

**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.

<b>Laboratory contact details</b>	Netherlands Institute for Vectors, Invasive plants and Plant health P.O. Box 9102, 6700 HC Wageningen, Netherlands
<b>Short description of the test</b>	Duplex-PCR tests for the identification <i>X. euvesicatoria</i> , <i>X. vesicatoria</i> , <i>X. gardneri</i> and <i>X. perforans</i> .
<b>Date, reference of the validation report</b>	2012-04-17 - MOVA nummer: 2011.molbio.004
<b>Validation process according to EPPO Standard PM7/98?</b>	yes
<b>Is the lab accredited for this test?</b>	no
<b>Was the validated data generated in the framework of a project?</b>	
<b>Description of the test</b>	
<b>Organism(s)</b>	<i>Xanthomonas hortorum</i> pv. <i>gardneri</i> (XANTGA) <i>Xanthomonas euvesicatoria</i> pv. <i>perforans</i> (XANTPF) <i>Xanthomonas vesicatoria</i> (XANTVE) <i>Xanthomonas euvesicatoria</i> pv. <i>euvesicatoria</i> (XANTEU)
<b>Detection / identification</b>	identification
<b>Method(s)</b>	Molecular Conventional PCR
<b>Method: Molecular Conventional PCR</b>	
<b>Reference of the test description</b>	
<b>As or adapted from an EPPO diagnostic protocol</b>	yes
<b>EPPO Diagnostic Protocol name</b>	PM 7/110 <i>Xanthomonas</i> spp. ( <i>Xanthomonas euvesicatoria</i> , <i>Xanthomonas gardneri</i> , <i>Xanthomonas perforans</i> , <i>Xanthomonas vesicatoria</i> ) causing bacterial spot of tomato and sweet pepper (version 1)
<b>Name of the test</b>	Conventional PCR (according to Koenraadt et al., 2009)
<b>Other information</b>	
<b>Reaction type</b>	Duplex
<b>Other details on the test</b>	Two conventional duplex-PCR tests
<b>Performance Criteria :</b>	

<b>Organism 1.:</b>	<b>Xanthomonas hortorum pv. gardneri(XANTGA)</b>
<b>Analytical sensitivity</b>	
<b>What is smallest amount of target that can be detected reliably?</b>	For the X. gardneri it is found to be $2,5 \times 10^6$ cfu/ml
<b>Analytical specificity - inclusivity</b>	
<b>Number of strains/populations of target organisms tested</b>	53 isolates of Xcv
<b>Specificity value</b>	
<b>Analytical specificity - exclusivity</b>	
<b>Number of non-target organisms tested</b>	6 non-target organisms (in total 21 isolates): Clavibacter michiganensis subsp. michiganensis, Ralstonia solanacearum, Pseudomonas syringae pv. tomato, Pseudomonas syringae pv. syringae, Agrobacterium tumefaciens, Pseudomonas corrugata
<b>Specificity value</b>	Five of the 21 related isolates gave weak (non-specific) amplicon(s). It refers to 2 P.syringae pv tomato isolates and 3 Pseudomonas corrugata isolates. This underlines the risk for wrong identifications for Xg or Xp.
<b>Cross reacts with</b>	Pseudomonas syringae pv. tomato Pseudomonas corrugata
<b>Reproducibility</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	100%
<b>Repeatability</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	93%
<b>Organism 2.:</b>	<b>Xanthomonas euvesicatoria pv. perforans(XANTPF)</b>
<b>Analytical sensitivity</b>	
<b>What is smallest amount of target that can be detected reliably?</b>	for X. perforans $1,9 \times 10^7$ cfu/ml
<b>Analytical specificity - inclusivity</b>	
<b>Number of strains/populations of target organisms tested</b>	53 isolates of Xcv
<b>Specificity value</b>	
<b>Analytical specificity - exclusivity</b>	
<b>Number of non-target organisms tested</b>	6 non-target organisms (in total 21 isolates): Clavibacter michiganensis subsp. michiganensis, Ralstonia solanacearum, Pseudomonas syringae pv. tomato, Pseudomonas syringae pv. syringae, Agrobacterium tumefaciens, Pseudomonas corrugata

