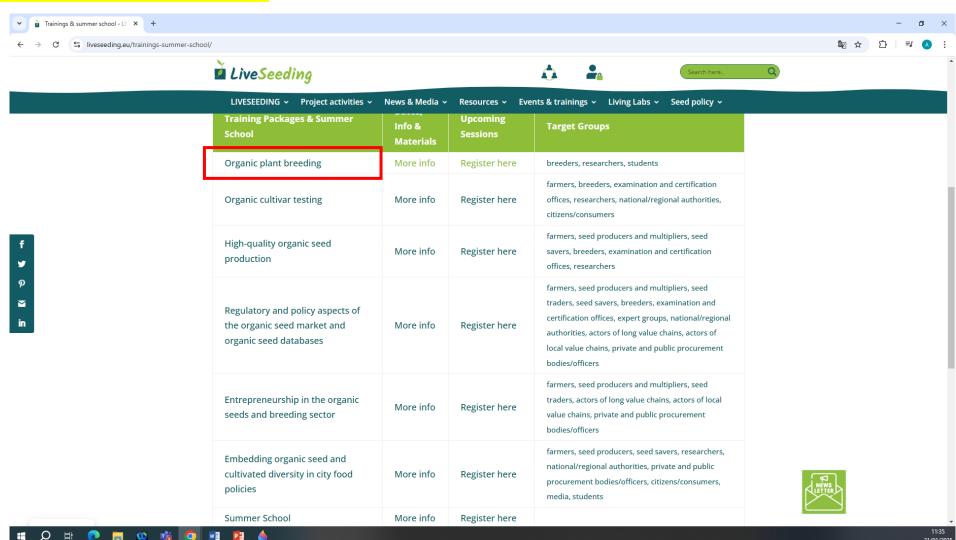


TRAINING IN ORGANIC BREEDING!



CONTEXT: Training in LIVESEEDING project

https://liveseeding.eu/trainings-summer-school/





Training in organic breeding organized in 5 Modules

- 1. Module 1 Plant Genetic Resources (PGRs): collection, conservation and exchange to support the increase of agrobiodiversity in farming systems
- 2. Module 2 Phenomics: approaches and tools for genetic resources and breeding material characterization FEBRUARY 3rd 2025, 9:00 to 17:30 CET
- 3. Module 3 Breeding methods fundamentals FEBRUARY 13th 2025, 9:00 to 18:00 CET
- **4. Module 4** Development and application of molecular methods in organic breeding MARCH 4th 2025, 9:00 to 18:00 CET
- 5. Module 5 Organic heterogeneous material (OHM) design and development MARCH 7th 2025, 9:00 to 18:00 CET





February 3rd 2025 - 9:00 to 17:30 CET



- Unit 2.1: Main descriptors used worldwide in characterizing plant genetic resources
 - 9:00-10:30 UPV (Adrian Rodríguez-Burruezo)
 - 10:30-11:00 Break
- Unit 2.2: Intro to ShineMas: a web tool dedicated to Seed Lots History, Phenotyping and Cultural Practices
 - 11:00-12:30 INRAe (Yannick de Oliveira, Isabelle Goldringer)
 - 12:30-14:00 Lunch Break
- Unit 2.3: Guidelines and examples of good practices in data management
 - 14:00-15:30 INRAe (Yannick de Oliveira, Isabelle Goldringer)
 - 15:30-16:00 Break
- Unit 2.4: Methods for phenotyping and selection of agronomic traits of interest in organic farming
 - 16:00-17:30 IPC (Pedro Mendes Moreira)

1 - An extra practical session to use the tool with own data is scheduled for FEB 10th (9-12h)





Module 2: Phenomics: approaches and tools for genetic resources and breeding material characterisation

Unit 2.1: Main descriptors used worldwide in characterizing plant genetic resources

Author: Adrian Rodríguez-Burruezo





Funded by the European Union, the Swiss State Secretariat for Education, Research and Innovation (SERI) and UK Research and Innovation (UKRI).





Planned for today

DYNAMIC MIXTURE OF:

- 1. Presentation about main topics on phenomic descriptors: utility, types, examples of descriptors, management of data, knowledge, additional material (50 min)
- 2. Guided virtual visits (about 10-15 min)
- 3. Fast quiz (about 10 min) ***
- 4. Debate, Wrap up & Proposed homework (about 10 min) ***
- 5. QUESTIONS: THROUGH THE CHAT (Petra Jelincic will manage)



Characterization of plant genetic resources (PGRs)

- > PGRs can be: Landraces/heirlooms, breeding lines, wild relatives,...
- For their efficient management, it is of paramount importance... being aware of their diversity
- ➤ HOW? Catalogue PGRs based on objective description
- > Actors using:

Seedbanks (essential) Level 2 info: "Characterization data"

Breeders, farmers

Seed companies



Descriptors?

