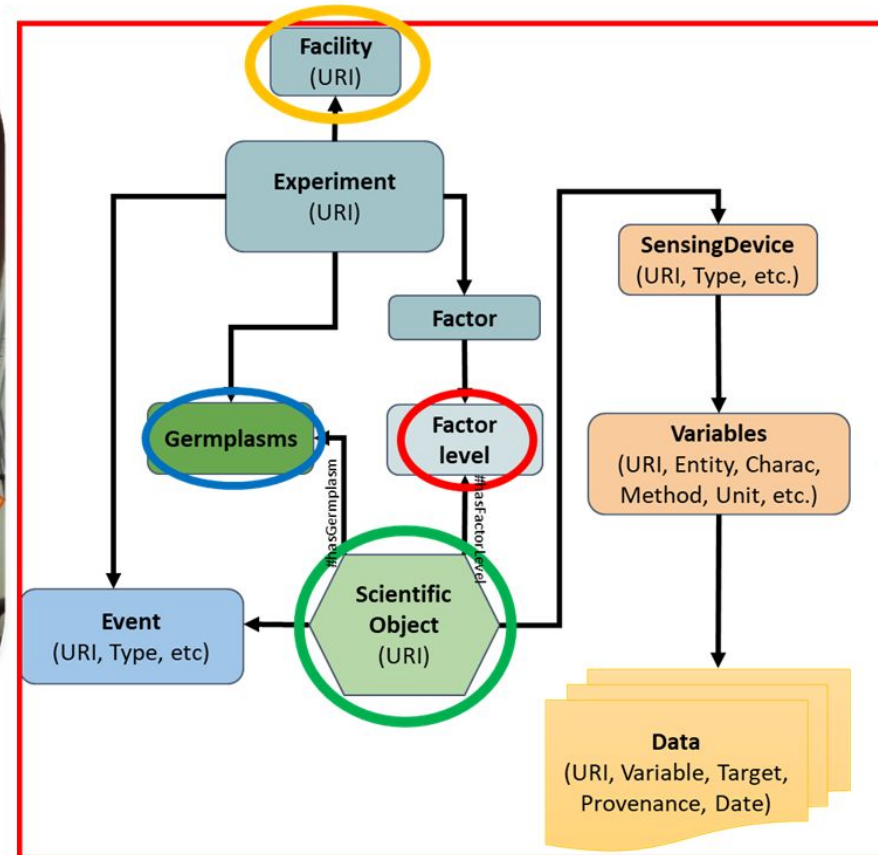
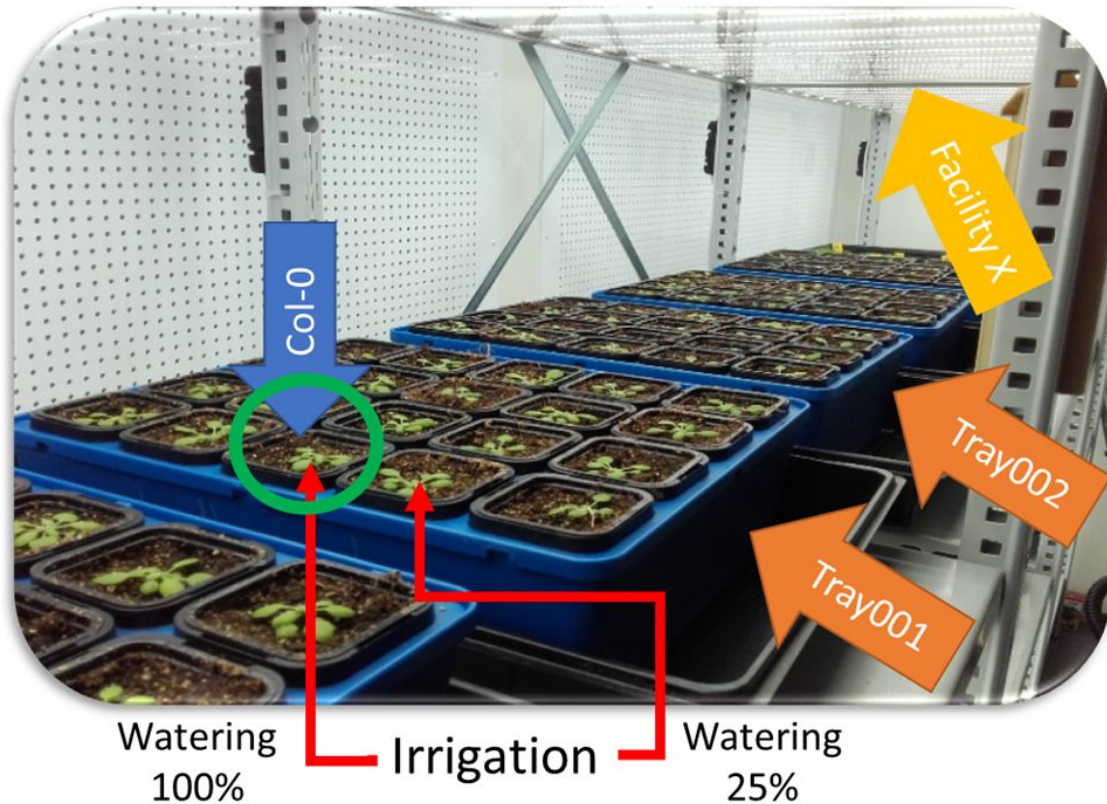
# PHIS Identification & Semantics

