

Tools and methods for the evaluation of Plant Genetic Resources

Entry-level training school on Plant Genetic Resources (PGR)

5-6 October, Mediterranean Agronomic Institute of Chania, Chania, Greece

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SPAIN

Content

- The need for evaluation of plant genetic resources
- Characterization and evaluation
- Agro-morphological characterization: the use of descriptors
- Evaluation for resistance to pests and diseases
- Evaluation for adaptation to abiotic stresses
- Speeding evaluation by high throughput phenotyping techniques

Characterization and evaluation is the key to assess the potential and actual value of germplasm

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Breeding for adaptation to climate change



Disease resistance



Cytoplasmic male sterile lines for hybrid breeding

Plant Genetic Resources



Insect-pest resistance



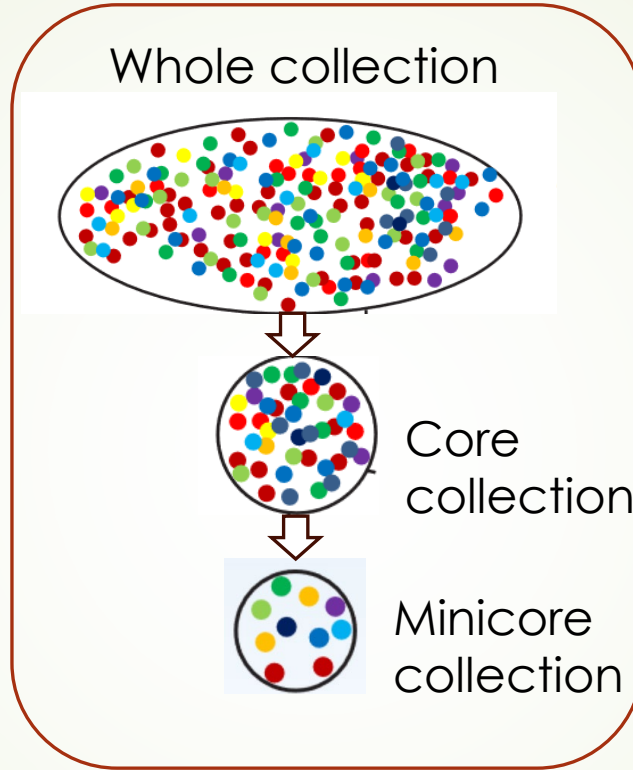
Breeding for non-conventional seasons



productivity enhancement

Germplasm characterization and evaluation in the broad sense and in the context of genetic resources is the description of a particular accession

Primary characterization by universally accepted descriptors



Categorization of the collection, core and minicore collections, reference sets of germplasm

EVALUATION

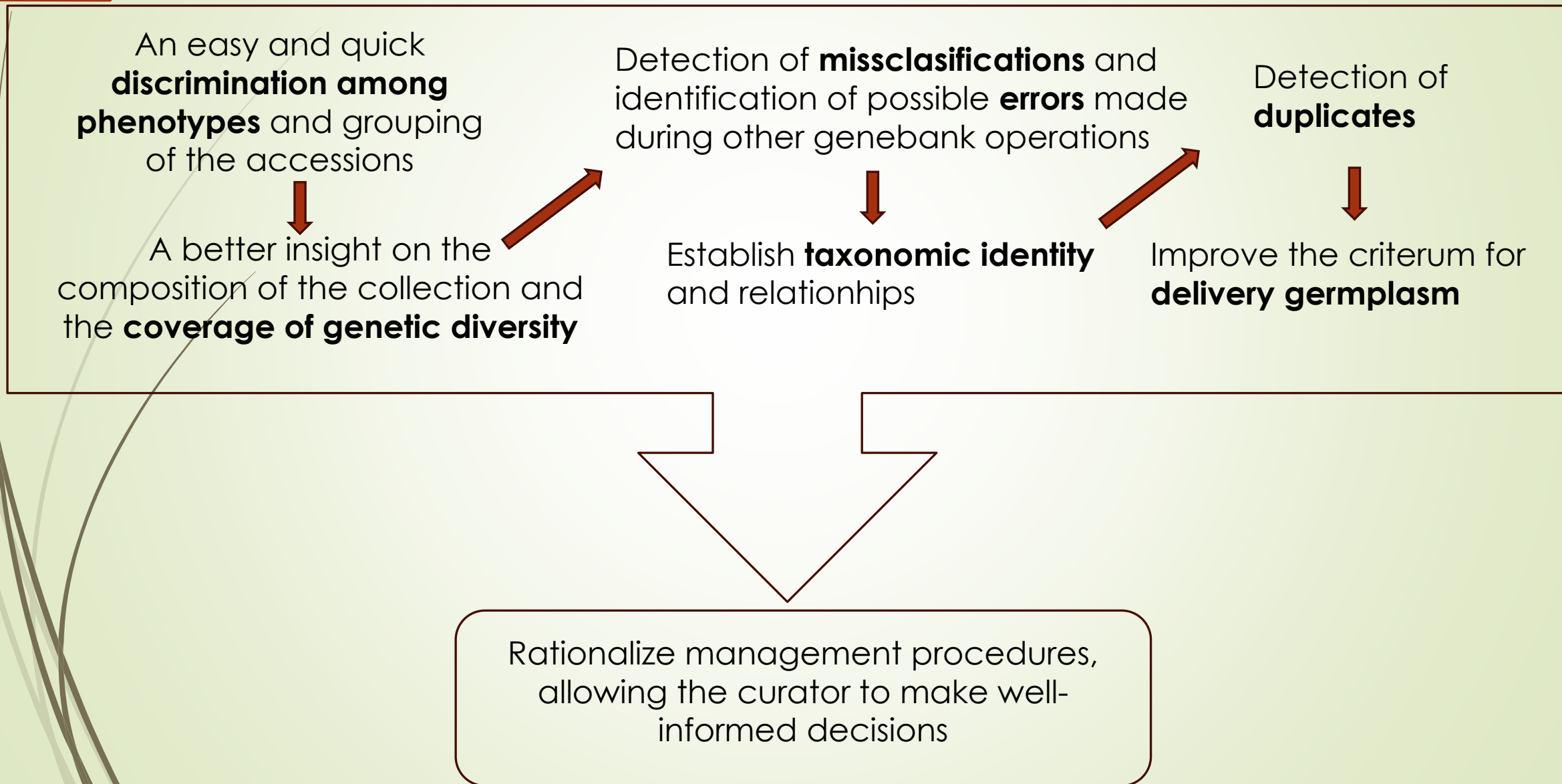
Agronomic and quality

Resistance to pests and diseases

Adaptation to abiotic stresses

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Characterization is necessary to:



Characterization and evaluation

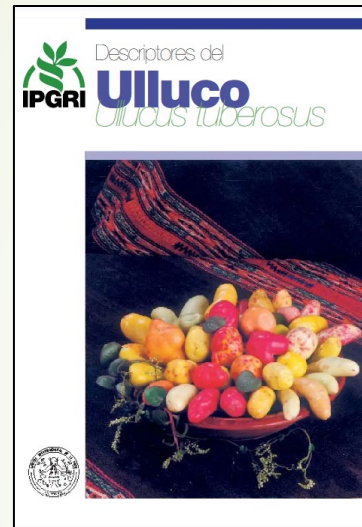
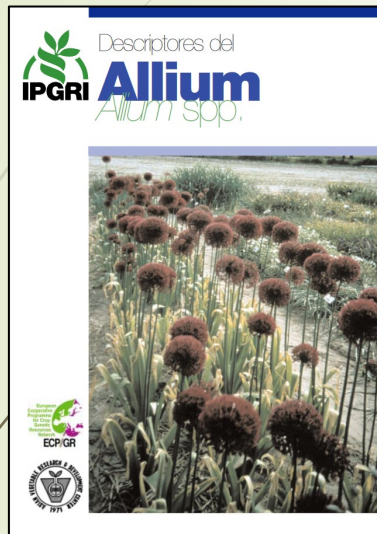
- The **characterization** of germplasm deals with the understanding and recording of **highly heritable traits** which are generally expressed in all the environments. Therefore, it can be performed in a single environment. It ranges from morphological features to seed proteins and molecular and biochemical markers
- Germplasm **evaluation** deals with the assessment of the agronomic potential of an accession including quality parameters and response to various abiotic and biotic stresses



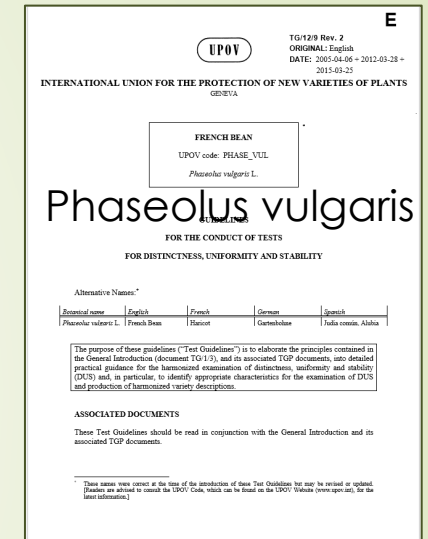
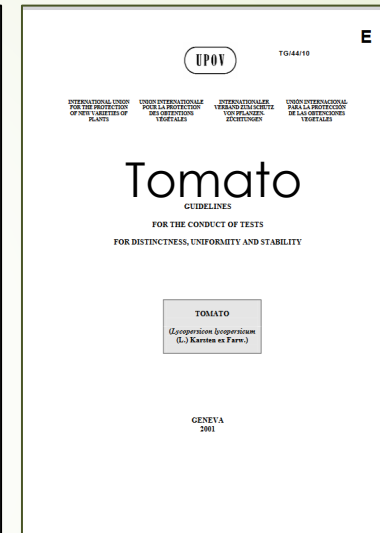
Morphological characterization

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IPGRI Descriptors



UPOV Guidelines




- Passport descriptors
- Management descriptors
- Environment and site descriptors
- Characterization descriptors
- Evaluation descriptors