- ➤ Many ways to describe variation:..... ¿?
- DESCRIPTORS = Phenological traits*
- *High inheritance (which means.....??)
- Compiled in a list, ordered by blocks

1. Plant vegetative traits:

General development

Branches

Leaves



2. Plant reproductive traits:

Flowers/inflorescences

Fruits/infrutescences

Seeds

















Descriptors. Why?

Essential to manage agrobiodiversity/PGRs in hands of:
Seedbanks (germplasm collections)
Breeders (germplasm, prebreeding materials, breeding lines, etc.)
☐ Farmers (landraces, traditional varieties, ecotypes, etc.)
Being aware of the materials you have
Check the level of diversity you have
Comparison to other collections in hands of other actors
Detecting duplicates, establishing core collections, know the types you need to enrich the diversity of your collections
LIMITATION: there are many descriptors, but better to work with those which best encompass the diversity present on a crop and its relatives (comprehensive & discriminating ability)



Examples of descriptors

☐ BIOVERSITY International (former IPGRI)

Alliance

https://alliancebioversityciat.org/





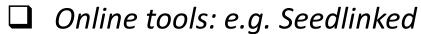


☐ High throughput digital tools

e.g. Tomato analyser

Developed in E. Van der Knapp lab

https://vanderknaaplab.uga.edu/tomato-analyzer/





"We connect plant growers and their data to help breed, source, and harvest the best seeds"

https://seedlinked.com



PRACTICAL:

- 1. Guided visit to BIOVERSITY Descriptors
 - + example with excel compiled data
- 2. Guided visit to TOMATO ANALYZER webpage
 - + example of UPV lab results with own data
- 3. Seedlinked (short)



1. VISIT TO:

Alliance





https://alliancebioversityciat.org/

PLEASE.

STILL DO NOT ENTER!!!



Public/Institutional Seedbanks Module 1 – Unit 1

1. VISIT TO:

EXAMPLE UPV



TYPE Original Research PUBLISHED 30 October 2024 DOI 10.3389/fpls.2024.1435427

Alliance







OPEN ACCESS

EDITED BY

Francesco Sunseri. Mediterranea University of Reggio Calabria,

REVIEWED BY Andrea Mazzucato. University of Tuscia, Italy Lorenzo Raggi,

University of Perugia, Italy

*CORRESPONDENCE

Adrián Rodríguez-Burruezo adrodbur@upvnet.upv.es

RECEIVED 20 May 2024 ACCEPTED 17 September 2024 PUBLISHED 30 October 2024

Genetic diversity, population structure, and phylogeny of insular Spanish pepper landraces (Capsicum annuum L.) through phenotyping and genotypingby-sequencing

Neus Ortega-Albero¹, Lorenzo Barchi², Ana Fita¹, Miguel Díaz¹, Felipe Martínez¹, Joana-Maria Luna-Prohens³ and Adrián Rodríguez-Burruezo1*



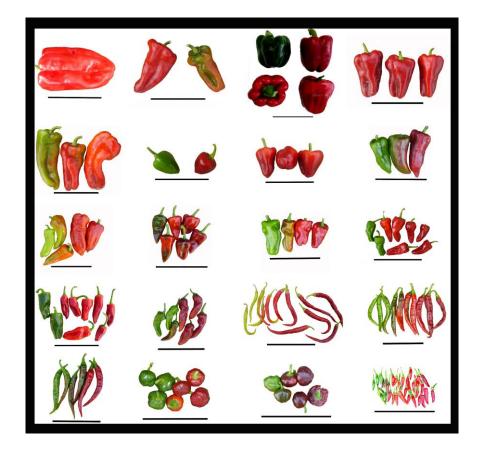
1. VISIT TO:

Alliance





Example fruit characterization. *Capsicum* peppers COMAV -UPV





2. VISIT TO:

TOMATO ANALYZER

Developed in E. Van der Knapp lab

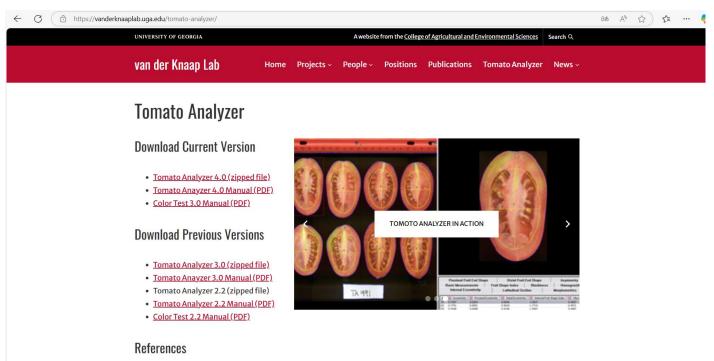
PLEASE.

