

International Union for the Protection of New Varieties of Plants

**Technical Working Party for Vegetables** 

TWV/54/7

Fifty-Fourth Session Brasilia, Brazil, May 11 to 15, 2020 Original: English

Date: April 23, 2020

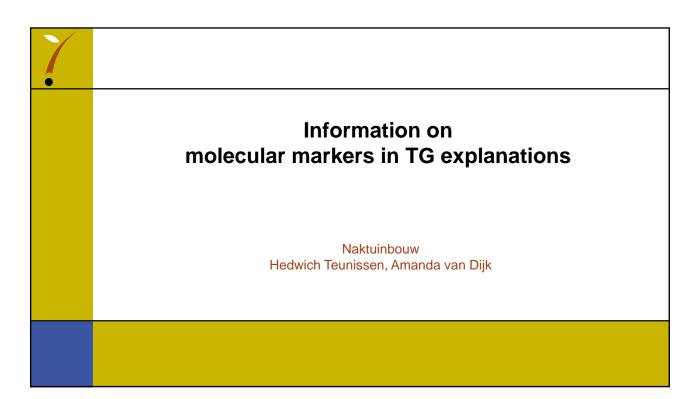
## **USE OF MOLECULAR TECHNIQUES IN DUS EXAMINATION**

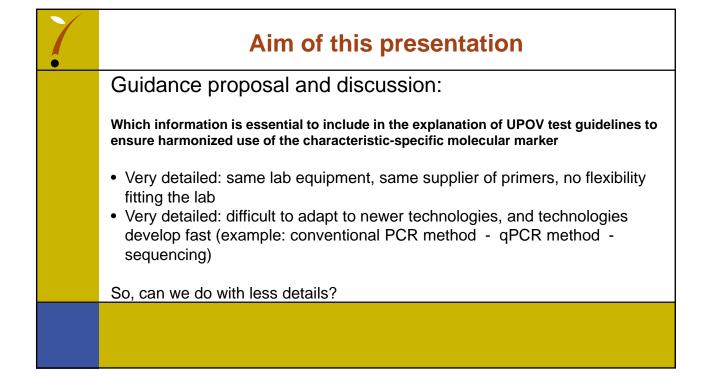
Document prepared by the Office of the Union

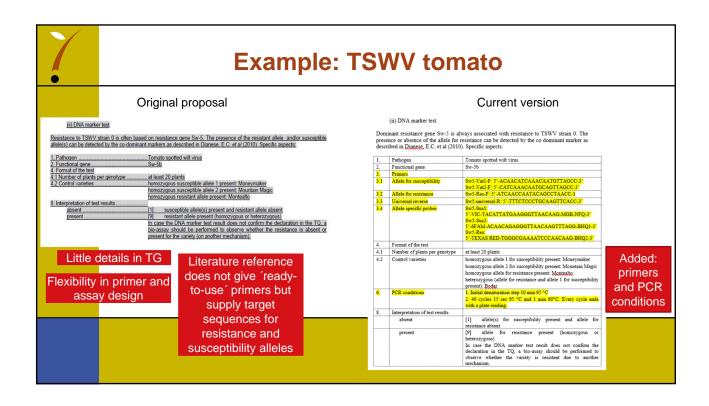
Disclaimer: this document does not represent UPOV policies or guidance

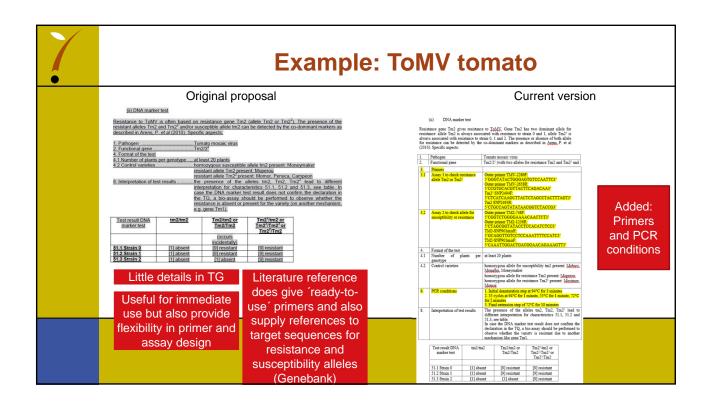
The annex to this document contains a copy of a presentation "Information on molecular markers in Test Guidelines explanations", prepared by an expert from the Netherlands, to be considered by the fifty-fourth session of the Technical Working Party for Vegetables (TWV).