## Identification

- Everything can be identified: plants, experiments, sensors, events, etc.
- Persistent, unambiguous, resolvable

## Semantics

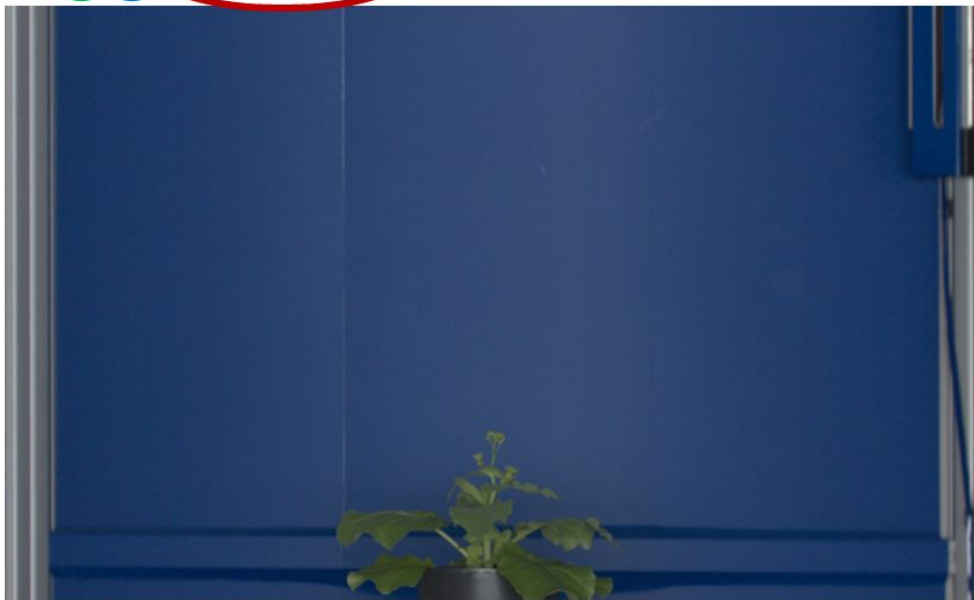
- Naming Conventions
- Controlled vocabulary
- Formalized relationships between entities
- Data annotation and enrichment



