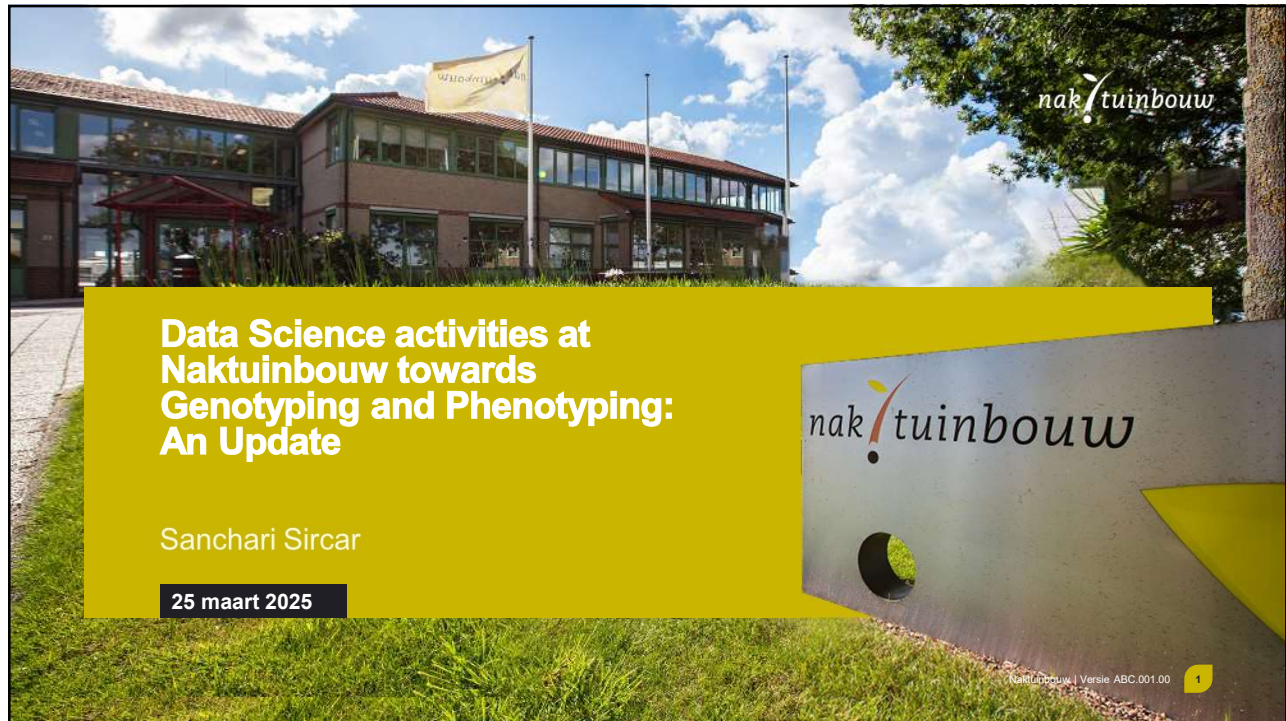


Technical Working Party on Testing Methods and Techniques**TWM/3/16****Third Session****Beijing, China, April 28 to May 1, 2025****Original:** English**Date:** March 26, 2025