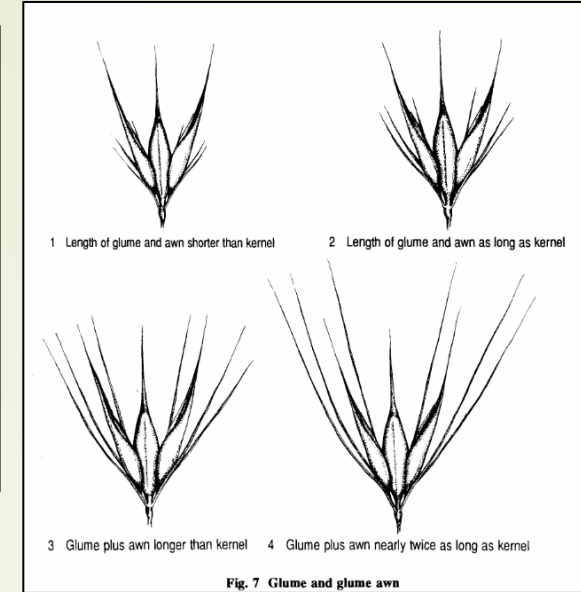
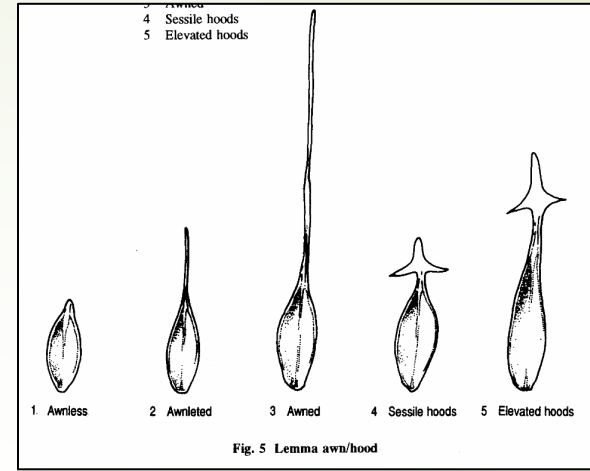
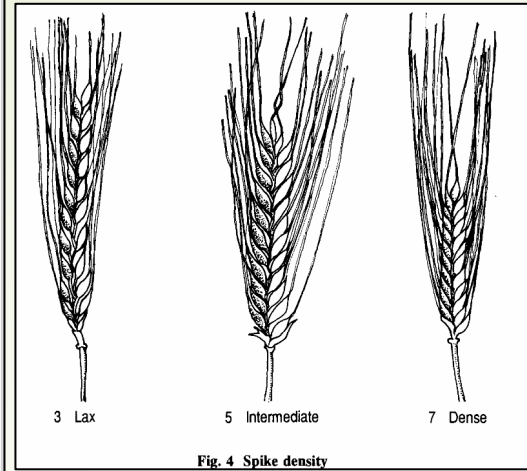
DEFINITIONS AND USE OF THE DESCRIPTORS

- **Passport descriptors:** These provide the basic information used for the general management of the accession and describe parameters that should be observed when the accession is originally collected.
Accession number
Donor name
...
- **Management descriptors:** These provide the basis for the management of accessions in the genebank and assist with their multiplication and regeneration.
Storage address
Storage date
...
- **Environment and site descriptors:** These describe the environmental and site-specific parameters that are important when characterization and evaluation trials are held.
Latitude, Longitude
Elevation
...
- **Characterization descriptors:** These enable an easy and quick discrimination between phenotypes. They are generally highly heritable, can be easily seen by the eye and are equally expressed in all environments.
Plant growth type
Foliage density
...
- **Evaluation descriptors:** Many of the descriptors in this category are susceptible to environmental differences but are generally useful in crop improvement and others may involve complex biochemical or molecular characterization. They include yield, agronomic performance, stress susceptibilities and biochemical and cytological traits
Yield
Agronomic performance
Biotic stresses
Abiotic stresses
...

Descriptors for
Barley
(*Hordeum vulgare* L.)



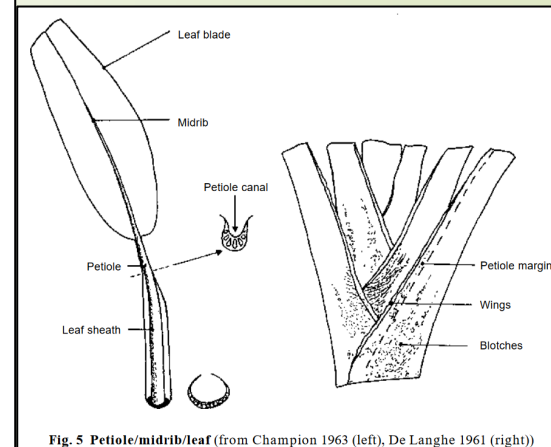
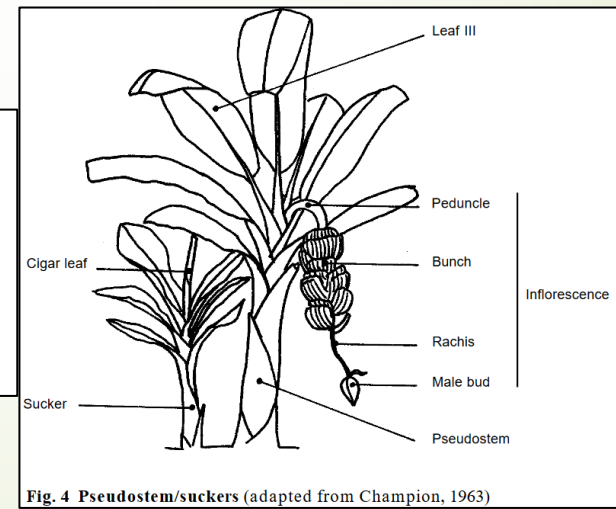
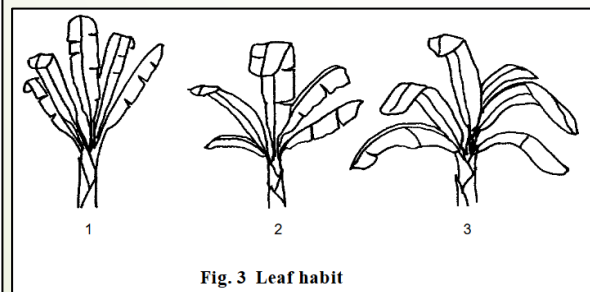
IPGRI



Descriptors for
Banana
(*Musa* spp.)



CIRAD
inbap
IPGRI



Descriptors for <i>Allium</i> spp.		
Passport		Number descriptors
	Accessions	12
	Collecting	17
Management		10
Multiplication/regeneration		11
Environment and site		11
Characterization		
	Plant descriptors	
	Vegetative	28
	Inflorescence and fruit	10
	Seeds	2
Evaluation		
	Plant descriptors	
	Vegetative	9
	Inflorescence and fruit	3
	Abiotic stresses	5
	Biotic stresses	6
	Biochemical markers	Isozymes
	Molecular markers	5

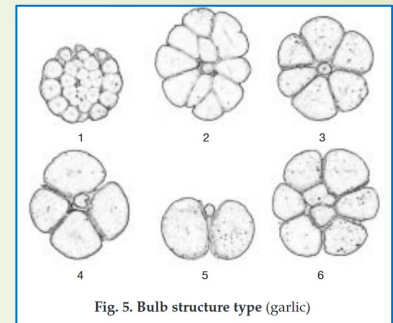
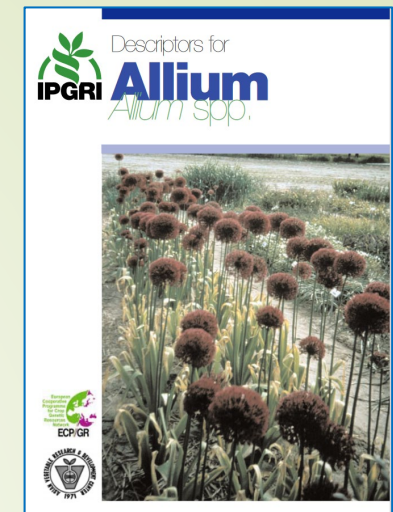


Fig. 5. Bulb structure type (garlic)

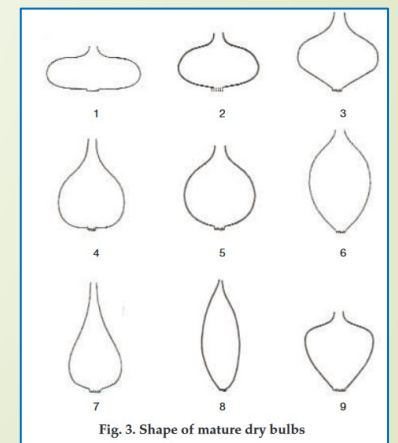


Fig. 3. Shape of mature dry bulbs

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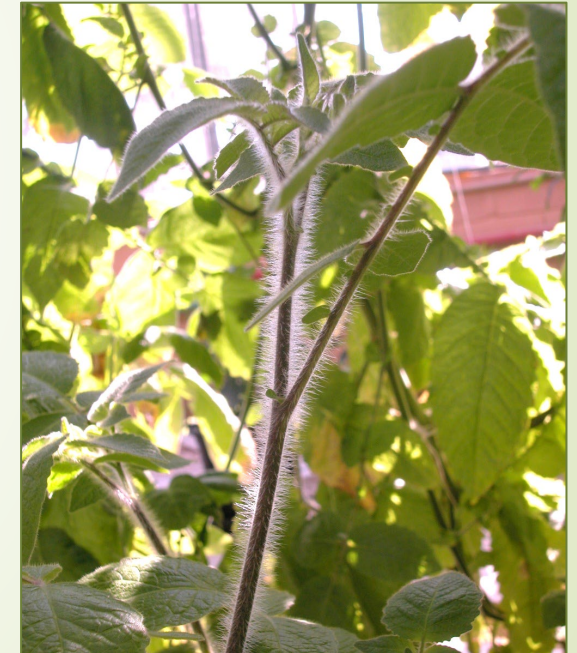
Biological status of the germplasm should be known in advance to determine the characterisation strategy