# Example of NaPPI: Data structure outputs

69 25 180806.Bra.37-~~RGB1-000-FishEyeCorrected.png~~

69 42 180806.Bra.37-~~RGB1-000-FishEyeCorrected.png~~

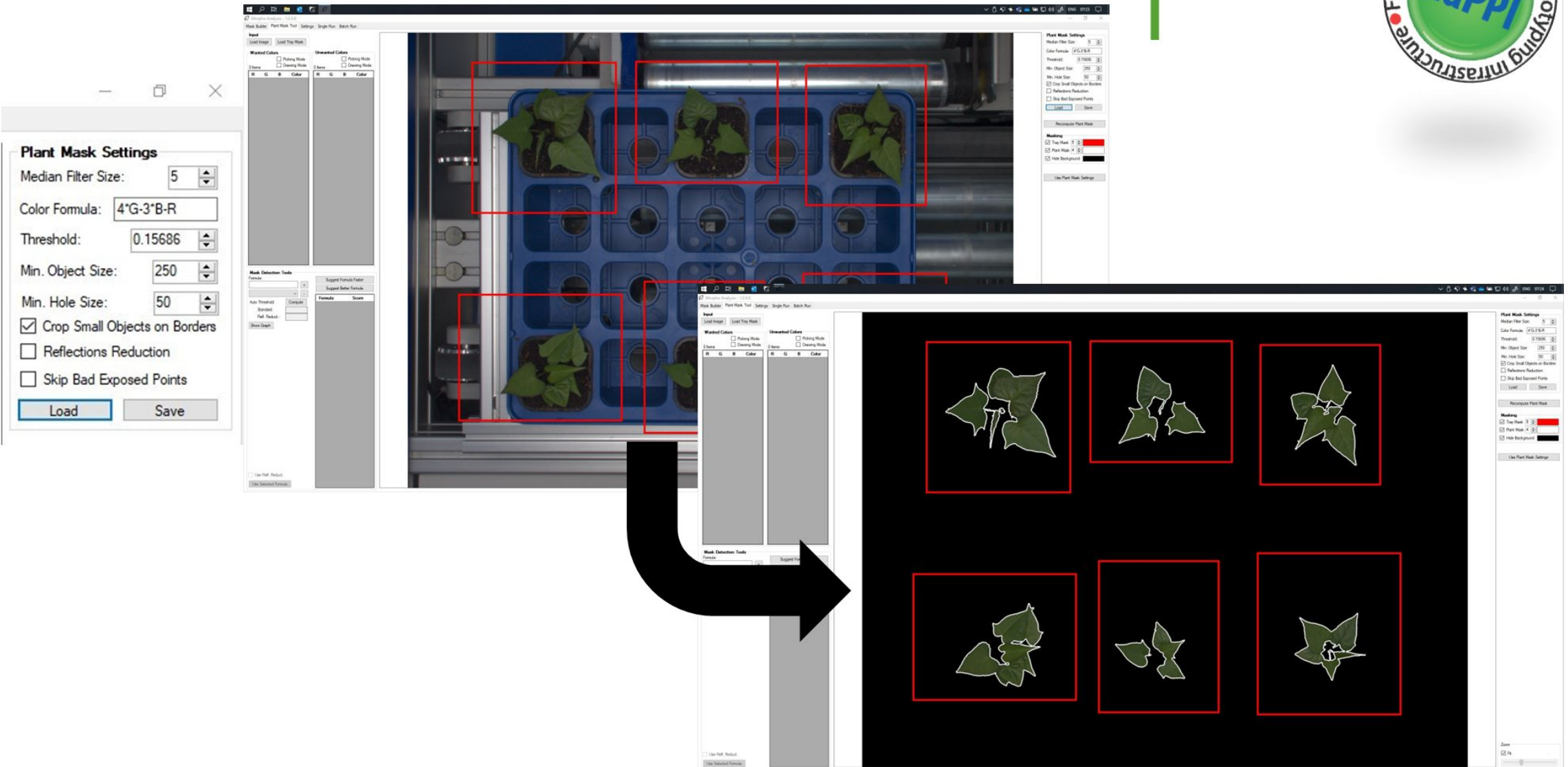


Measuring Date	Experiment ID	Round Order	Tray ID	Plant ID	Geno	Germplasm	PID	Position	Angle	AREA_MM	PERIMETER_MM
2018-08-31	69	14	180806.Bra.37	BRA04_037	SemiDwarf	BC4F2_4003_SemiDwarf	RGB1	A1	0	4990.566734	745.6099917
2018-09-02	69	19	180806.Bra.37	BRA04_037	SemiDwarf	BC4F2_4003_SemiDwarf	RGB1	A1	0	8288.080108	1142.58486
<b>2018-09-06</b>	<b>69</b>	<b>25</b>	<b>180806.Bra.37</b>	<b>BRA04_037</b>	<b>SemiDwarf</b>	<b>BC4F2_4003_SemiDwarf</b>	<b>RGB1</b>	<b>A1</b>	<b>0</b>	<b>26405.13174</b>	<b>2902.298909</b>
2018-09-08	69	34	180806.Bra.37	BRA04_037	SemiDwarf	BC4F2_4003_SemiDwarf	RGB1	A1	0	40337.01069	4089.33801
2018-09-10	69	34	180806.Bra.37	BRA04_037	SemiDwarf	BC4F2_4003_SemiDwarf	RGB1	A1	0	53688.43063	6107.038481
2018-09-12	69	37	180806.Bra.37	BRA04_037	SemiDwarf	BC4F2_4003_SemiDwarf	RGB1	A1	0	64638.33809	7413.620659
2018-09-13	69	39	180806.Bra.37	BRA04_037	SemiDwarf	BC4F2_4003_SemiDwarf	RGB1	A1	0	70297.66953	6896.141705
<b>2018-09-14</b>	<b>69</b>	<b>42</b>	<b>180806.Bra.37</b>	<b>BRA04_037</b>	<b>SemiDwarf</b>	<b>BC4F2_4003_SemiDwarf</b>	<b>RGB1</b>	<b>A1</b>	<b>0</b>	<b>76262.47543</b>	<b>7984.092591</b>
2018-09-17	69	41	180806.Bra.37	BRA04_037	SemiDwarf	BC4F2_4003_SemiDwarf	RGB1	A1	0	97027.26002	9549.016183
2018-09-18	69	41	180806.Bra.37	BRA04_037	SemiDwarf	BC4F2_4003_SemiDwarf	RGB1	A1	0	102117.0628	10932.23031