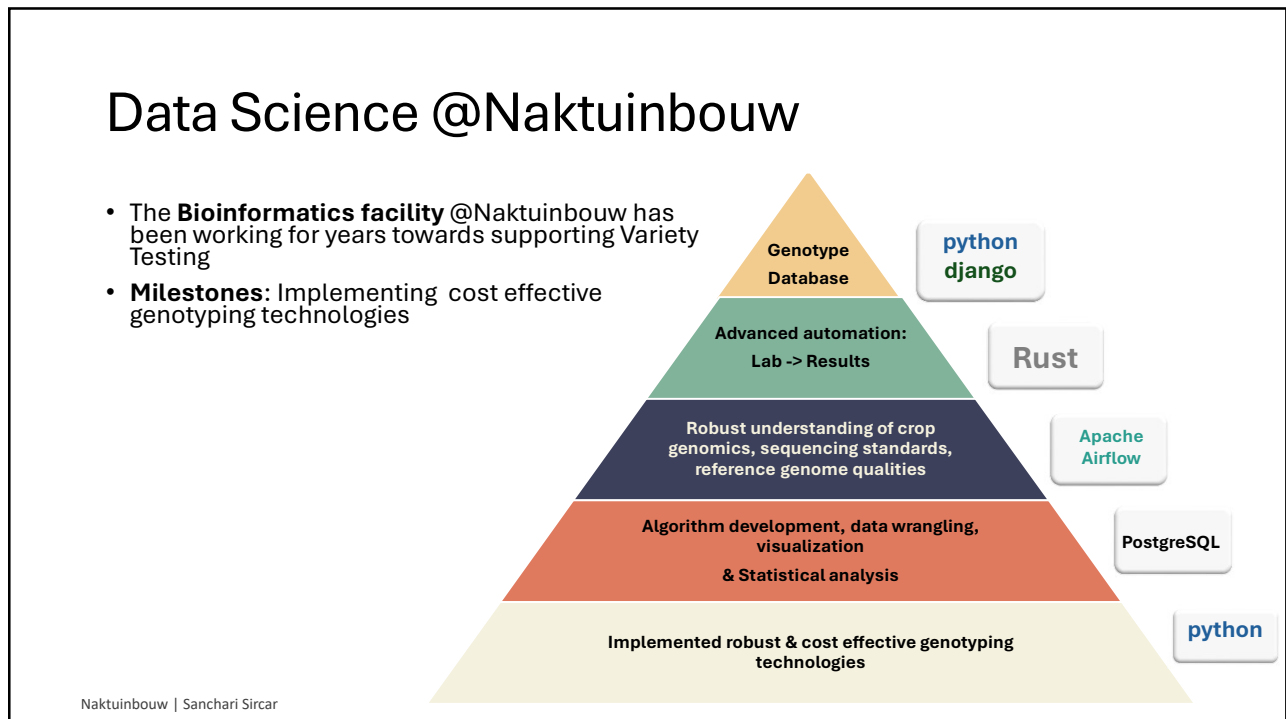
DATA SCIENCE ACTIVITIES AT NAKTUINBOUW TOWARDS GENOTYPING AND PHENOTYPING: AN UPDATE*Document prepared by an expert from the Netherlands (Kingdom of)**Disclaimer: this document does not represent UPOV policies or guidance*

1. The Bioinformatics facility at Naktuinbouw has been instrumental in supporting variety identification and research. Our work includes identifying molecular markers using next-generation sequencing, developing algorithms for optimization and automation, and conducting statistical analysis and reporting. This has driven us to create uniform data platforms that enable meaningful comparisons.
2. Building on this foundation, our next step is to standardize image analysis and phenotyping at Naktuinbouw. This presentation will offer a sneak peek into our upcoming projects on image analysis, workflow developments, AI and other collaborative efforts
3. The annex to this document contains a copy of a presentation “Data Science activities at Naktuinbouw towards Genotyping and Phenotyping: An Update”, to be made by an expert from the Netherlands (Kingdom of), at the third session of the TWM.

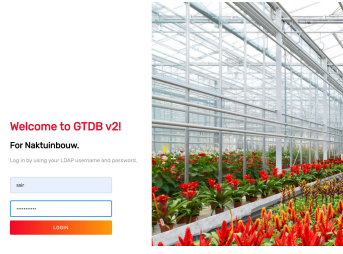
[Annex follows]



1



2



Welcome to GTDB v2!
For Naktuinbouw.
Log in by using your LDAP username and password.