Heterogeneous traditional varieties maintained by farmers



Uniform breeding lines and improved materials



Specific traits: wild species

The basic morphological characterization: plant traits



Indeterminate



Semideterminate



Determinate



Present



Absent

- Seedling: anthocyanin coloration of Hypocotyl
- Plant: growth type
- Plant: number of inflorescences on main stem
- Inflorescence: number of flowers per inflorescence
- Stem: anthocyanin coloration
- Stem: length of internode
- Plant: height

The basic morphological characterization: leaf traits



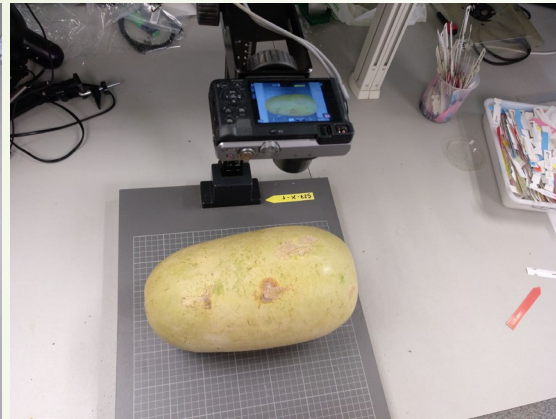
- Leaf: attitude
- Leaf: length
- Leaf: width
- Leaf: type of blade
- Leaf: size of leaflets
- Leaf: intensity of green colour
- Leaf: glossiness
- Leaf: blistering
- Leaf: attitude of petiole of leaflet



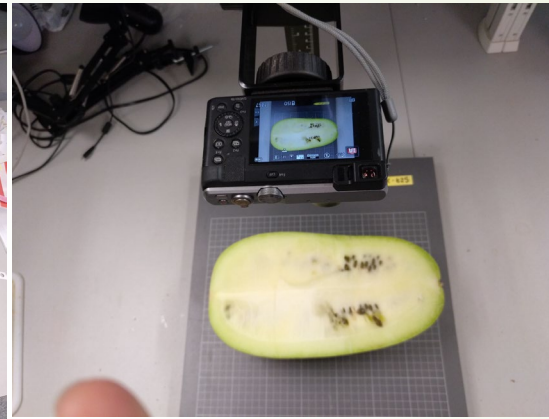
The basic morphological characterization: fruit



Fruit weight



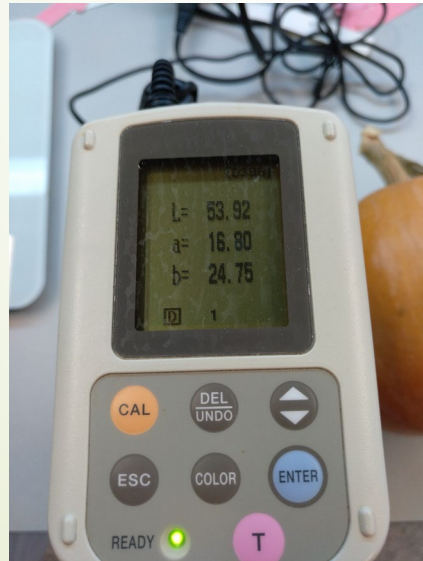
Images



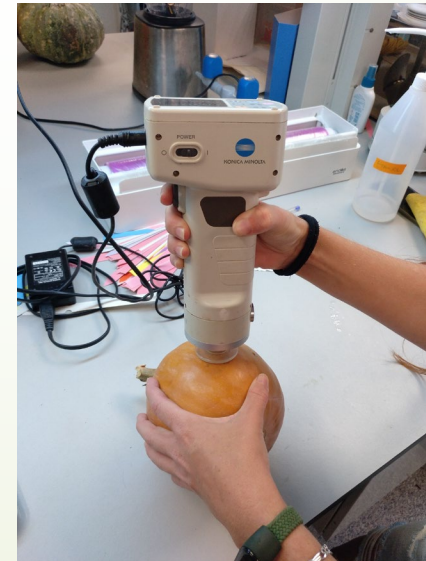
Fruit length and with



^a Brix



Colorimeter



The need for standardization

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- Standardization for:
 - Traits: IPGRI descriptors, UPOV, etc.
 - MIAPPE (**Minimum Information About Plant Phenotyping Experiments**): MIAPPE is an open, community driven, data standard designed to **harmonize data from plant phenotyping experiments**. MIAPPE provides a specification including a checklist and a data model of metadata required to adequately describe plant phenotyping experiments.
 - Ontologies: The Plant Ontology is a **structured vocabulary and database resource that links plant anatomy, morphology and growth and development to plant genomics data**. The PO is under active development to expand to encompass terms and annotations from all plants.

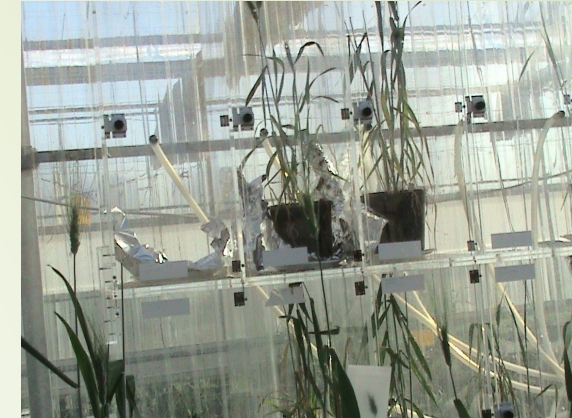
Evaluation is needed for

- **Identify the appropriate germplasm with a target trait for their further utilization.**
- There is a need for a systematic evaluation in order to know its various morphological, physiological and developmental characters
- Evaluation of germplasm is a **multi-disciplinary approach** and it should be done in collaborative mode involving germplasm **curator, plant breeder, physiologist, pathologist, entomologist, biochemist** etc.
- The germplasm accessions are usually evaluated for **two consecutive years** for an adequate documentation
- For effective evaluation of germplasm, a **close organization and personal contact between curator and breeder** is necessary: It servers **to meet the demands of companies**

Evaluation for resistance to diseases.

Previous concepts:

- The response of a plant depends on:
 - The host
 - The pathogen race
 - The environment



Source: F. Martínez

- The identification of a resistance source against a particular race/strain/isolate/biotype within a particular location does not guarantee its resistance response in other locations as **race/strain/isolate/biotype** may vary depending upon the **agro-meteorological conditions** and the presence of **resistance genes** in the commercial varieties

Evaluation for resistance to diseases.

Aspects to take into account

- ▶ Natural infection conditions vs. controlled conditions: mimic the conditions of development of the disease and the infection mode of the pathogen
- ▶ Wild vs. cultivated species: the presence of natural mechanisms of avoidance or antixenosis
- ▶ Definition of the inoculation technique
- ▶ Standard Evaluation Systems (SES): Development of a reliable diagnostic technique
 - ▶ Evaluation of symptoms
 - ▶ Quantification of pathogen

Evaluation for resistance to diseases: The tomato yellow leaf curl virus.

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Controlled conditions with the
vector (*Bemisia tabaci*)



Agroinoculation

Inoculation techniques

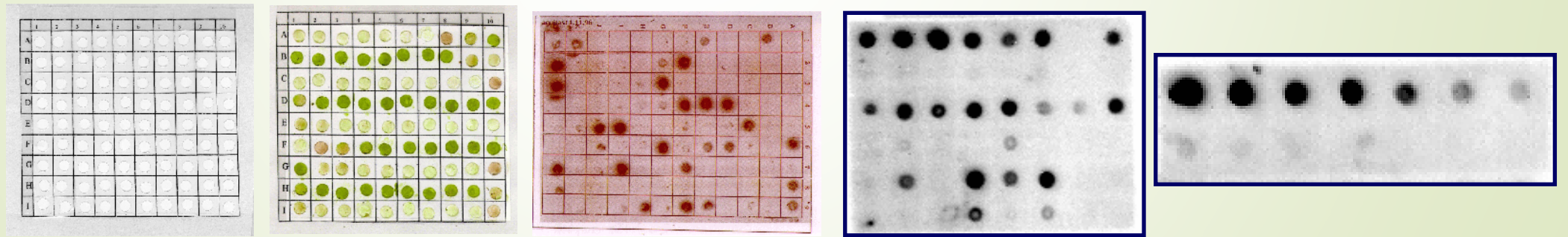


Natural infection conditions

Standard Evaluation System: severity of symptoms at 15, 25, 35, 45 and 55 after inoculation



Virus quantification by squash blot or dot blot



Squash blot

Dot blot

The problem of wild relatives for evaluation of resistance

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- Antixenosis or antibiotic mechanisms: presence of dense trichomes



