

**EUROPEAN AND MEDITERRANEAN PLANT PROTECTION ORGANIZATION**  
**ORGANISATION EUROPEENNE ET MEDITERRANEENNE POUR LA PROTECTION DES PLANTES**  
(11-17239)

**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>Target Organism</b>	Bursaphelenchus xylophilus	
<b>Short description</b>	detection of Bursaphelenchus xylophilus by real-time PCR	
<b>Laboratory contact details</b>	National Reference Centre, National Plant Protection Organization P.O. Box 9102, 6700 HC Wageningen, Netherlands	
<b>Date and reference of the validation report</b>	2014-07-21 - F-315-000-004	
<b>Validation process according to EPPO Standard PM 7/98:</b>	Yes	
<b>Reference of the test description</b>	PM 7/004	
<b>Is the test the same as described in the EPPO DP?</b>	Modified Use of SYBR green real-time PCR from ClearDetections	
<b>Is the lab accredited for this test?</b>	No	
<b>Plant species tested (if relevant)</b>		
<b>Matrices tested (if relevant)</b>	nematode suspension from pine bark	
<b>List of methods used</b>		
<b>Method for extraction / isolation / baiting of target organism from matrix</b>		
<b>Molecular methods, e.g. hybridization, PCR and real time PCR</b>	X	Three different isolation methods were first compared before the lab validation was carried out further. The three methods tested from the EPPO protocol were the QIAamp DNA mini kit Qiagen, FES protocol Agowa sbeadex plant mini kit using the KingFisher and ClearDetections Nematode DNA extraction & purification kit. Taqman real-time PCR from the EPPO protocol PM7/4(3) and SYBR green real-time PCR from ClearDetections
<b>Serological methods: IF, ELISA, Direct Tissue Blot Immuno Assay</b>		
<b>Plating methods: selective isolation</b>		
<b>Bioassay methods: selective enrichment in host plants, baiting,</b>		

<b>plant test and grafting.</b>		
<b>Pathogenicity test</b>		
<b>Fingerprint methods: protein profiling, fatty acid profiling &amp; DNA profiling</b>		
<b>Morphological and morphometrical methods intended for identification</b>		
<b>Biochemical methods: e.g. enzyme electrophoresis, protein profiling</b>		
<b>Other</b>		
<b>Analytical sensitivity (= limit of detection)</b>		
<b>What is smallest amount of target that can be detected reliably?</b>	The detection limit for the Taqman real-time PCR, for the Sbeadex extraction method, is 10 <i>B. xylophilus</i> per 2 ml of nematode suspension from wood (St=0). The detection limit for SYBR green real-time PCR is 1 <i>B. xylophilus</i> per 2 mL of nematode suspension from wood for both the Sbeadex and the ClearDetections extraction method.	
<b>Diagnostic sensitivity</b>		
<b>Proportion of infected/infested samples tested positive compared to results from the standard test , see appendix 2 of PM 7/98</b>	Not evaluated	
<b>Specify the standard test</b>		
<b>Analytical specificity</b>		
<b>Specificity value</b>		
<b>Number of strains/populations of target organisms tested</b>	4 populations of target organisms were tested: FR-04-415-1 <i>Bursaphelenchus xylophilus</i> Frankrijk, collectie LNPV <i>Botrytis cinerea</i> E4757-1 <i>B. xylophilus</i> Canada Hout, Pinus E4757-2 <i>B. xylophilus</i> Canada Hout, Pinus F0426 <i>B. xylophilus</i> Portugal verpakkingshout	
<b>Number of non-target organisms tested</b>	10 F0881 <i>B. mucronatus</i> China emballagehout E9674-1 <i>B. mucronatus</i> Rusland Larix F0335-1 <i>B. fungivorus</i> Portugal Schors van Pinus F0444 <i>B. minutus</i> Portugal Schors van Pinus E9674-2 <i>B. mucronatus</i> China meubels F0228 <i>A. blastophtorus</i> Nederland Crocus E9805 <i>A. saprophilus</i> Nederland Pinus E9597 <i>A. subtenuis</i> Nederland Crocus E9192 <i>A. besseyi</i> Turkije <i>Oryza sativa</i> E7072-2 <i>Aphelenchoides</i> sp. Portugal verpakkingshout	
<b>Cross reacts with (specify the species)</b>	These 10 non-targets were negative in the Tagman real-time PCR. In the SYBER green real-time PCR, 1 of the 4 biological samples from <i>Bursaphelenchus minutus</i> (F0444) and 1 of the four biological samples from <i>Bursaphelenchus</i> sp. F0881 gave a false positive result. It is possible that the samples are contaminated, but it is no longer possible to determine where this happened. The reliability of the samples is called into	

	question and not the final assessment.
<b>Diagnostic Specificity</b>	
<b>Proportion of uninfected/uninfested samples (true negatives) testing negative compared to results from a standard test</b>	
<b>Specify the standard test</b>	
<b>Reproducibility</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	
<b>Repeatability</b>	
<b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>	
<b>Test performance study</b>	
<b>Test performance study?</b>	No
<b>Include brief details of the test performance study and its output. If available, provide a link to published article/report</b>	
<b>Other information</b>	
<b>Any other information considered useful e.g. robustness, ease of performing the test, etc.</b>	<p>The results show that the Sbeadex and ClearDetections extraction methods perform better than those of QiaAmp extraction methods. The differences between the Sbeadex and ClearDetections extraction methods are small.</p> <p>No additional wood species have been included for selectivity.</p>
<b>The following complementary files are available online:</b>	<ul style="list-style-type: none"> <li>• <a href="#">Short validation and implementation of real-time PCR detection B xylophilus in nematode suspension of wood bark final</a></li> </ul>