Log in

Run Info

Run ID: 5122724 Run Date: 14-05-2025 Species: Tomato Machine ID: ST-63077 Run number: 1285 Flowcell: H830CER2 Lane: 1


TotalClustersRaw: 0 TotalClustersPF: 0 Total reads: 105 Users: 1 Run type (GSE): High-Output

Primer Info

Run number	Species	SNP name	registered	Start	End	Type	Comment	temperature	C9 (Flow)	strand	count	count	count	sequence
98	tomato	5122724_5122846_PCR_product	<input type="checkbox"/>	5122724	5122846	PCR_product		128	0.3442622958819672	37	42	27	15	TGAGGAAAGTGTATGATGGGTTGA
98	tomato	5122724_5122746_forward_primer	<input type="checkbox"/>	5122724	5122746	forward_primer		66	0.375	5	9	3	8	TGAGGAAAGTGTATGATGGG
98	tomato	5122822_5122846_reverse_primer	<input type="checkbox"/>	5122822	5122846	reverse_primer		148	0.47826889552174	5	7	10	1	CTGACACCACTATCTATGAC
98	tomato	5122821_5122822_SNP	<input type="checkbox"/>	5122821	5122822	SNP		2	0	0	1	0	0	A

SNP Info

Marker frequency plot




Check ploidy
Check reference base
Find sequences
PCR product
Flanking sequence
Primer sequences

Naktuinbouw | Sanchari Sircar

3

Run statistics

- Optimization
 - Merker team
- Flowcell info
- Fastp reports
 - Raw data filtering info
 - Read lengths
 - Read quality
- Summaries of read data
- Base distributions
- Flowcell distributions
- And much more.



25 March 2025

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4

GFF3 data

All gff info stored

Show 10 entries

Run number	Species	SNP name	rejected	Start	End	Type	Comment	temperature	CG (float)	tcount	acount	ccount	gcount	sequence
98	tomato	SL3.0Ch11:5122821	<input type="checkbox"/>	5122724	5122846	PCR_product		328	0.3442622950819672	37	43	27	15	TAAAGAAAAGTGGTCATGATTGGGGTCCA
98	tomato	SL3.0Ch11:5122821	<input type="checkbox"/>	5122724	5122748	forward_primer		66	0.375	6	9	1	8	TAAAGAAAAGTGGTCATGATTGGG
98	tomato	SL3.0Ch11:5122821	<input type="checkbox"/>	5122823	5122846	reverse_primer		68	0.4782608695652174	5	7	10	1	CGTCACCCCAATCATCTAC

SL3.0Ch11:5122821

SL3.0Ch11:5122821

SL3.0Ch11:5122821

SL3.0Ch11:5122821

SL3.0Ch11:5122821

SL3.0Ch11:5122821

- Primer Optimization
 - SNP Discovery -> SNP Validation
 - SNPs thrown out
 - Rerun
 - Repetitive process
- Interactions between
 - Primers for this run
 - Visualizes bindings with Sankey Plots