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BASED ON:

CUT FRUIT SCANNED PICTURES

https://vanderknaaplab.uga.edu/tomato-analyzer/





OTHER EXAMPLES OF Tomato Analyzer:

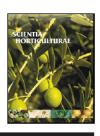
Scientia Horticulturae 164 (2013) 625-632

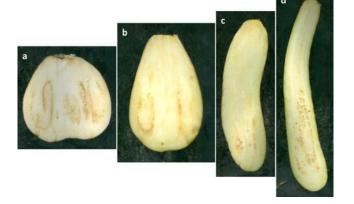


Contents lists available at ScienceDirect

Scientia Horticulturae

journal homepage: www.elsevier.com/locate/scihorti



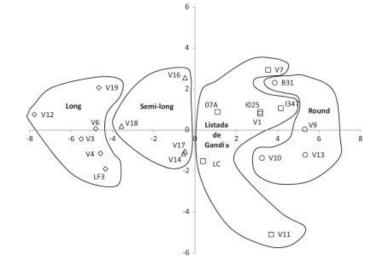


Phenomics of fruit shape in eggplant (Solanum melongena L.) using Tomato Analyzer software

Maria Hurtado, Santiago Vilanova, Mariola Plazas, Pietro Gramazio, F. Javier Herraiz, Isabel Andújar, Jaime Prohens*

Instituto de Conservación y Mejora de la Agrodiversidad Valenciana, Universitat Politècnica de València, Camino de Vera 14, 46022 Valencia, Spain







3. SHORT VISIT TO:



https://seedlinked.com

PLEASE.

STILL DO NOT ENTER!!!



EXAMPLE OF UTILITY BY USING BOTH:

Descriptors & Tomato Analyzer (alone and together)

Scientia Horticulturae 265 (2020) 109245



Contents lists available at ScienceDirect

Scientia Horticulturae

journal homepage: www.elsevier.com/locate/scihorti



Phenomics of elite heirlooms of peppers (*Capsicum annuum* L.) from the Spanish centre of diversity: Conventional and high-throughput digital tools towards varietal typification



Leandro Pereira-Dias^a, Ana Fita^{a,*}, Santiago Vilanova^a, Elena Sánchez-López^b, Adrián Rodríguez-Burruezo^b

^b Instituto Murciano de Investigación Agraria y Alimentaria, C/Mayor s/n, La Alberca-Murcia, 30150, Spain

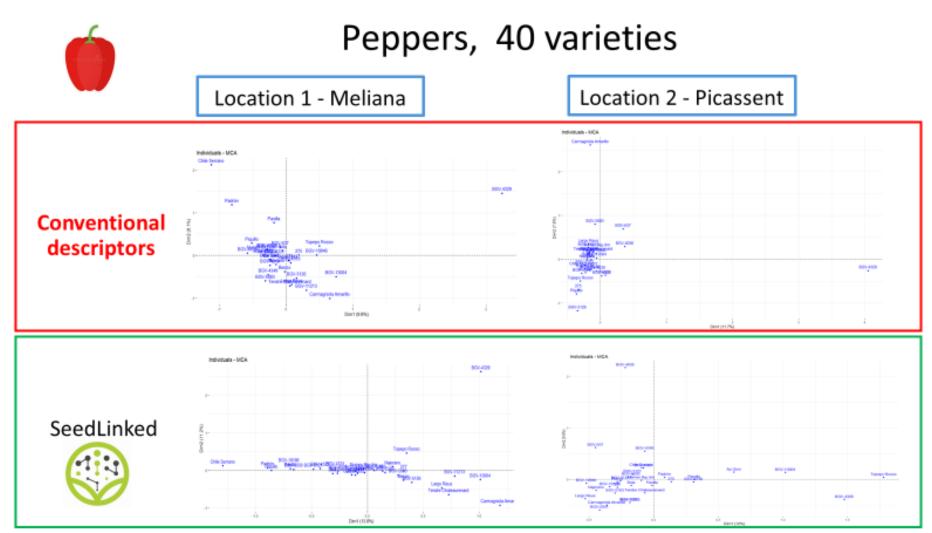


^a Instituto de Conservación y Mejora de la Agrodiversidad Valenciana, Universitat Politècnica de València, Camino de Vera s/n, 46022 Valencia, Spain