[Annex follows]











## **EU situation vs UPOV**

#### In general

 In EU the UPOV TGs are converted into European Technical Protocols (TPs) wich are mandatory

#### In the tomato situation

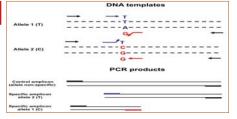
- In parallel with UPOV partial revision of tomato, CPVO revision of technical protocol of tomato.
- Based on the discussions in TWV and VEM the molecular markers for ToMV and TSWV were added to the CPVO TP including all technical details on primers sequences and PCR conditions

CPVO-TP/044/4 Rev.3 Ad. 55 for TSWV resistance; CPVO-TP/044/4 Rev.3 Ad. 48.1-48.3 for ToMV resistance



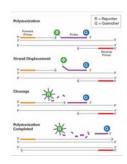
# Improvements of the method to detect the marker





### Improved method:

TaqMan SNP detection Only 1 assay: Using tm2, Tm2 and Tm2<sup>2</sup> specific probes

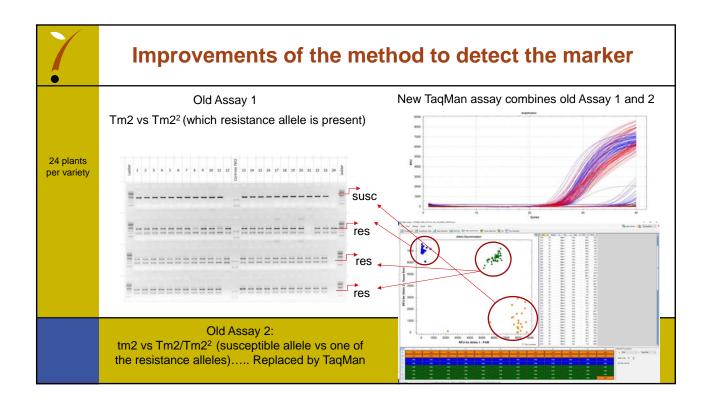


#### Original method:

ARMS-PCR SNP detection

1st assay: Tm2 vs Tm22 (which resistance allele is present)

2<sup>nd</sup> assay: tm2 vs Tm2/Tm2<sup>2</sup> (susceptible allele vs one of the resistance alleles)



	Which information is essential?		
		Essential?	Proposal or remark
1	Pathogen	YES	
2.1	Functional gene	YES	Reference to literature and/or position on ref genome/sequence in public databases
2.2	Functional allele for resistance	YES	Reference to literature and/or position on ref genome/sequence in public databases
2.3	Functional allele for susceptibility	YES	Reference to literature and/or position on ref genome/sequence in public databases
3	Primers	NO	Sequences, reference to accessions and sequences in public databases (Genebank numbers), litter
3.1	Primers to detect allele susceptibility	NO	Sequences corresponding to allele(s) for expression A (susceptibility)
3.2	Primers to detect allele for resistance	NO	Sequences corresponding to allele(s) for expression B (resistance)
4	Format of the test		
4.1	Number of plants	YES	A minimal number of plants required
4.2	Control varieties	YES	Control varieties (same as in bioassay) as standards representing alleles homozygous for suscepti homozygous for resistance, heterozygous.
5	Preparation	NO	Depending on the method used. Not in the TG but detailed protocol(s) as an example in annex or a on request from the institute that developed the marker
6	PCR conditions	NO	Depending on the method used. Not in the TG but detailed protocol(s) as an example in annex or a on request from the institute that developed the marker
7	Observations	NO	Depending on the method used. Not in the TG but detailed protocol(s) as an example in annex or a on request from the institute that developed the marker
8	Interpretation of the test results	YES	Relation between alleles and expressions (with its notes)



[End of Annex and of document]