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 assay for detection and preliminary identification of pospiviroids (expect CLVd) by Pospi1-FW/Pospi1-RE
Date, reference of the validation report	2013-09-17 - NRC-ref: 2010.molbio.033
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	
Organism(s)	Pospiviroid (1POSPG) Potato spindle tuber viroid / Pospiviroid fusituberis (PSTVD0)
Detection / identification	detection and identification
Matrix(ces) tested	Leaves leaves of Celosia cardiophyllum and Solanum lycopersicum
Plant species tested	Celosia sp., Solanum lycopersicum
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 Verhoeven et al. (2004)
Other information	
Other details on the test	section 3.3.3.3 Conventional RT-PCR using the primers of Verhoeven et al. (2004) J.Th.J. Verhoeven, C.C.C. Jansen, T.M. Willemen, L.F.F. Kox, R.A. Owens and J.W. Roenhorst (2004) Natural infections of tomato by Citrus exocortis viroid, Columnea latent viroid, Potato spindle tuber viroid and Tomato chlorotic dwarf viroid. European Journal of Plant Pathology 110: 823–831 RT-PCR by Pospi1-FW/Pospi1-RE 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(1POSPG)**
Analytical sensitivity	
What is the smallest amount of target that can be detected reliably?	Undiluted infected leaf sap was considered 100% infected. Starting with undiluted plant sap from infected hosts up to 10^7 times dilutes in sap of healthy tomato leaves, all species were detected up to a relative infection rate of 2,5 % (=40 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	12 pospiviroid strains, see table 4 in validation report
Specificity value	1

Analytical specificity - exclusivity	
Number of non-target organisms tested	10 (4 Avsunviroidae and 6 Pospiviroidae), see table 8 in validation report
Specificity value	
Diagnostic Specificity	
Proportion of uninfected/uninfested samples (true negatives) testing negative compared to results from a standard test	100 %
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 undiluted, 2x, 4x, 10x and 100 diluted
Repeatability	
Provide the calculated % of agreement for a given level of the pest (see PM 7/98)	100% at undiluted, 2x, 4x, 10x and 100 diluted
Organism 2.:	**Pospiviroid fusituberis(PSTVD0)**
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 assay for detection and preliminary identification of pospiviroids (expect CLVd) by Pospi1-FW/RE. (In Dutch)

Creation date: 2014-05-28 00:00:00 - Last update: 2021-05-10 17:13:19