Solanum pennellii

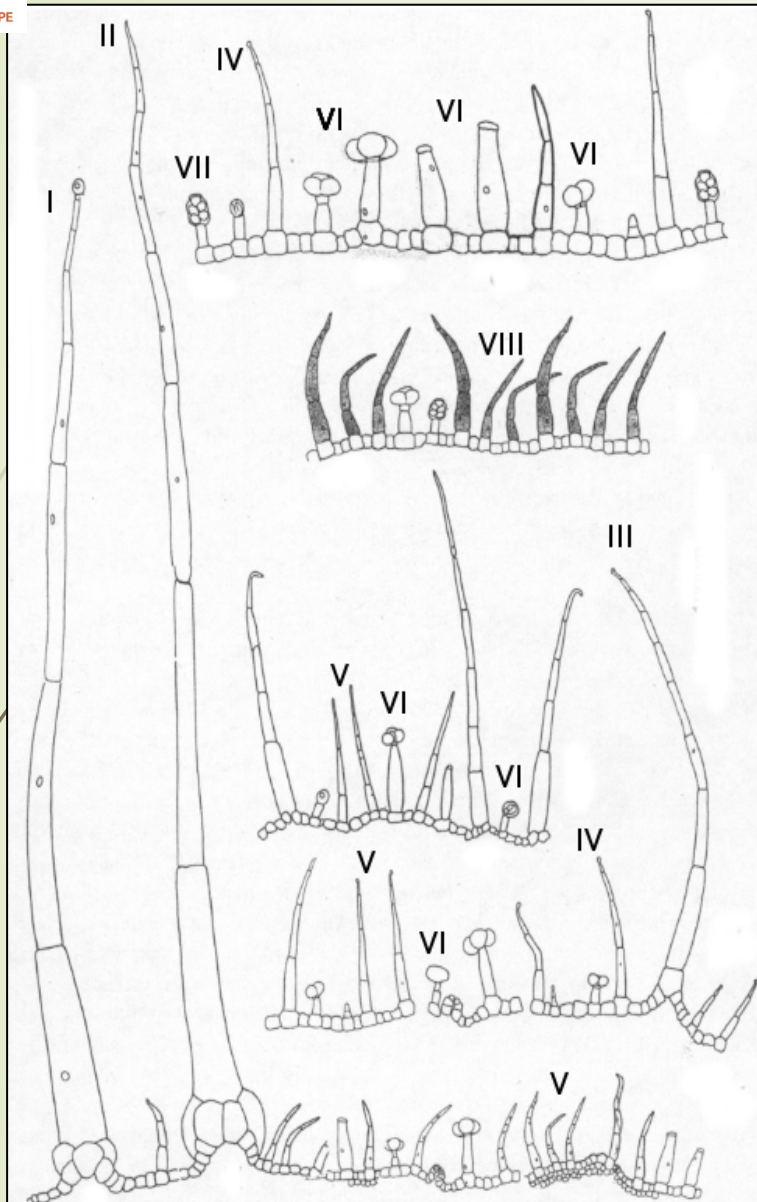


Solanum habrochaites

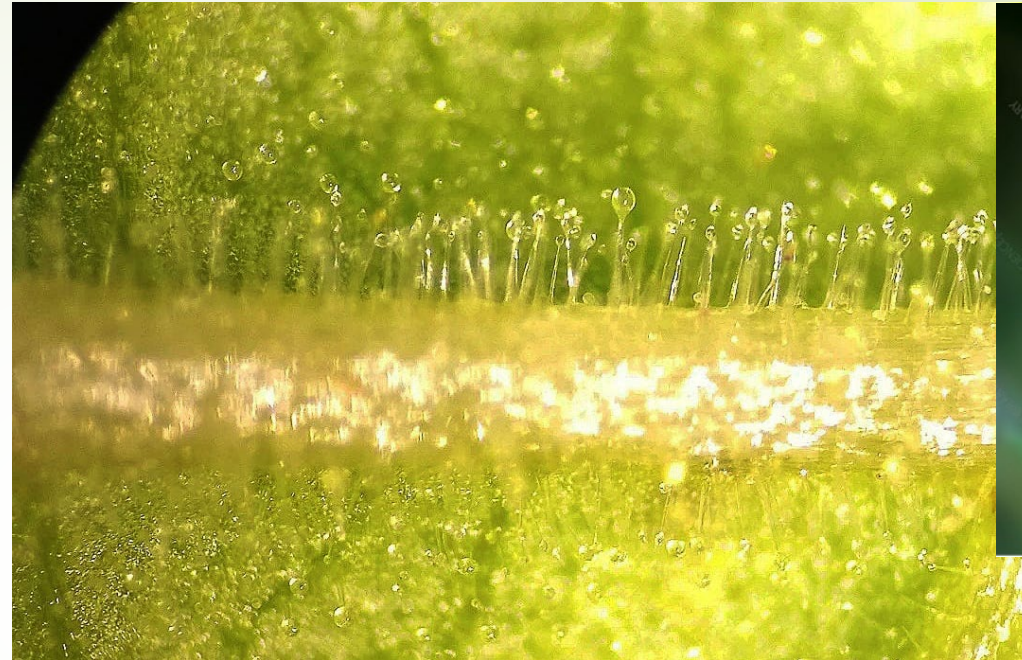
- Symptomatic and asymptomatic infected plants



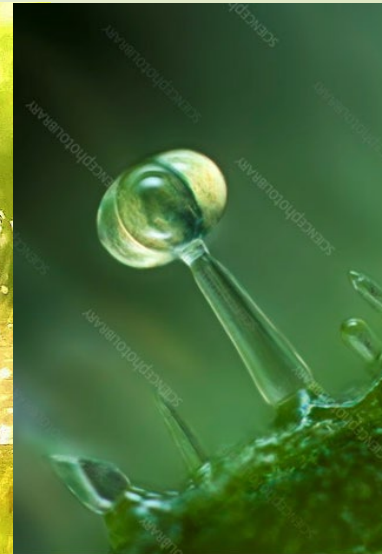
Solanum peruvianum

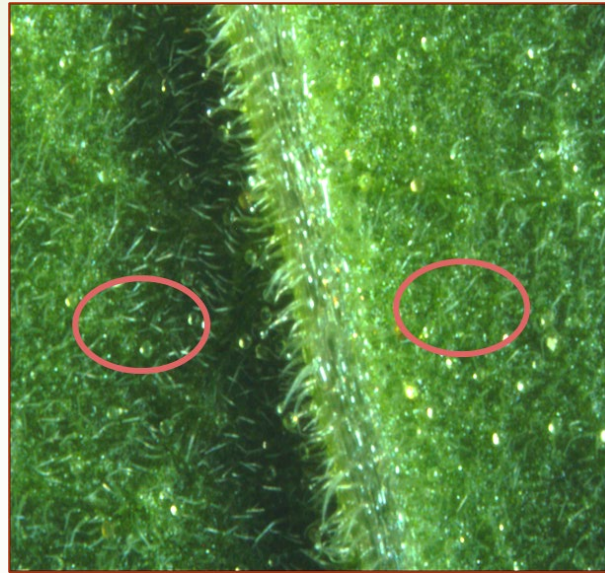
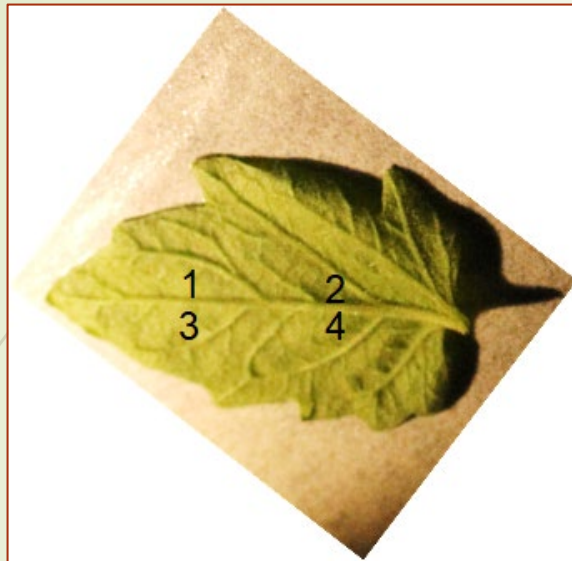


Types of trichomes in tomato and wild relatives (Modified Peralta, Spooner & Knapp, 2008)



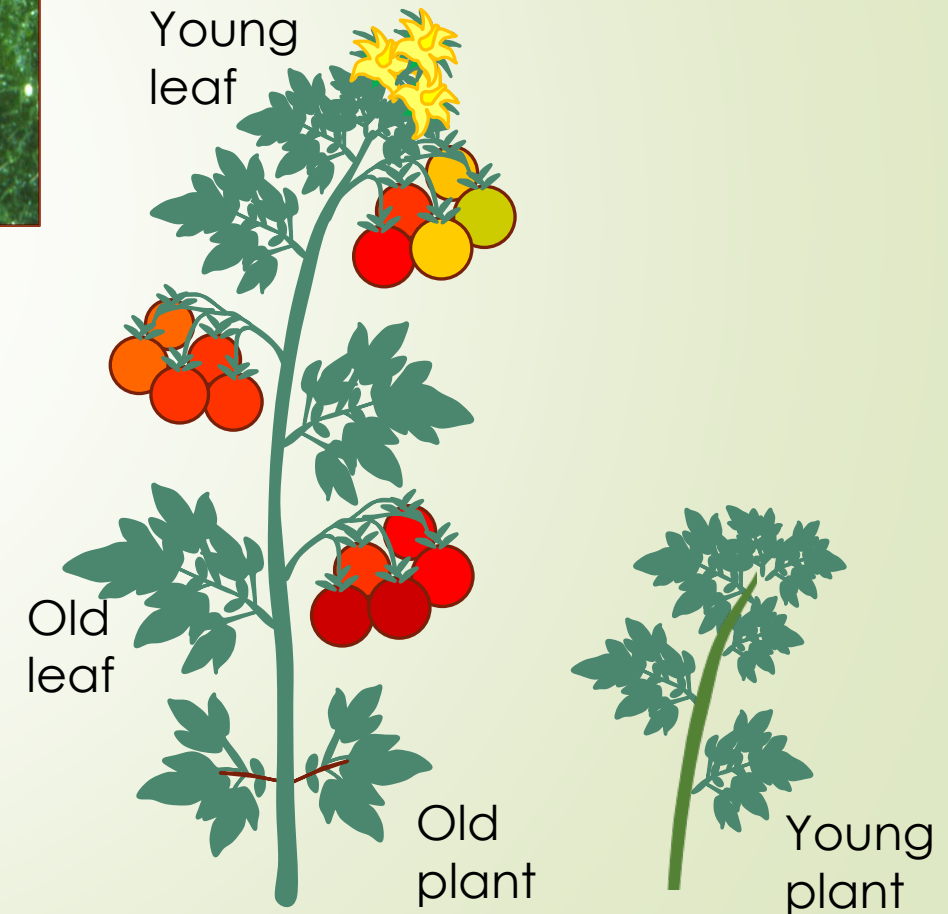
Glandular trichomes





► Factors affecting the development of trichomes:

