



International Union for the Protection of New Varieties of Plants

Technical Working Party on Testing Methods and Techniques TWM/1/19

First Session Virtual meeting, September 19 to 23, 2022 Original: English

Date: August 31, 2022

VARIETY TRACER: FRAUDULENT USE OF PARENTAL LINES

Document prepared by an expert from the Netherlands

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The annex to this document contains a copy of a presentation on "Variety Tracer: Fraudulent use of parental lines", prepared by an expert from the Netherlands, to be made at the first session of the TWM.

[Annex follows]



Variety Tracer Fraudulent use of parental lines

With examples in pepper and broccoli

UPOV-TWM/1 – September 19-23, 2022



DNA as a tool to find your parents



DNA Onbekend - Corrie en Gerard: worstelen met vragen

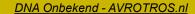


DNA Onbekend: Cyril en Yvonne - wie is mijn vader?





DNA Onbekend: Jolanda en Marijke - wie is mijn vader en wie is mijn opa?





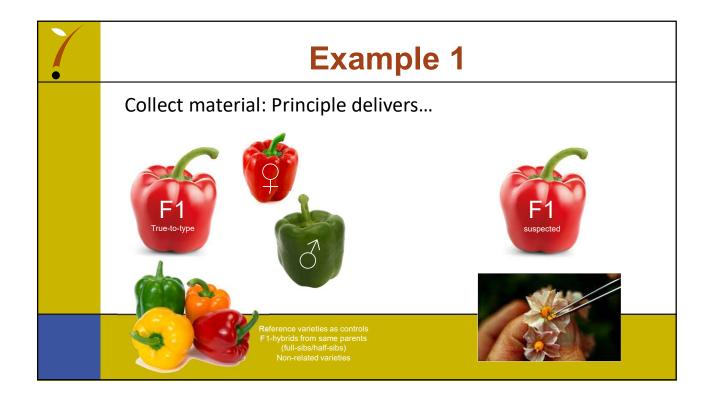
DNA Onbekend - Thea en Pieter op zoek naar de waarheid

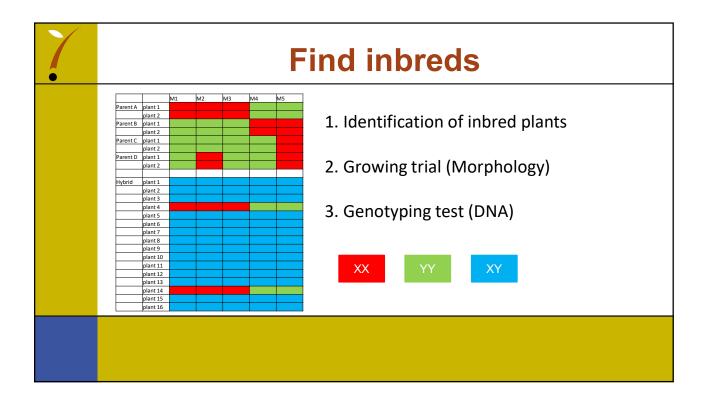
Variety Tracer

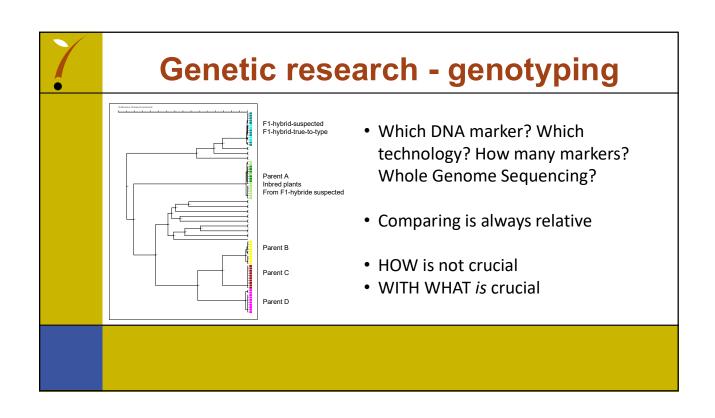
How to proof that a protected parent is (illegally) used to produce a F1-hybrid variety?

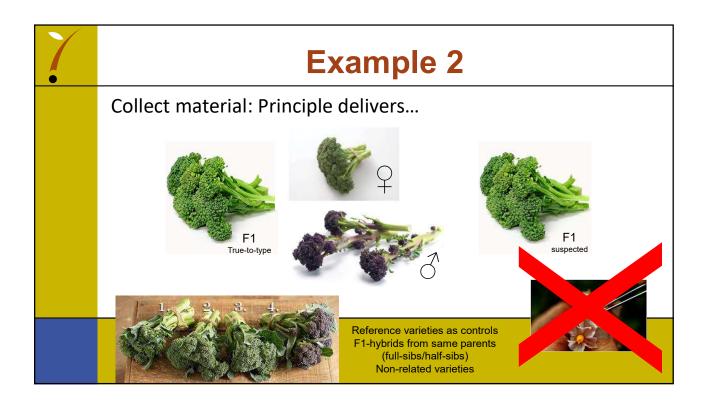
Strategies:

- 1. Find inbred plants in the F1 and compare with potential true-to-type parents.
- 2. Statistical 'parent-offspring' test.
- 3. Multiple Sequence Alignments inheritance of haploblocks.