<b>Specificity value</b>	Five of the 21 related isolates gave weak (non-specific) amplicon(s). It refers to 2 P.syringae pv tomato isolates and 3 Pseudomonas corrugata isolates. This underlines the risk for wrong identifications for Xg or Xp.
<b>Cross reacts with</b>	Pseudomonas syringae pv. tomato Pseudomonas corrugata
<b>Reproducibility</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	100%
<b>Repeatability</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	93%
<b>Organism 3.:</b>	<b>Xanthomonas vesicatoria(XANTVE)</b>
<b>Analytical sensitivity</b>	
<b>What is smallest amount of target that can be detected reliably?</b>	for X. vesicatoria: 1,6x10 <sup>6</sup> cfu/ml.
<b>Analytical specificity - inclusivity</b>	
<b>Number of strains/populations of target organisms tested</b>	53 isolates of Xcv
<b>Specificity value</b>	
<b>Analytical specificity - exclusivity</b>	
<b>Number of non-target organisms tested</b>	6 non-target organisms (in total 21 isolates): Clavibacter michiganensis subsp. michiganensis, Ralstonia solanacearum, Pseudomonas syringae pv. tomato, Pseudomonas syringae pv. syringae, Agrobacterium tumefaciens, Pseudomonas corrugata
<b>Specificity value</b>	Five of the 21 related isolates gave weak (non-specific) amplicon(s). It refers to 2 P.syringae pv tomato isolates and 3 Pseudomonas corrugata isolates. This underlines the risk for wrong identifications for Xg or Xp.
<b>Cross reacts with</b>	Pseudomonas syringae pv. tomato Pseudomonas corrugata
<b>Reproducibility</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	100%
<b>Repeatability</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	93%
<b>Organism 4.:</b>	<b>Xanthomonas euvesicatoria pv. euvesicatoria(XANTEU)</b>
<b>Analytical sensitivity</b>	

<b>What is smallest amount of target that can be detected reliably?</b>	for <i>X. euvesicatoria</i> $5,5 \times 10^5$ cfu/ml
<b>Analytical specificity - inclusivity</b>	
<b>Number of strains/populations of target organisms tested</b>	53 isolates of Xcv
<b>Specificity value</b>	
<b>Analytical specificity - exclusivity</b>	
<b>Number of non-target organisms tested</b>	6 non-target organisms (in total 21 isolates): <i>Clavibacter michiganensis</i> subsp. <i>michiganensis</i> , <i>Ralstonia solanacearum</i> , <i>Pseudomonas syringae</i> pv. <i>tomato</i> , <i>Pseudomonas syringae</i> pv. <i>syringae</i> , <i>Agrobacterium tumefaciens</i> , <i>Pseudomonas corrugata</i>
<b>Specificity value</b>	Five of the 21 related isolates gave weak (non-specific) amplicon(s). It refers to 2 <i>P.syringae</i> pv <i>tomato</i> isolates and 3 <i>Pseudomonas corrugata</i> isolates. This underlines the risk for wrong identifications for Xg or Xp.
<b>Cross reacts with</b>	<i>Pseudomonas syringae</i> pv. <i>tomato</i> <i>Pseudomonas corrugata</i>
<b>Reproducibility</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	100%
<b>Repeatability</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	93%
<b>Test performance study</b>	
<b>Test performance study?</b>	no
<b>Other information</b>	
<b>Any other information considered useful</b>	This PCR test has been found robust for variations in the DNA extraction method. Three different DNA extraction methodologies have been performed and scored equally well: 1. QuickPick Plant DNA Kit (Bio-Nobile, KingFisher method) 2. High Pure PCR Template preparation kit (Roche) 3. Boiling method
The following complementary files are available online:	
	<ul style="list-style-type: none"> <li>• <a href="#">Validation report 2011.molbio.004</a></li> </ul>

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