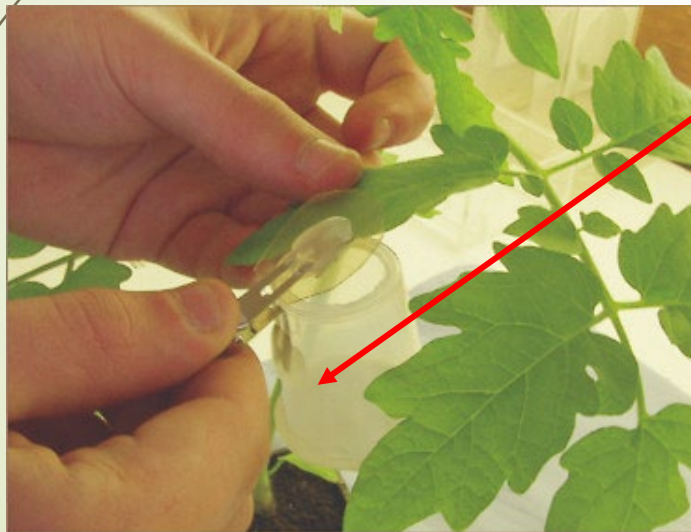
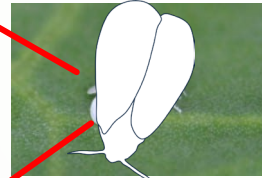
- Planta age
- Leaf age
- Temperature
- Humidity



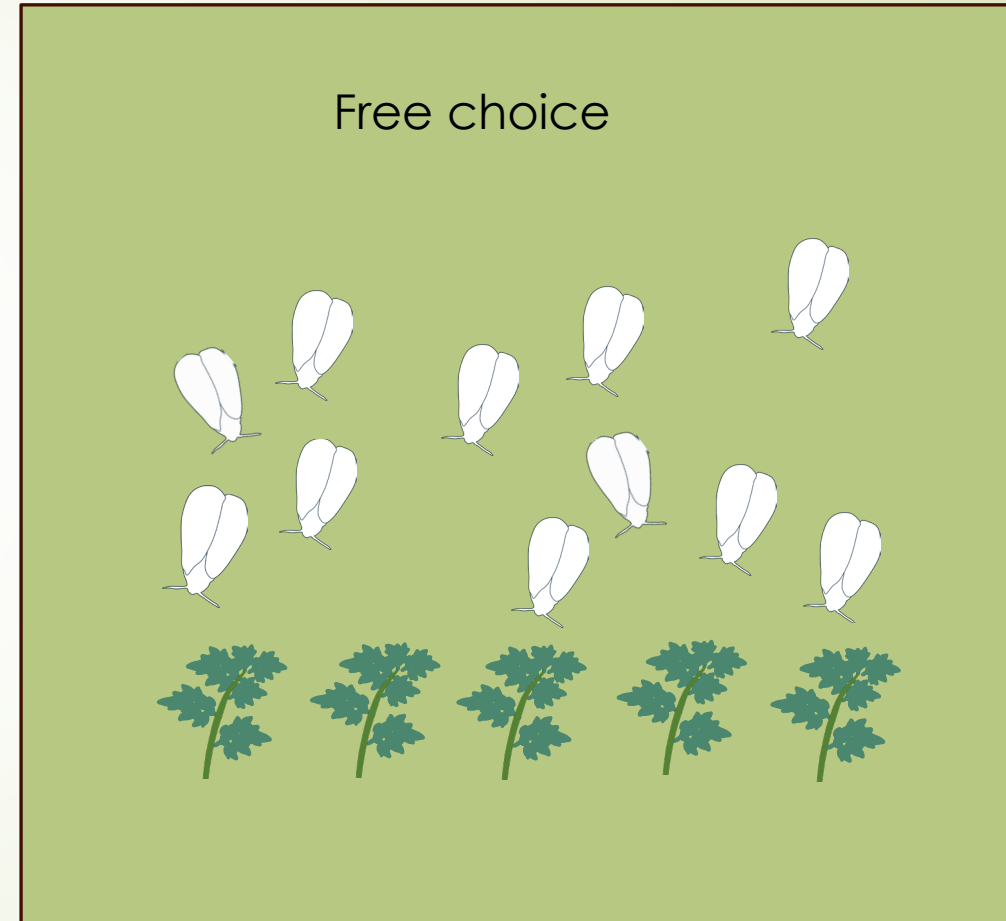
Looking deeper: - the resistance mechanism: antibiosis or antixenosis



Controlled infestation



Antibiosis mechanisms: mortality, fecundity

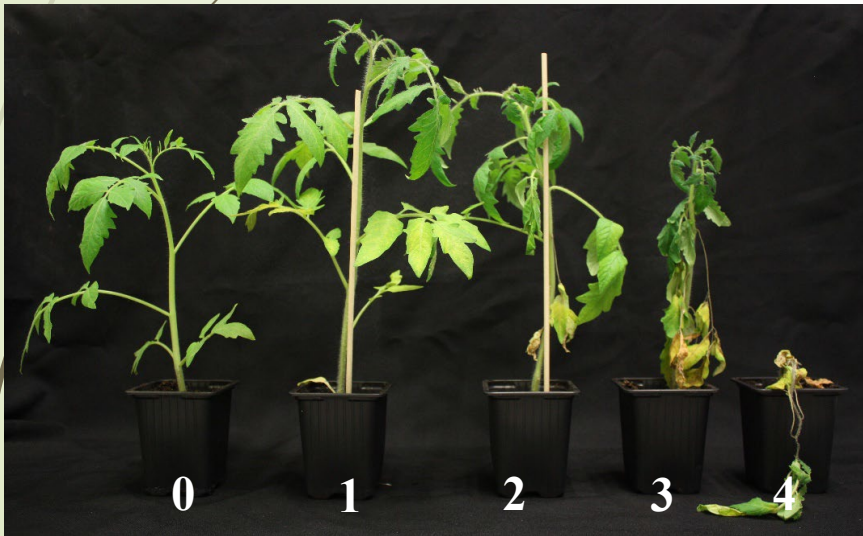
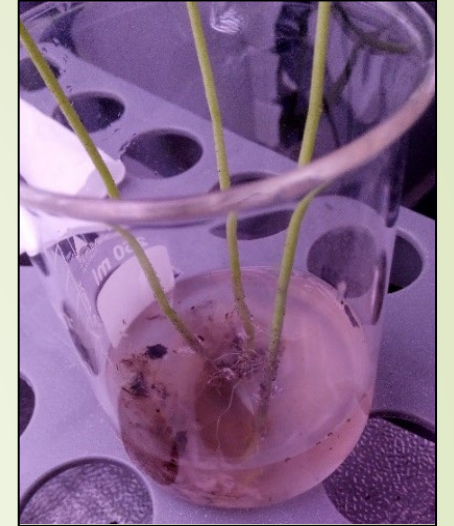


Free choice

Antixenosis mechanisms: epidemiology

Screening for *Fusarium oxysporum* f. sp. *lycopersici*

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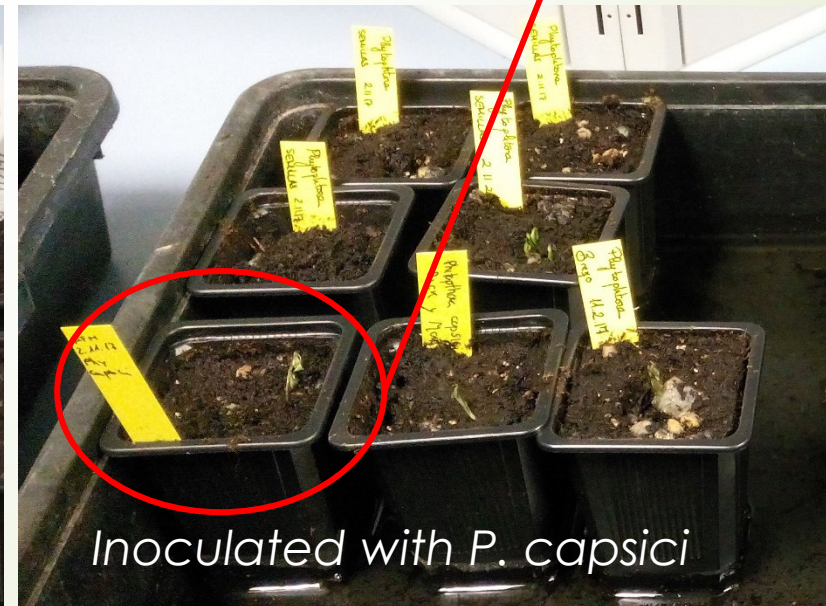
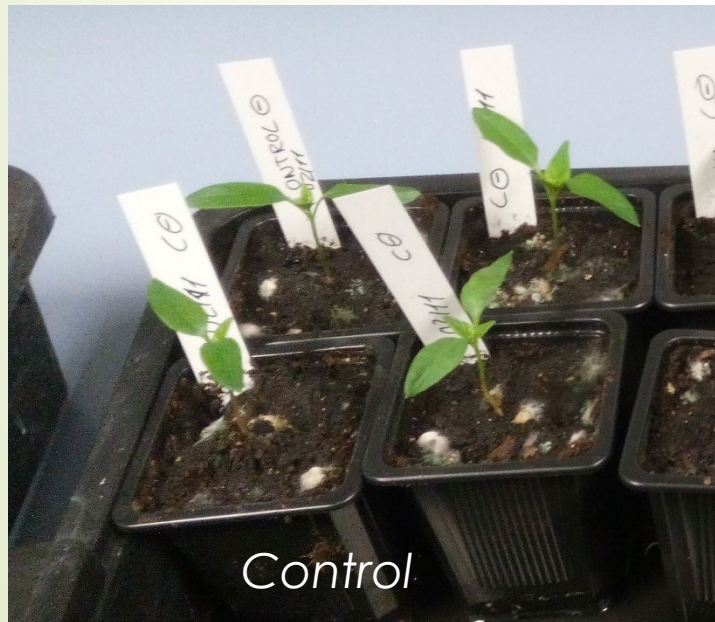
Symptoms scoring