SL3.0Ch11:5122821

SL3.0Ch11:5122821

SL3.0Ch11:5122821

SL3.0Ch11:5122821

SL3.0Ch11:5122821

SL3.0Ch11:5122821

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5

5

Heatmap

Heatmap

Heatmap

Heatmap

Heatmap

Heatmap

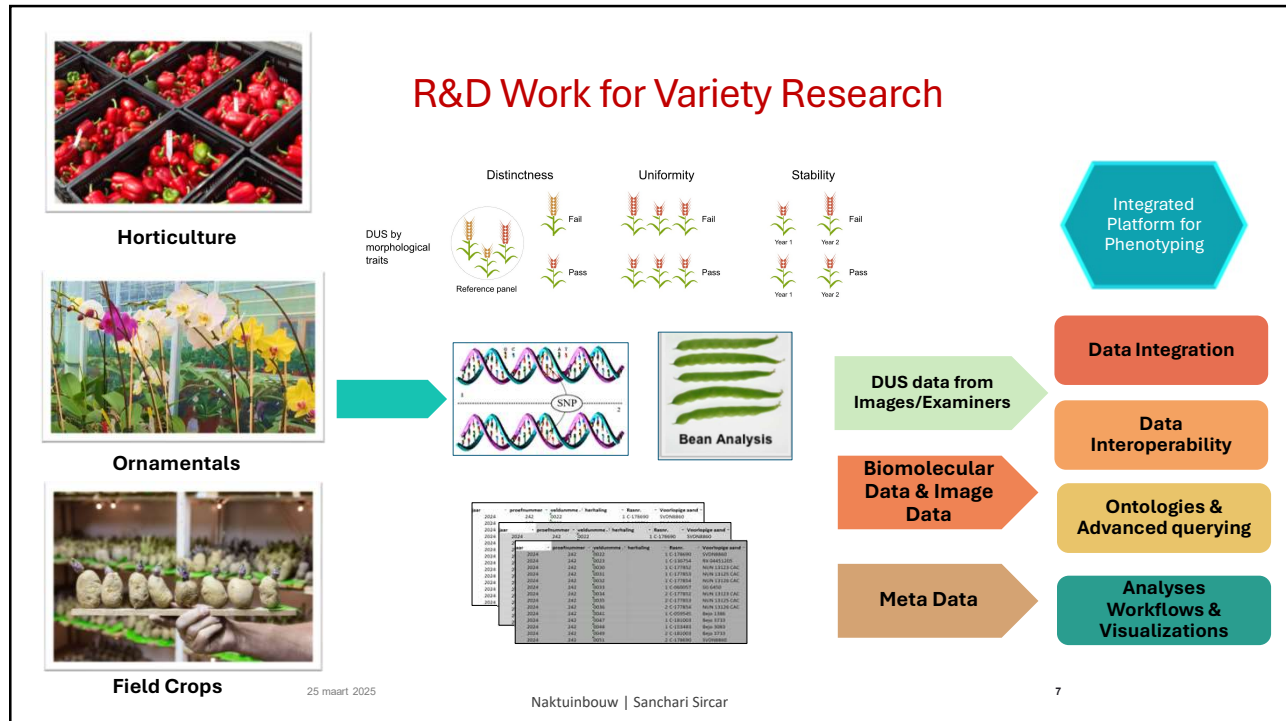
- For a given species, compare within and between runs
- Dynamic dendrogram and heatmap
 - Collapsible nodes
 - Clickable names (and searchable)
 - Lots of filtering
 - Depth
 - Individual sample selection
 - Heatmap hovertools, showing haplotypes or depth.