Statistical 'Parent-Offspring'-test

- Input DNA information: Brassica-specific genotyping 15K array with 13714 SNPs
- Looked at the possible genetic contribution of each individual potential parent to the genotype of each offspring plant (F1-hybrid).

Parent	Offspring	Score
XX	XX	1
XX	XY	1
XX	YY	0
XY	XX	1
XY	XY	1
XY	YY	1
YY	XX	0
YY	XY	1
YY	YY	1

C = b / (b+c)

Where C is the contribution of one potential parent to a single offspring F1 hybrid variety, b is the number of 1s and c is the total number of 0s for all SNPs between the parent and the offspring.

Max C = 0.5

Can this parent have contributed to this F1 offspring plant?



Statistical 'Parent-Offspring'-test

	F1-suspected-sample 1	F1-suspected-sample 2	F1-suspected (harvested material)	F1-true-to-type-principle	F1-true-to-type (DUS material)	х,	٨,	א,	,8.	,כ,	,۵,	. 3.	.2.
	1a	1b	1c	4	5	8	10	14	15	16	17	18	13
	2 0,499	0,499	0,499	0,499	0,499	0,499	0,433	0,419	0,428	0,434	0,432	0,433	0,438
	3 0,475	0,474	0,474	0,499	0,499	0,455	0,469	0,451	0,470	0,473	0,466	0,466	0,439
	9 0,454	0,454	0,454	0,458	0,458	0,499	0,468	0,447	0,461	0,460	0,465	0,466	0,449
1		0,457	0,456	0,459	0,458	0,454	0,499	0,467	0,488	0,471	0,483	0,483	0,450
1		0,460	0,460	0,468	0,468	0,469	0,499	0,456	0,465	0,463	0,472	0,499	0,455

1abc = 2?

Research question



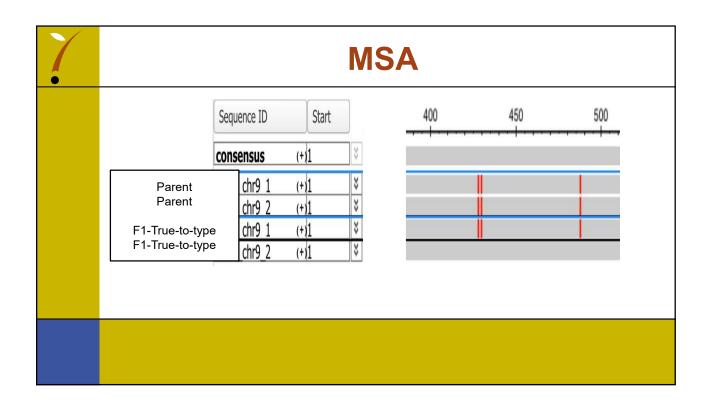
Smoking-Gun Evidence

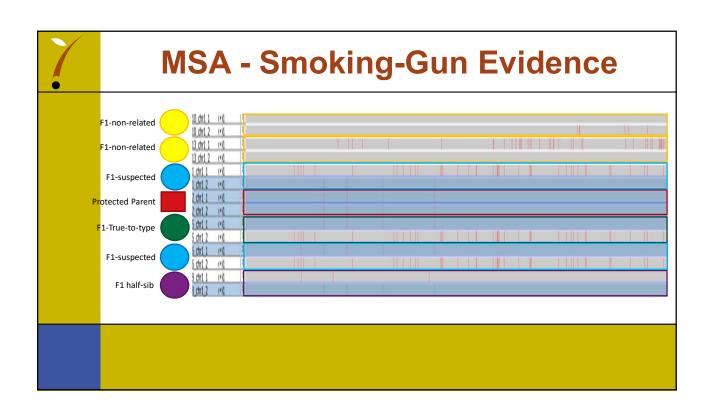
Multiple Sequence Alignments – inheritance of haploblocks (physically linked SNPs) from parent to offspring

What do we need?

- WGS: DNA sequences of long strands of DNA (e.g. nanopore)
- Reliable and high quality reference genome for mapping.
- · Identification of regions with physically linked SNPs
- High coverage and high reliable sequencing quality (Illumina)
- · Highly skilled bioinformaticians with a lot of spare time
- Enough space for data storage









Summary

- There are several strategies to proof that a protected parent has (illegally) been used to produce a F1-hybrid.
- The relative simple and cost-effective strategies need inbred plants and/or produces statistical evidence.
- The innovative and expensive strategy produces 'Smoking-Gun' evidence.





Principles

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