Necrosis in a susceptible plant



Screening for resistance to *Phytophthora capsici* in pepper

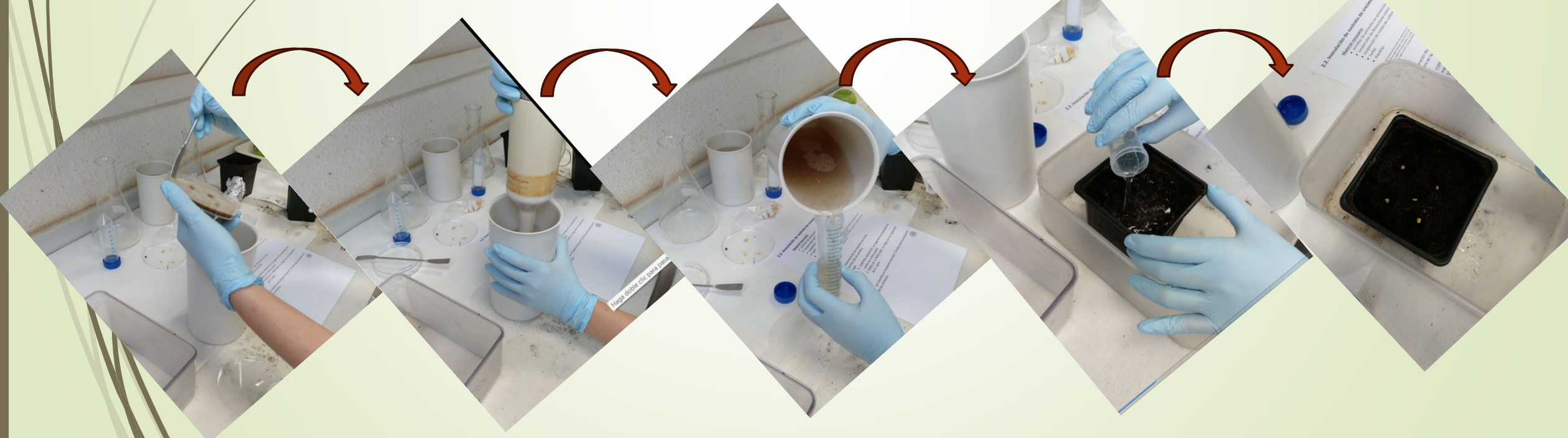


Screening for *Phytophthora capsici* resistance in pepper

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Control in a wet box

Progress of infection in inoculated stem



Dealing with abiotic stresses

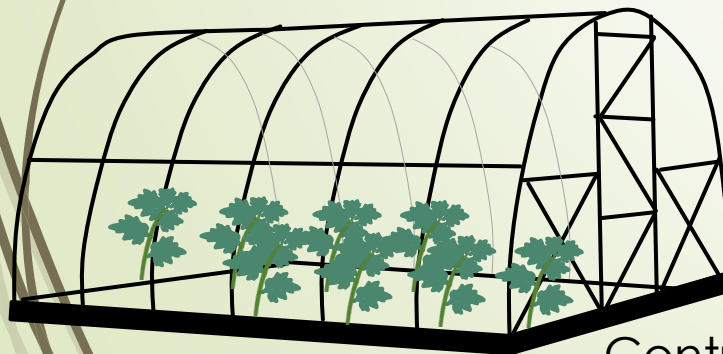
Aspects to take into account

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- Preliminary screening/phenotyping should be done with large number of accessions in field conditions specific to the stress
- Alternatively, trials in lab conditions can be carried out if reliable protocols are available
- Assays should be conducted under well-defined controlled conditions: the optimum or different levels of the stress can be applied



Screening for drought tolerance in the lab



Control



Treatment 1



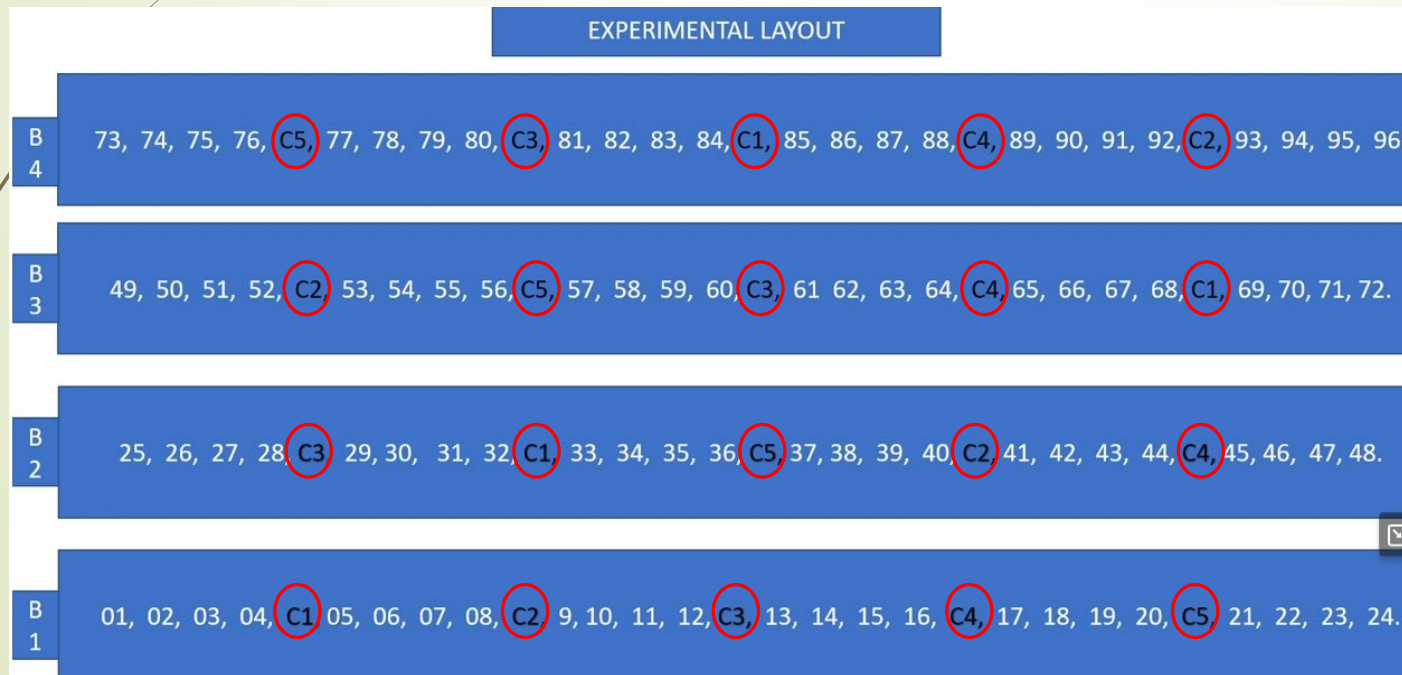
Treatment 2

Dealing with abiotic stresses

Aspects to take into account

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- Evaluation should be conducted with proper **experimental desing** depending on the **number of accessions** to be evaluated.
 - **Augmented block desing** (ABD) is being practiced in large number of accessions
 - For few acesions evaluation should be done in **randomized block design** (RBD) where the checks should be randomized along with the accessions in each replication

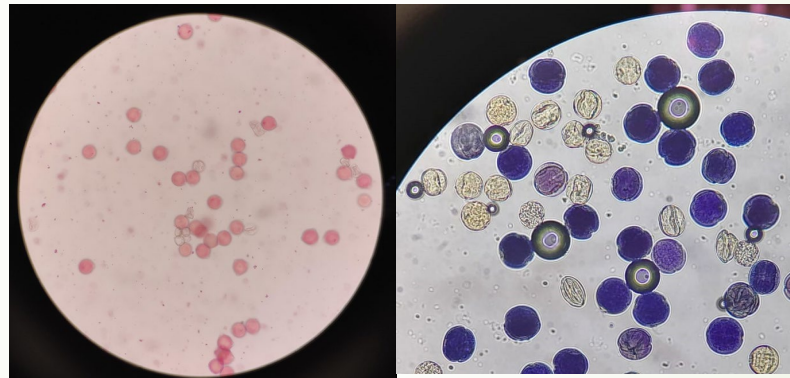


- In ABD, the **checks should be replicated in each block after separate randomization of checks within a block**. The number of checks will depend upon the crop and the parameters under study and representative of the type of germplasm. Three or more checks in which one national as well as one locally adapted check used for comparative assessment of germplasm

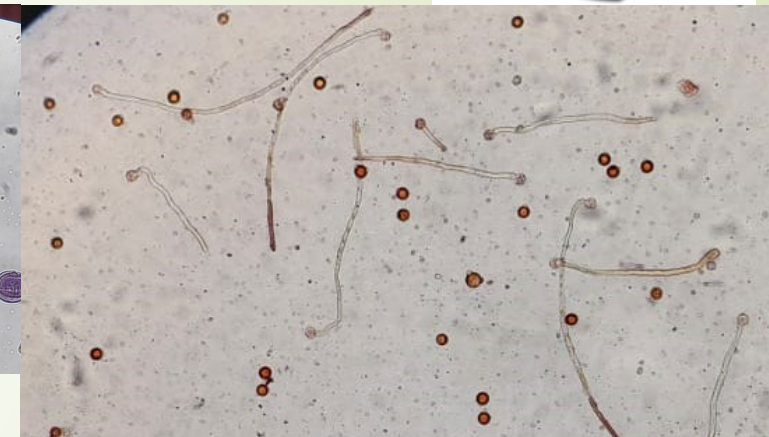
Dealing with abiotic stresses

Aspects to take into account: traits to be recorded

- ▶ Plant growth
 - ▶ Plant height, fresh weight, dry weight,
- ▶ Flowering
 - ▶ Earliness (male and female flowers in monoecious plants)
- ▶ Fruit set
- ▶ Pollen quality
- ▶ Fruit traits
 - ▶ Fruit weight
 - ▶ Fruit size
 - ▶ Fruit quality traits
- ▶ Physiological traits
 - ▶ Content of proline, glycine betaine, etc.
- ▶ Roots