Dendrogram

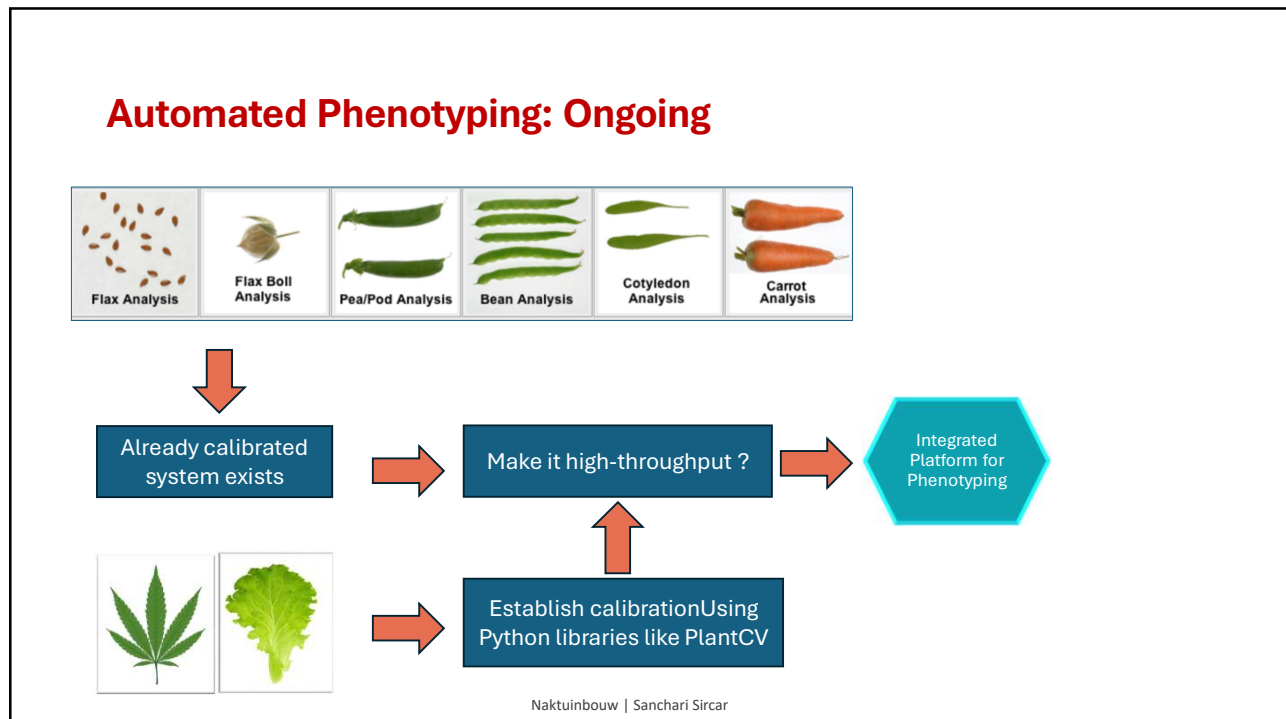
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6

6

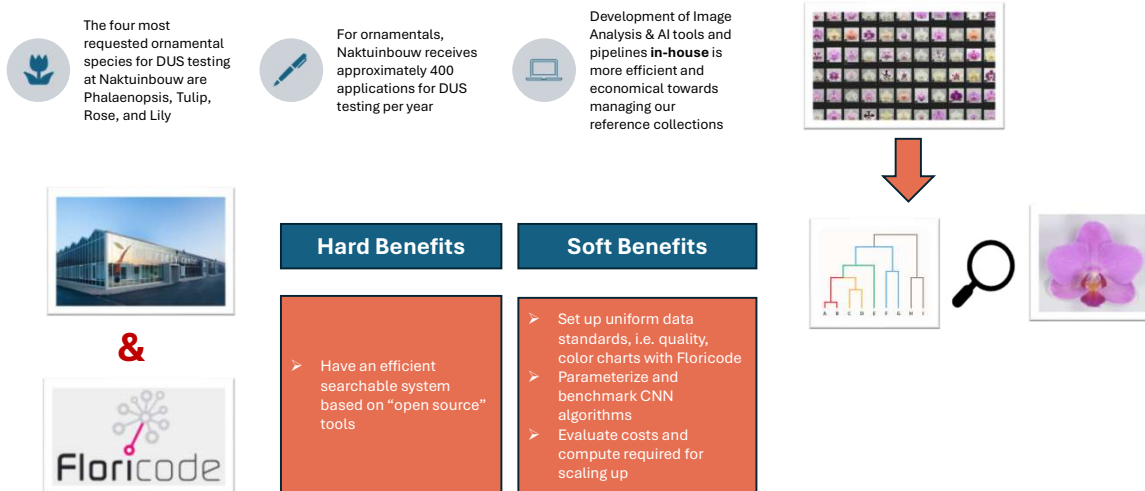


7



8

Managing Our PhotoDatabases: Ongoing



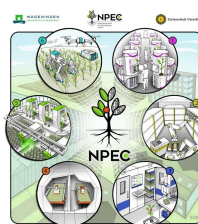
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9