Scientific Object

Germplasm

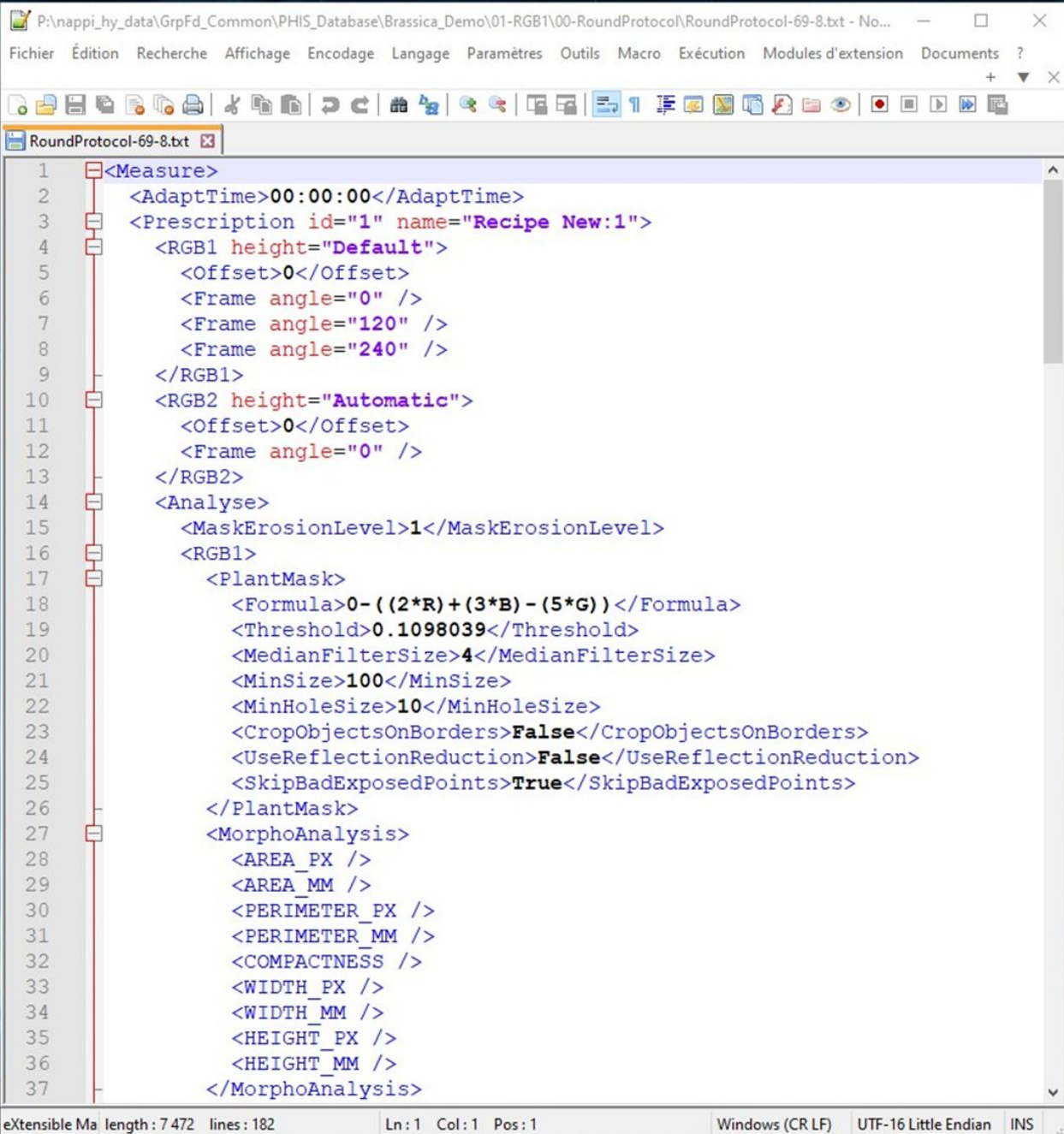