Acetocarmine and
tetrazolium staining



Pollinic tube stained
with acetocarmine

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Angle of the root in the upper part



Length of the main roots



Diameter neck



Testing adaptation to drought: fine roots



Dealing with abiotic stresses

Looking for a needle in a haystack

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Dealing with abiotic stresses

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Genetic diversity and **digitalization** to save water resources in the cultivation of Cucurbitaceae



100% irrigation



50% irrigation



Salinity conditions

Platforms and Sensors



DJI Matrice 300 RTK + Zenmuse H20T

- **MTOM:** 9000 g
- **Optic sensor RGB:** CMOS 1/2.3", 12 MP
 - DFOV: 82.9°
 - Focal length: 4.5 mm
 - Aperture: f/2.8
 - Focus: 1 m a ∞
- **Thermal sensor:** Microbolometer Vox
 - DFOV : 40.6°
 - Focal length: 13.5 mm (equivalence a 58 mm)
 - Aperture: f/1.0
 - Focus: 5 m a ∞
- **Spectral band:** 8-14 μm

DJI Phantom 4 RTK Multispectral

- **MTOM:** 1487 g
- **Sensor:**
- Six sensors CMOS de 1/2.9", including RGB sensor for the visible spectrum and five monocrom sensors for multiespectral image
- **Bands**
 - Blue (B): 450 nm \pm 16 nm;
 - Green (G): 560 nm \pm 16 nm;
 - Red (R): 650 nm \pm 16 nm;
 - Red border (RE): 730 nm \pm 16 nm;
 - Near infrared (NIR): 840 nm \pm 26 nm

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Coded targets

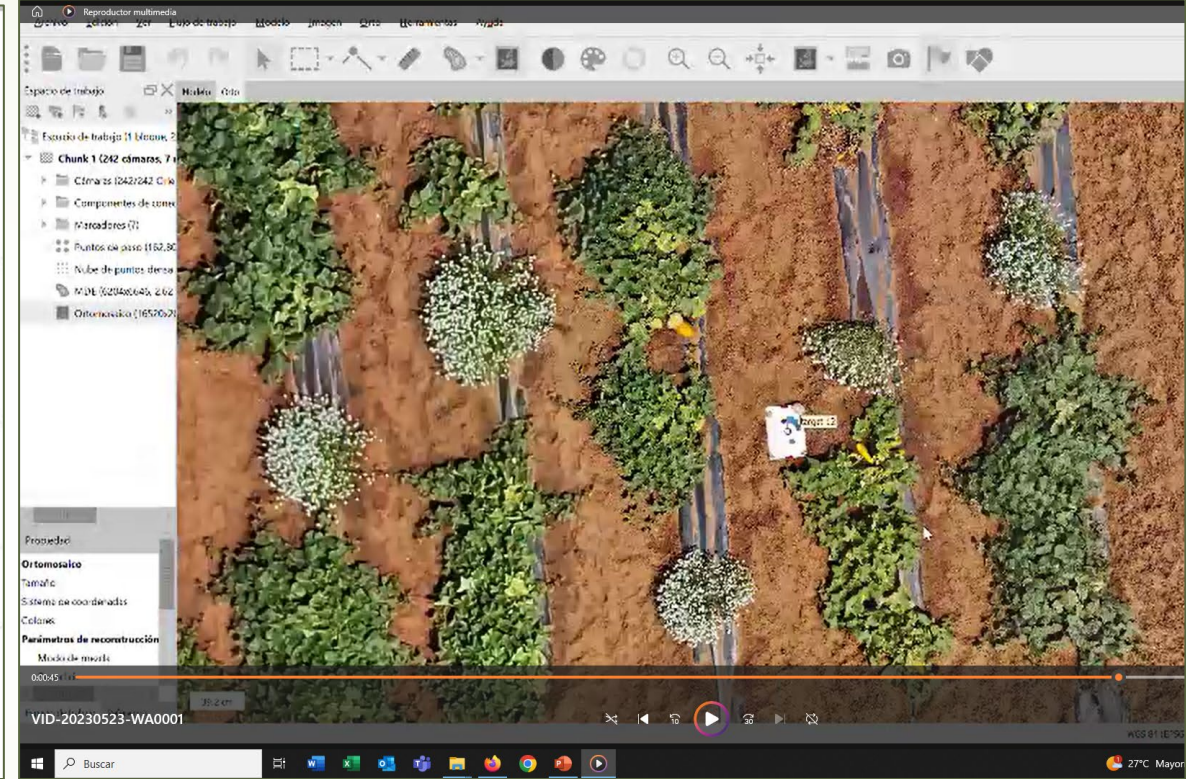
Data collection



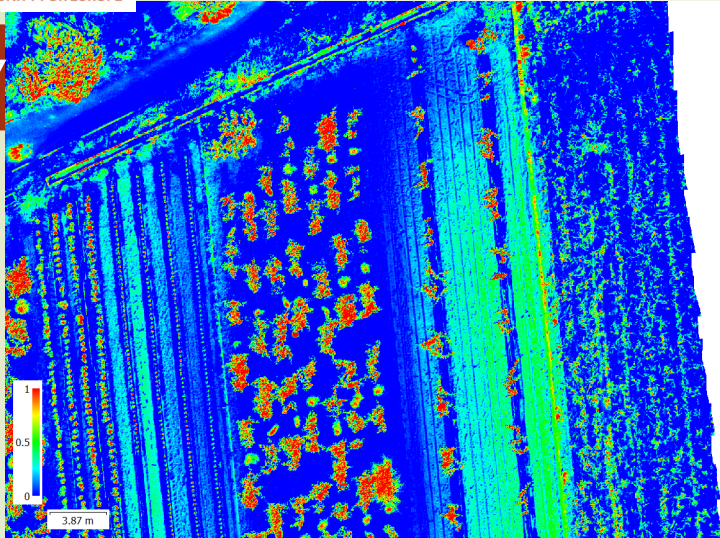
Flight
itinerary

Assessing abiotic stresses

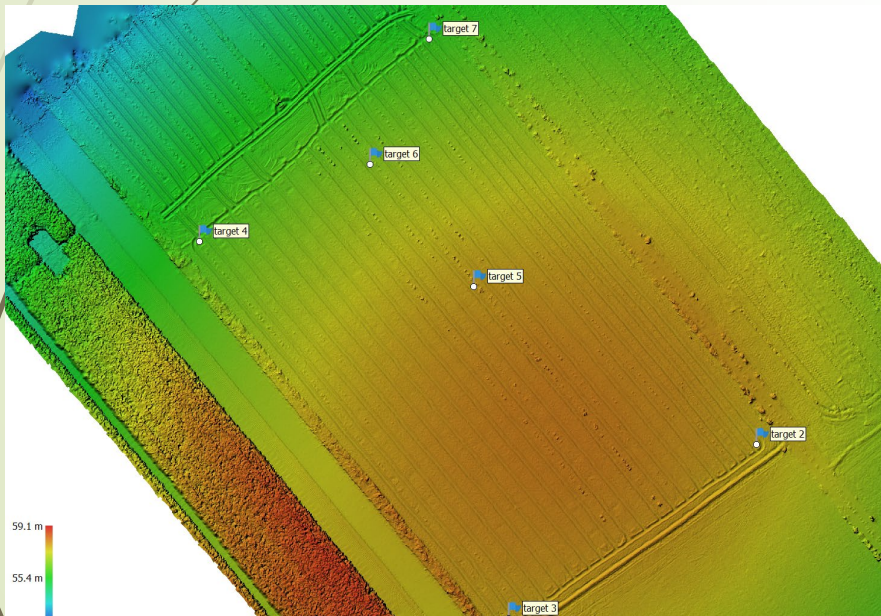
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Orthophoto in false color for sampling of the Vegetative Index



Digital elevation model for extraction of the digital terrain model and calculation of plant mass volumes



High resolution orthophoto for counting: 8.43 mm/pix

Data processing

Processing is carried out with Agisoft Metashape Professional in order to obtain **high-resolution orthophotos** and **digital elevation models** with which the **volumes of plant mass** will be calculated.

