EUROPEAN AND MEDITERRANEAN PLANT PROTECTION ORGANIZATION ORGANISATION EUROPEENNE ET MEDITERRANEENNE POUR LA PROTECTION DES PLANTES Summary sheet of validation data for a diagnostic test

The EPPO Standard PM 7/98 Specific requirements for laboratories preparing accreditation for a plant pest diagnostic activity describes how validation should be conducted. It also includes definitions of performance criteria.

Laboratory contact details	Netherlands Institute for Vectors, Invasive plants and Plant health P.O. Box 9102, 6700 HC Wageningen, Netherlands	
Short description of the test	Validation of a conventional RT-PCR test for detection and identification of PSTVd, TCDVd, MPVd and TPMVd using primers 2H1/3H1 described by Shamoul et al. (1997)	
Date, reference of the validation report	2013-09-17 - NRC-ref: 2011.molbio.006	
Validation process according to EPPO Standard PM7/98?	yes	
Is the lab accredited for this test?	no	
Was the validated data generated in the framework of a project?		
Description of the test		
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Organism(s)	Tomato chlorotic dwarf viroid / Pospiviroid chloronani (TCDVD0) Potato spindle tuber viroid / Pospiviroid fusituberis (PSTVD0)	
Detection / identification	detection and identification	
Method(s)	Molecular Extraction DNA RNA Molecular Conventional RT PCR Molecular other	
Method: Molecular Extraction DNA RNA		
Reference of the test description		
Kit		
Is a kit used	yes	
Manufacturer name	QIAGEN	
Specify the kit used	RNeasy Plant Mini Kit	
Kit used following the manufacturer's instructions?		
Other information		
Method: Molecular Conventional RT PCR		
Reference of the test description		
As or adapted from an IPPC diagnostic protocol	yes	

IPPC diagnostic Protocol name	ISPM 27 Annex 07 DP 07: Potato spindle tuber viroid (version 2006)	
Name of the test	Conventional RT-PCR using the primers of Shamloul et al. (1997)	
Other information		
Other details on the test	section 3.3.4.1 Conventional RT-PCR using the primers of Shamloul et al. (1997) Shamloul AM, Hadidi A, Zhu SF, Singh RP, Sagredo B (1997) Sensitive detection of potato spindle tuber viroid using RT-PCR and identification of a viroid variant naturally infecting pepino plants. Can. J. Pl. Pathol. 19: 89-96 RT-PCR by 2H1/3H1 primers and sequencing of the amplicon	
Method: Molecular other		
Reference of the test description		
Other information		
Other details on the test	Sequencing	
Are the performance characteristics included in the EPPO diagnostic protocol?	no	
Performance Criteria :		
Organism 1.:	Pospiviroid chloronani(TCDVD0)	
Analytical sensitivity		
What is smallest amount of target that can be detected reliably?	Undiluted leaf sap was considered 100% infected. Starting with undiluted plant sap from PSTVd and TCDVd infected hosts up to 10^7 times dilutes in sap of healthy tomato leaves, all PSTVd and TCDVd strains were detected up to a relative infection rate of 10% (=10 times diluted).	
Diagnostic sensitivity		
Proportion of infected/infested samples tested positive compared to results from the standard test, see appendix 2 of PM 7/98	1	
Standard test(s)	PCR sequencing of the complete viroid genome is considered the standard test. This can be achieved with primers Vid-FW/Vid-RE (CLVd, PSTVd and TCDVd) and 3H1/2H1 (PSTVd, TCDVd, MPVd and TPMVd). For other pospiviroids primers Pospi1-FW/Pospi1-RE are used to sequence the partial viroid genome.	
Analytical specificity - inclusivity		
Number of strains/populations of target organisms tested	7 pospiviroid strains, see table 4 validation report	
Specificity value	1	
Analytical specificity - exclusivity		
Number of non-target organisms tested	10 (4 Avsunviroidae and 6 Pospiviroidae strains), see table 6 validation report	

Specificity value		
<u>Diagnostic Specificity</u>		
Proportion of uninfected/uninfested samples (true negatives) testing negative compared to results from a standard test	1	
Specify the test(s)	PCR sequencing of the complete viroid genome is considered the standard test. This can be achieved with primers Vid-FW/Vid-RE (CLVd, PSTVd and TCDVd) and 3H1/2H1 (PSTVd, TCDVd, MPVd and TPMVd). For other pospiviroids primers Pospi1-FW/Pospi1-RE are used to sequence the partial viroid genome.	
Reproducibility		
Provide the calculated % of agreement for a given level of the pest (see PM 7/98)	100% at undilluted, 2x and 4x diluted	
Repeatability		
Provide the calculated % of agreement for a given level of the pest (see PM 7/98)	100% at undilluted, 2x and 4x diluted	
Organism 2.:	Pospiviroid fusituberis(PSTVD0)	
Analytical sensitivity		
What is smallest amount of target that can be detected reliably?	Undiluted leaf sap was considered 100% infected. Starting with undiluted plant sap from PSTVd and TCDVd infected hosts up to 10^7 times dilutes in sap of healthy tomato leaves, all PSTVd and TCDVd strains were detected up to a relative infection rate of 10% (=10 times diluted).	
Diagnostic sensitivity		
Proportion of infected/infested samples tested positive compared to results from the standard test, see appendix 2 of PM 7/98	1	
Standard test(s)	PCR sequencing of the complete viroid genome is considered the standard test. This can be achieved with primers Vid-FW/Vid-RE (CLVd, PSTVd and TCDVd) and 3H1/2H1 (PSTVd, TCDVd, MPVd and TPMVd). For other pospiviroids primers Pospi1-FW/Pospi1-RE are used to sequence the partial viroid genome.	
Analytical specificity - inclusivity		
Number of strains/populations of target organisms tested	7 pospiviroid strains, see table 4 validation report	
Specificity value	1	
Analytical specificity - exclusivity		
Number of non-target organisms tested	10 (4 Avsunviroidae and 6 Pospiviroidae strains), see table 6 validation report	
Specificity value		
Diagnostic Specificity		

Proportion of uninfected/uninfested samples	1	
(true negatives) testing negative compared to results from a standard test		
Specify the test(s)	PCR sequencing of the complete viroid genome is considered the standard test. This can be achieved with primers Vid-FW/Vid-RE (CLVd, PSTVd and TCDVd) and 3H1/2H1 (PSTVd, TCDVd, MPVd and TPMVd). For other pospiviroids primers Pospi1-FW/Pospi1-RE are used to sequence the partial viroid genome.	
Reproducibility		
Provide the calculated % of agreement for a given level of the pest (see PM 7/98)	100% at undilluted, 2x and 4x diluted	
Repeatability		
Provide the calculated % of agreement for a given level of the pest (see PM 7/98)	100% at undilluted, 2x and 4x diluted	
Test performance study		
Test performance study?	no	
Other information		
Any other information considered useful	To determine the robustness of the test two different reaction mixes were used to generate PCR products: OneStep RT-PCR kit (Qiagen) and SuperScript One-Step RT-PCR System with Platinum Taq (Invitrogen). The latter yielded the best results. Different extraction buffers were used. Using a GH plus buffer with a heating step yielded the best results. Freezing the samples prior to analysis did not influence the qualitative results obtained.	
The following complementary files are available online:	 Validation of a conventional RT-PCR test for detection and identification of PSTVd, TCDVd, MPVd and TPMVd using primers described by Shamoul et al. (1997). (In Dutch) 	

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