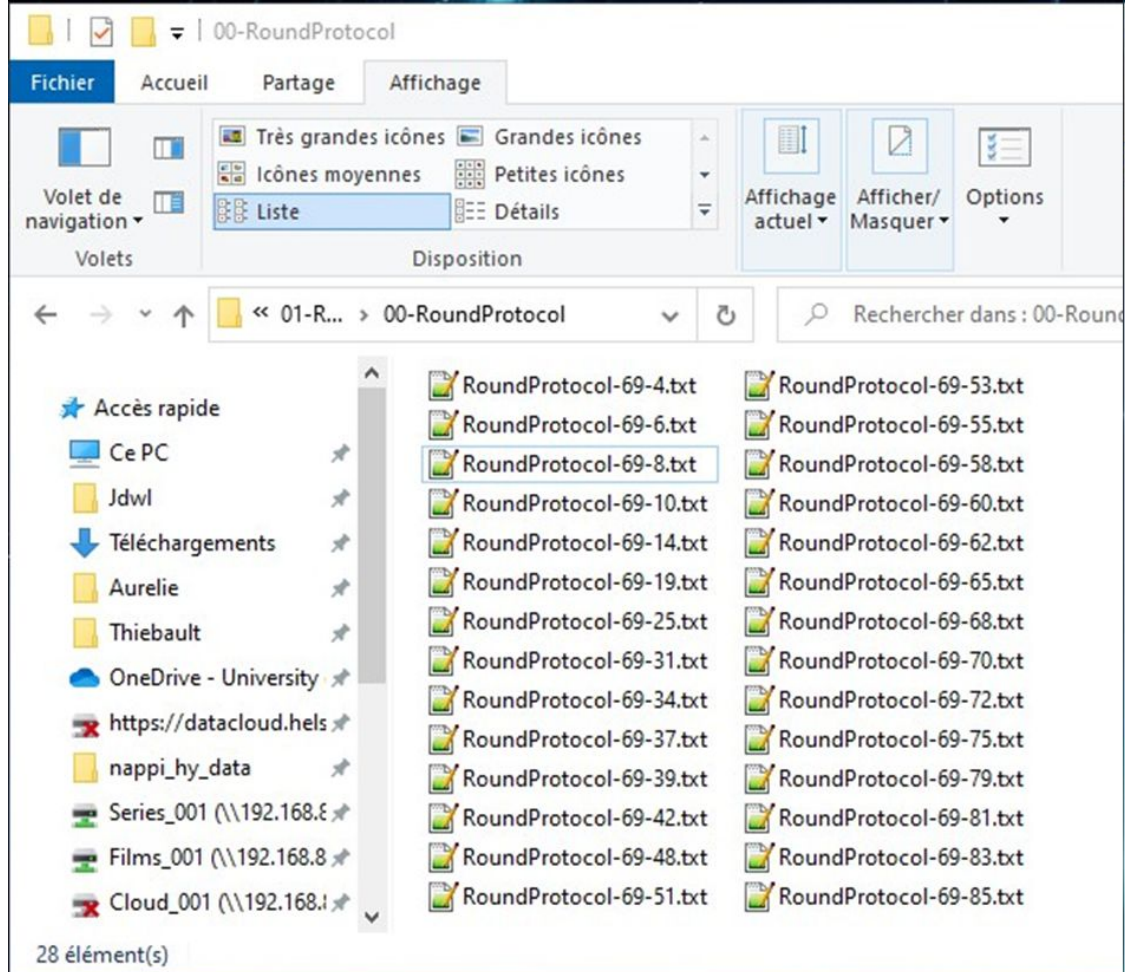
# Extracting info from images with PSI software