With the **high resolution orthophoto** it can be obtained together with the field data:

- Number of flowers per plant– statistical estimation
- Number of fruits– statistical estimation
- Fruit size– Manually (8mm resolution)
- Number of dead plants- Manually

With **multispectral shots and IR sensors**, information on:

- Vegetative aerial part (vegetative index), surface, color, and chlorophyll content
- Vegetative aerial part (vegetative index), temperature
- Vegetative aerial part (vegetative index), water content

Testing adaptation to drought in the laboratory: the poliethilenglicol

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Screening for adaptation to water deficit by *in vitro* culture

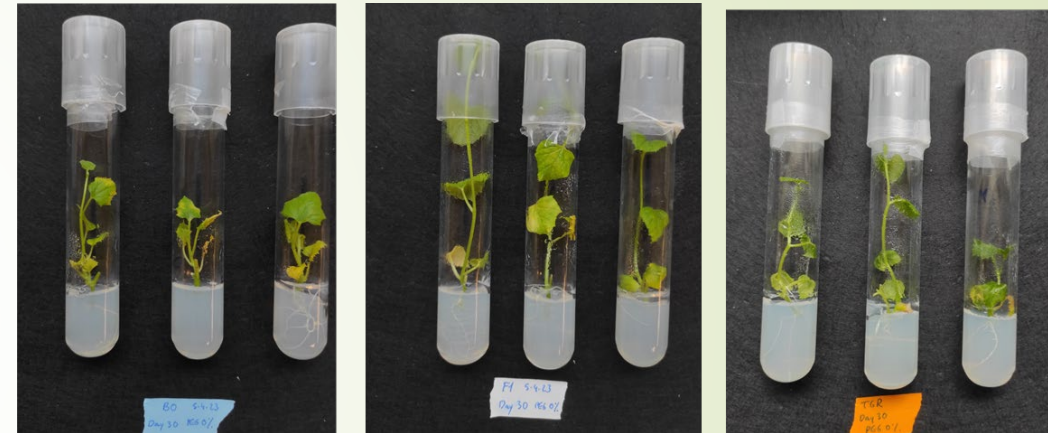
- ❑ Establishing *in vitro* culture: germination and micropropagation
- ❑ Determining PEG and culture media for *in vitro* selection

40

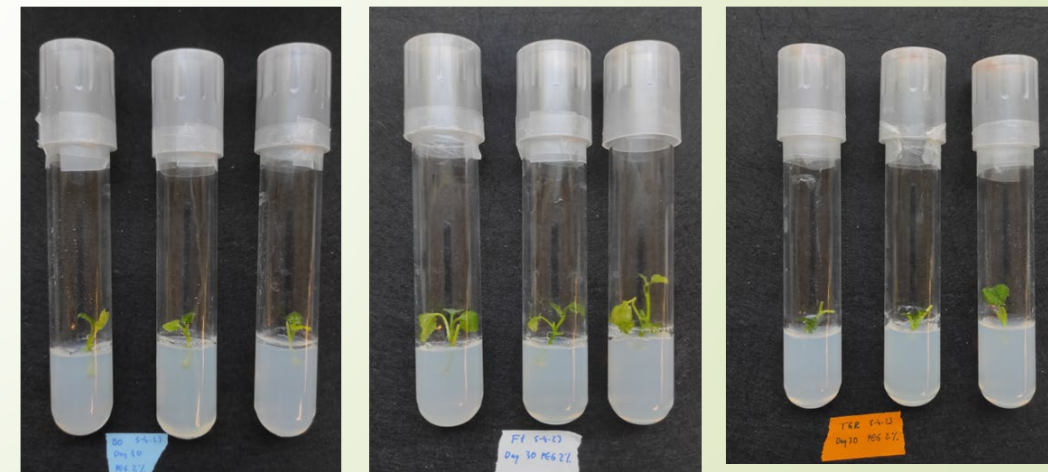
MATERIAL :

- ❑ Bola de oro (BO; *Cucumis melo* ssp. *melo*)
- ❑ 'TGR-1551' (TGR; *Cucumis melo* ssp. *agrestis*)
- ❑ F1 (BO x TGR)

Control: plants at 30 days in vitro culture without PEG



Plants at 30 days in vitro culture with PEG 2%



BO

F1

TGR

Study on the adaptation of germplasm to water stress in *in vitro* culture

Traits to be recorded

Genotype	% sprouting	% rooting	Index root development	Length aerial part	Number of leaves	Inhibition apical development
BO PEG 0	100	100	4,25	9,68	7,13	0
BO PEG 1	100	100	1,57	2,17	3,42	77,56
BO PEG 1,5	100	98,5	1,13	1,36	3,75	85,92
BO PEG 2	100	62,5	0,87	1,2	3,88	87,6
BO PEG 2,5	87,5	62,5	0,63	1,05	4,57	89,15
F1 PEG 0	100	100	4,63	12,07	9	0
F1 PEG 1	100	100	2,17	4,47	5	62,98
F1 PEG 1,5	100	100	1,8	2,74	5,6	77,29
F1 PEG 2	100	100	2	2,23	4,33	81,49
F1 PEG 2,5	87,5	87,5	1,43	2,24	4,86	81,41
TGR PEG 0	100	100	4,13	8,38	7,75	0
TGR PEG 1	100	100	1,88	2,13	5	74,49
TGR PEG 1,5	100	100	1,83	1,52	4,17	81,9
TGR PEG 2	100	100	1,5	1,18	3,13	85,92
TGR PEG 2,5	100	62,5	0,83	1,35	4,67	83,89

Results at 45 days

The PEG 1 concentration is already selective: it inhibits root formation and plant development.

The F1 shows more vigor and shows less inhibition.

There are no major differences between BO and TRG

Focused Identification of Germplasm Strategy (FIGS)

- It is an efficient technique that combines passport and agroclimatic data for a large number of accessions.

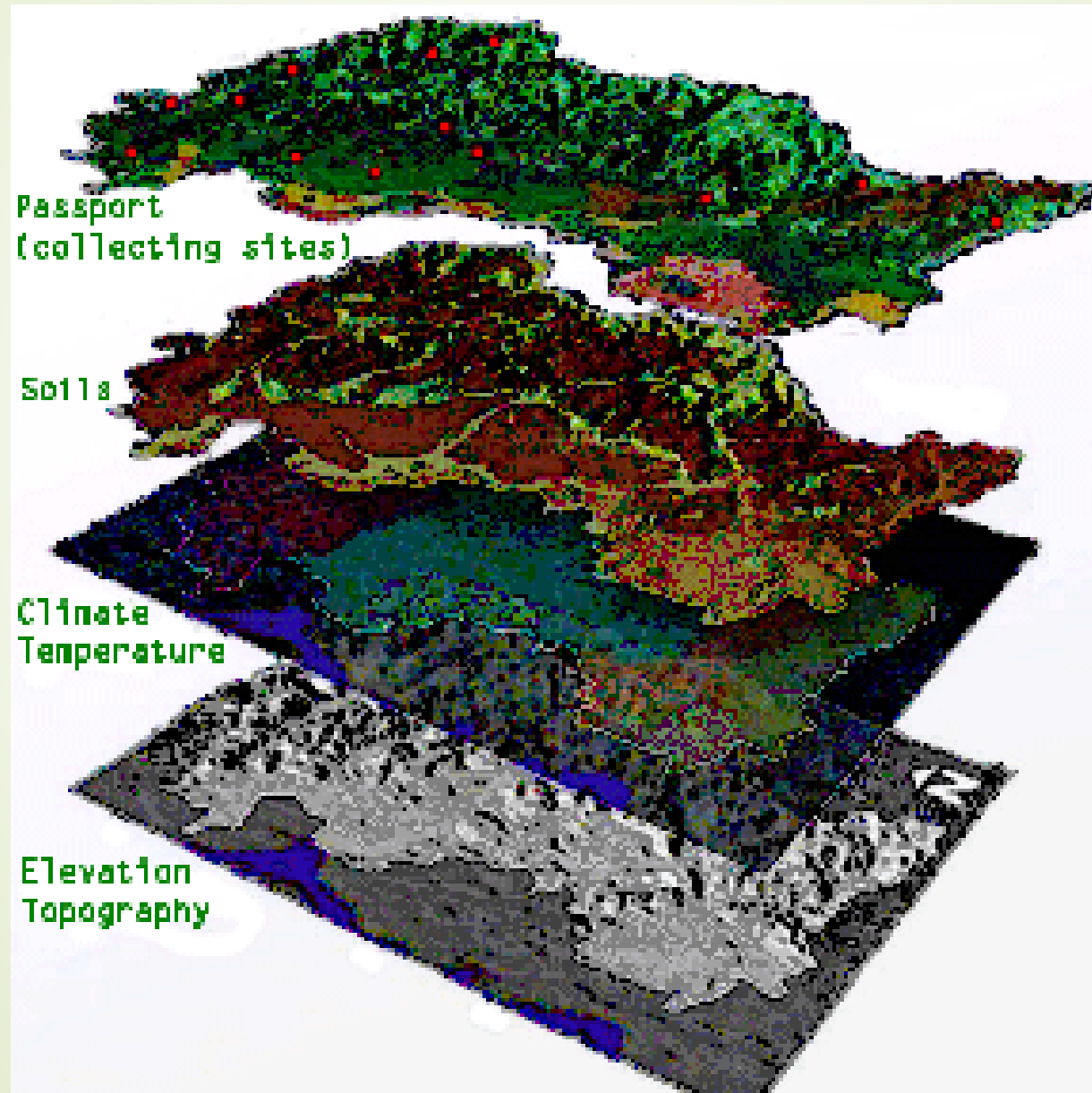
The breeder looks for specific characters in uncharacterized material in germplasm banks

Hipotesis:

If we can identify **geographic regions** in which **limiting environmental conditions** (biotic or abiotic stresses) have occurred during the evolutionary process, we can identify those accessions in a germplasm bank **originating from such regions** or from others subject to the same selection pressures.

Provision of passport, soil, climate and topography data

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Methodology (I)

- 1. Collect all passport data in a single database, verifying geographical coordinates
- 2. Obtaining environmental data from each collection location through geographic information systems (GIS)
- 3. Environmental information from collection sites can be used to create trait-specific subsets of materials for evaluation.

Methodology (II)

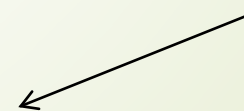
Take the 400 entries from the bank with known resistance to downy mildew.



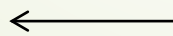
Use climatological and ecological data where they evolved to **determine the environmental profile associated with resistance**



Search the full database (16089) for **entries collected in places with similar environmental profiles**



Identification of 1,320 entries from Iran, Türkiye and Afghanistan



Screening: 16% of samples showed resistance (211 of 1320)

Methodology (III)

- Molecular characterization to determine allelic variants of the *Pm3* gene
 - More than half (111 of 211) had the resistance gene, some with unknown variants: 7 different alleles

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