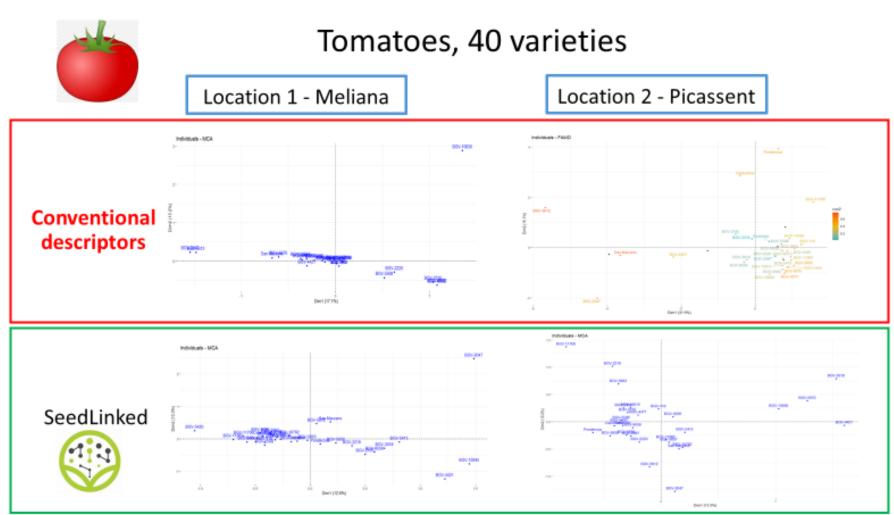
EXAMPLE OF UTILITY BY USING BOTH:

Descriptors & SeedLinked (alone and together)











FAST QUIZ

- Question 1: Why are descriptors useful?
- Question 2: Name 5 species covered by BIOVERSITY descriptors
- ☐ Question 3: What raw info is essential to work with Tomato Analyzer
- Question 4: What is SEEDLINKED?

Send to:

<u>adrodbur@doctor.upv.es</u> and <u>petra.jelincic@ips-konzalting.hr</u>

<u>In 10 min</u>



DEBATE

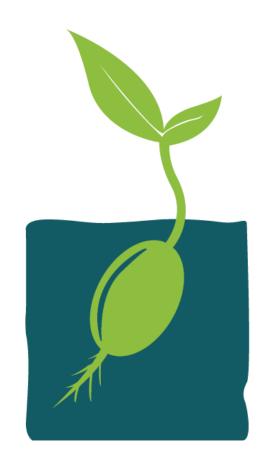
☐ Revise the quiz in common

☐ Have you heard about these tools before?

- ☐ What's the most useful for your work at this moment
- Other questions and doubts



WRAP UP



LiveSeeding

- What we have learned today?
- ☐ Proposed homework: Case of study. 2-3 accessions of one species of your interest, close to you, and characterise as much as possible, according to the corresponding BIOVERSITY descriptors. Prepare an excel with the data compiled.

And send to:

<u>adrodbur@doctor.upv.es</u> and <u>petra.jelincic@ips-konzalting.hr</u>

By next Monday 10th february

Additional available materials

- 1. Operational Genebank Manual. Centre for Genetic Resources, The Netherlands (CGN-WUR). And others manuals: https://www.ecpgr.org/aegis/aquas/genebank-manuals
- 2. IPGRI/Bioversity international descriptors: https://cgspace.cgiar.org/collections/835fa638-0167-4669-9532-ffc488facc94
- 3. Gotor, E., Alercia, A., Rao, V.R., Watts, J., Caracciolo, F., 2008. The scientific information activity of Bioversity International: the descriptor lists. Genet. Resour. Crop Evol. 55: 757–772. https://doi.org/10.1007/s10722-008-9342-x
- 4. Tomato Analyzer: https://vanderknaaplab.uga.edu/tomato-analyzer/
- 5. Tomato Analyzer VIDEO REFERENCE: Rodriguez, G.R., Moyseenko, J.B., Robbins, M.D., Huarachi Morejon, .N, Francis, D.M., van der Knaap, E.(2010). Tomato Analyzer: A Useful Software Application to Collect Accurate and Detailed Morphological and Colorimetric Data from Two-dimensional Objects. JoVE 37.
- 5. Seedlinked: https://seedlinked.com/
- 6. Ortega-Albero N, Barchi L, Fita A, Diaz M, Martinez F, Luna-Prohens J-M and Rodriguez-Burruezo A (2024) Genetic diversity, population structure, and phylogeny of insular Spanish pepper landraces (Capsicum annuum L.) through phenotyping and genotyping-by-sequencing. Front. Plant Sci. 15:1435427. doi: 10.3389/fpls.2024.1435427
- 7. Pereira-Dias L, Fita A, Vilanova S, Sánchez-López E, Rodríguez-Burruezo A. 2020. Phenomics of elite heirlooms of peppers (Capsicum annuum L.) from the Spanish centre of diversity: Conventional and high-throughput digital tools towards varietal typification. Scientia Horticulturae 265: 109245. https://doi.org/10.1016/j.scienta.2020.109245.















































































