

## **WORKSHOP**

# Advancing Berry Breeding: From Phenotyping to Genomic Innovation



Also accessible online

### **AGENDA**

17:30

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13:30	REGISTRATION
14.00	WELCOME AND INTRODUCTION OF THE BREEDINGVALUE PROJECT Bruno Mezzetti, UPM
14:15	<b>WELCOME</b> Agata Gulisano, Plant Breeding and Genetic Resources Policy Officer, EC
14:20	BERRY GENETIC RESOURCES: Importance of a large germplasm and breeding material for BV; Results of phenotyping Klaus Olbricht, Hansabred
14:50	NEW PHENOTYPING METHODS – Application of NIR spectroscopy for the rapid quality assessment of different strawberry cultivars  Manuela Mancini, UPM
15:05	NEW PHENOTYPING METHODS – Strawberry root phenotyping Tuuli Haikonen, LUKE
15:20	NEW PHENOTYPING METHODS – Use of ground based hyperspectral imaging for high throughput field phenotyping Dominic Williams, JHI
15:35	How can we translate strawberry flavour into chemical compounds for breeding? Sonia Osorio, UMA
15:55	Coffee break
16:25	SENSORY QUALITY AND CONSUMER ACCEPTANCE OF BERRIES: Towards satisfied berry consumers – an introduction to activities Saila Karhu, LUKE
16:40	SENSORY QUALITY AND CONSUMER ACCEPTANCE OF BERRIES: Short and Long-Term Future Visions for Enhancing Fruit Quality in Cultivated Berries Terhi Latvala, LUKE
17:00	SENSORY QUALITY AND CONSUMER ACCEPTANCE OF BERRIES: Panel and consumer contribution to characterize and valorize berry GenRes Stefano Predieri, CNR-IBE
17:20	RESULTS FROMTHE OPEN CALL #4: Toolkits of sensorial qualityassessment for berry breeders Stefano Predieri, CNR-IBF

• Lithuanian Research Centrefor Agriculture and Forestry, Neringa Rasiukevičiūtė

Experiences and feedback from Open Call #4 participants:

Angus Soft Fruits, Lucy WilkinsG-Berries, Alessandro Gasparrini

17:45	From Bits to Bytes: Software Development for Genetics, Genetic Resources and Plant Pre- Breeding Sebastian Raubach, James Hutton Institute
18:00	Integrating and Maintaining BerryData: The Germinate Database and beyond Björn Usadel / Elisa Senger, Forschungszentrum Jülich
18:30	End of day 1
Thurso	day, 3rd April 2025
09:00	WORKSHOP ON MARKER-ASSISTED SELECTION, GENOME-WIDE ASSOCIATION STUDIES AND GENOMICSELECTION IN BERRY BREEDING PROGRAMS – WELCOME AND INTRODUCTION
09:15	Tracing strawberry population structures: The legacy of domestication and intertwined breeding  Jakob Junkers, NIBIO
09:35	Multi-model GWAS reveals key loci for horticultural traits in reconstructed garden strawberry Attiq Rehman / Tuuli Haikonen, LUKE
09:55	Genome-wide association studies (GWAS) in a diverse strawberry collection from IFAPA unveil loci controlling agronomic and fruit quality traits  Cristina Castillejo, IFAPA
10:15	The BreedingValue low-density array for MAS in strawberry Iraida Amaya, IFAPA
10:40	💪 Coffee break
11.10	RESULTS FROM THE OPEN CALL #1 Marker-Assisted Selection (MAS) in strawberry Juliette Bénéjam, INRAE
11:30	Perspectives, practical applications, challenges, and innovations from Open Call participant: • Angus Soft Fruits, Lucy Wilkins
11:40	Marker-assisted selection in raspberry: challenges and opportunities in the -omics era.  Brezo Mateos, JHI
12:00	RESULTS FROM OPEN CALL #3: Marker-Assisted Selection (MAS) and Genome wide association studies (GWAS). GWAS in Raspberries: Developmental and Fruit Traits.  Philip Greenspoon, JHI (BioSS)
12:20	<ul> <li>Experiences and feedback from Open Call #3 participants:</li> <li>Angus Soft Fruits, Lucy Wilkins</li> <li>G-Berries, Alessandro Gasparrini</li> </ul>
12:35	Lunch break
14:30	INTRODUCTION TO GENOMIC SELECTION (GS) AND AN EXAMPLE FROM THE BREEDINGVALUE project: Brief introduction on GS. What is it, how is done, what benefits for implementation in breeding programs  Jahn Davik, NIBIO
14:50	The benefits of GS in a cultivated strawberry breeding programme: an example of powdery mildew resistance

Juliette Bénéjam, INRAE

# 15:10 RESULTS FROM OPEN CALL#2: Genomic prediction for strawberry. Predicting strawberry powdery mildew susceptibility in strawberry breeding material

Jahn Davik, NIBIO

### **15:30** Experiences and feedback from Open Call #2 participant. Current and future benefits:

• Fresh Forward, Johan Willemsen

### 15:40 ROUND TABLE ON MAS AND GENOMIC SELECTION OF BERRIES

Moderators: Jahn Davik, Beatrice Denoyes, Iraida Amaya, Bruno Mezzetti.

With the objective of encouraging reflection on how we can apply genomic tools in plant breeding. Focus:

- How MAS and genomic selection are changing berry breeding strategies.
- Success storiesor challenges in using thesetechniques in real-world berry breeding programs.
- Integration of traditional breeding methods with genomic tools.
- The feasibility and challenges of integrating genomic selection in small-scale vs. large-scale berry breeding.
- Future trends in the use of genomics for disease resistance or yield improvement in berries.
- How has marker-assisted selection advanced berry breeding, and what limitations remain?
- What role does GWAS play in uncovering traits of interest in berries, and how can these findings be used in breeding programs?
- What are the biggest hurdles in applying genomic selection in commercial berry breeding?
- How can breeders ensure that genomictechniques are cost-effective and accessible to small-scale farmers?

### 17:00 Conclusions and Closing Remarks

Bruno Mezzetti