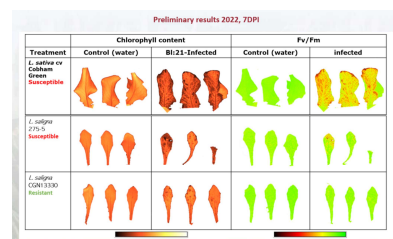
Future Collaborations

- Non-competitive Collaborations
- Comparing analyses
 - Pattern recognitions
 - Comparing data standards
 - Resources & Algorithms
 - Common controls

The Netherlands Plant Eco-phenotyping Centre: NPEC



P4P: Phenotyping 4 Profit For disease resistance in plants



Digital Phenotyping of downy mildew disease in Lettuce

PPS INSPIRE: INnovations for Smart Plant Insect Resistance Evaluation and testing



Detection of whitefly eggs from tomato & potato leaves using AI

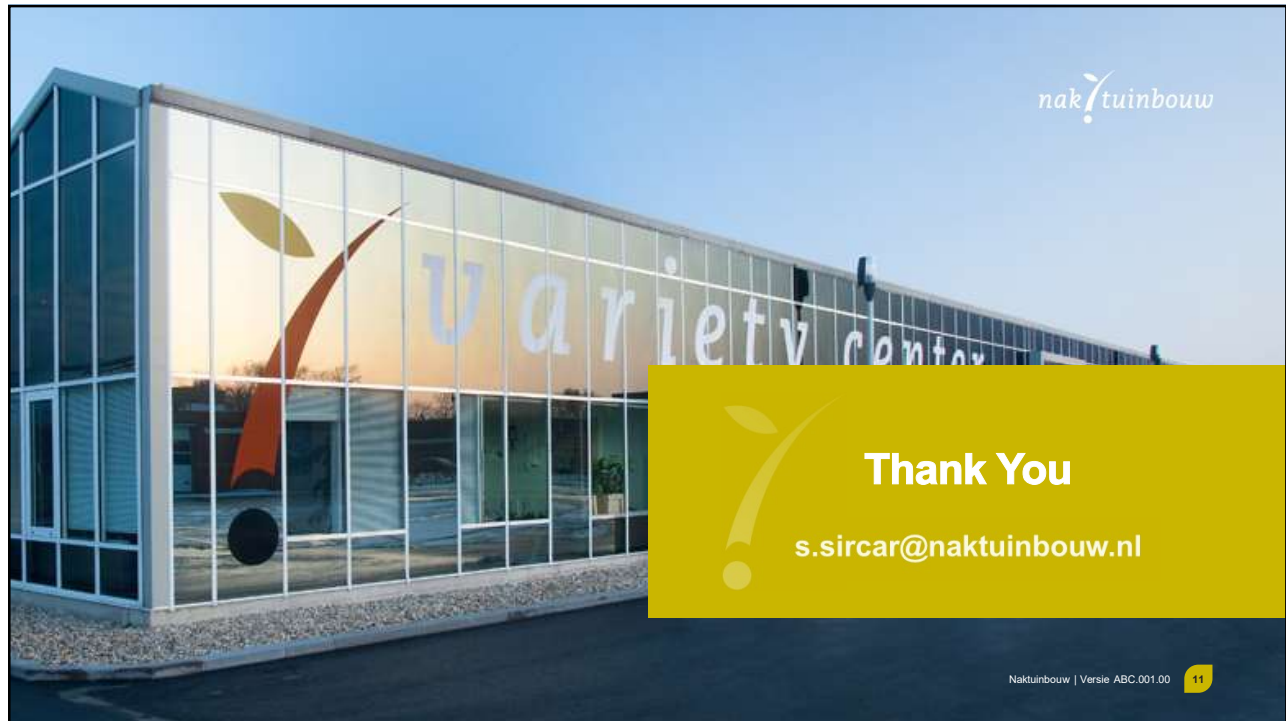
WAGeningen UNIVERSITY & RESEARCH



Registration and coding of floriculture products.
Description and photography of floriculture products

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10



11

[End of Annex and of document]