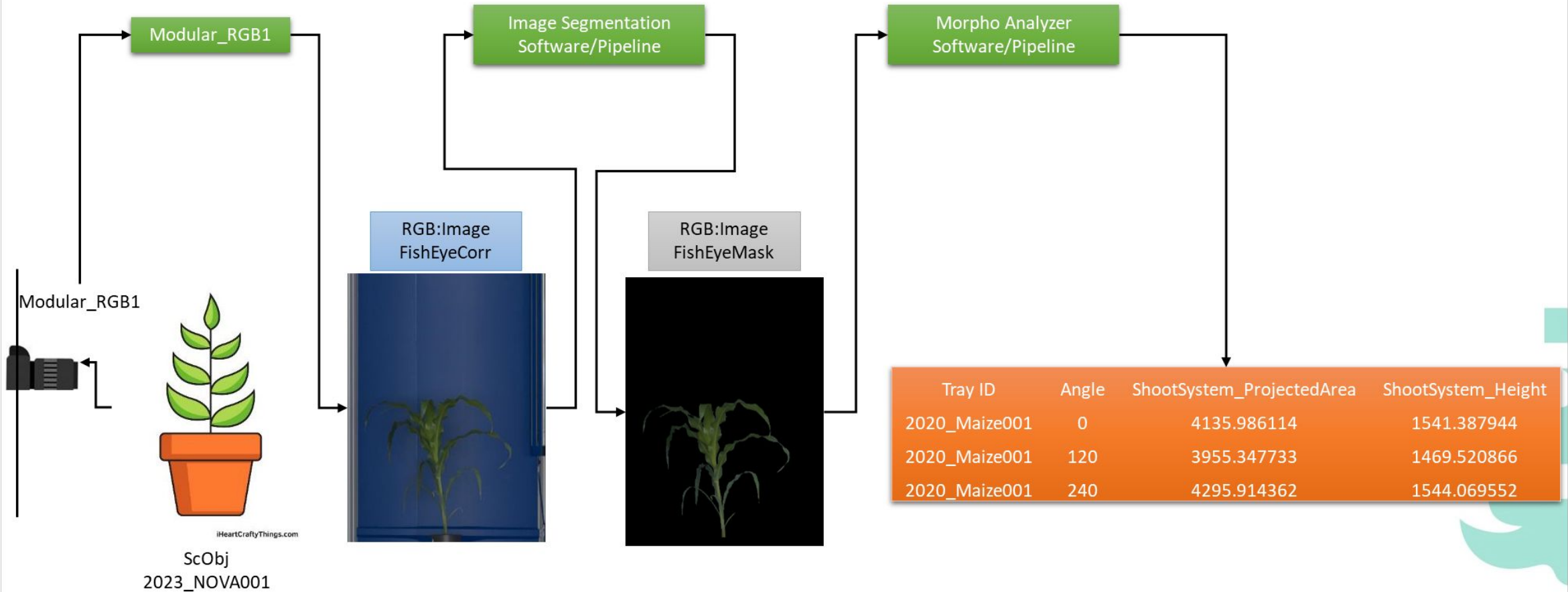
The screenshot illustrates the PSI software interface for plant mask extraction. On the left, the **Plant Mask Settings** panel is visible, containing the following controls:

- Median Filter Size: 5
- Color Formula: 4\*G-3\*B-R
- Threshold: 0.15686
- Min. Object Size: 250
- Min. Hole Size: 50
- Crop Small Objects on Borders
- Reflections Reduction
- Skip Bad Exposed Points
- Buttons: Load, Save

The main window displays a photograph of a blue tray with six plants. Red bounding boxes are drawn around each plant. A large black arrow points from this image to a zoomed-in view of the extracted masks. In this view, the background is black, and each plant is represented by a green mask with a white outline, all contained within red bounding boxes. The software interface also shows various toolbars and settings panels, including 'Mask Detection Tools' and 'Plant Mask Settings' on the right side.



# PHIS, Provenance



# Conclusion

## Take home message

- **Variables:** Take time to define your variables will allow you to understand them, reuse them, share them  
=> Portals exist to help you discovering Ontologies and Concepts! (Agroportal, BioPortal, Ontobee, etc.)
- **Datafiles:** study the storage solution best suited to **your needs and uses**  
*Remember you can have separate solutions for your information system!*
- **Provenance:** Make your data really **Reusable!**
- **Data Transfer Pipeline:** for improved automatization!



# Thank you for your attention!



<https://www.phenome-emphasis.fr/>



OpenSILEX Team - <http://opensilex.org/>



<http://www.phis.inrae.fr/>

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 EMPHASIS on Plant Phenomics



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EUROPEAN INFRASTRUCTURE  
FOR PLANT PHENOTYPING