

**EUROPEAN AND MEDITERRANEAN PLANT PROTECTION ORGANIZATION**  
**ORGANISATION EUROPÉENNE ET MEDITERRANÉENNE 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>	EUPH DNA Barcoding , , EUPHRESCO
<b>Short description of the test</b>	DNA Barcoding - Optimizing and validating DNA barcoding protocols for arthropods
<b>Date, reference of the validation report</b>	2016-06-30 - Final_report_DNA barcoding.doc
<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>	Euphresco
<b>If yes, please specify</b>	EUPHRESCO DNA Barcoding - Optimizing and validating DNA barcoding protocols for plant pests
<b>Description of the test</b>	
<b>Organism(s)</b>	Anoplophora glabripennis (ANOLGL) Bemisia tabaci (BEMITA) Vespa crabro (VESPCC) Spodoptera eridania (PRODER) Liriomyza huidobrensis (LIRIHU)
<b>Detection / identification</b>	identification
<b>Method(s)</b>	Molecular Sanger seq Molecular other
<b>Method: Molecular Sanger seq</b>	
<b>Reference of the test description</b>	
<b>Other information</b>	
<b>Method: Molecular other</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 7/129 DNA barcoding as an identification tool for a number of regulated pests (version 1)
<b>As or adapted from an IPPC diagnostic protocol</b>	no
<b>Is the test modified compared to the reference test</b>	no

<b>Kit</b>	
<b>Is a kit used</b>	no
<b>Other information</b>	
<b>Other details on the test</b>	BIO-X-ACT™ Short Mix (Bioline)
<b>Are the performance characteristics included in the EPPO diagnostic protocol?</b>	<b>yes</b>
<b>Performance Criteria :</b>	
<b>Organism 1.:</b>	<b>Anoplophora glabripennis(ANOLGL)</b>
<b>Analytical sensitivity</b>	
<b>What is smallest amount of target that can be detected reliably?</b>	Genomic DNA extract dilutions tested ranged from 0.3 ng/µl to 3.9 ng/µl for the arthropod samples and resulted in amplicons, high quality trace data and identifiable consensus sequences.
<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>	100%
<b>Organism 2.:</b>	<b>Bemisia tabaci(BEMITA)</b>
<b>Analytical sensitivity</b>	
<b>What is smallest amount of target that can be detected reliably?</b>	Genomic DNA extract dilutions tested ranged from 0.3 ng/µl to 3.9 ng/µl for the arthropod samples and resulted in amplicons, high quality trace data and identifiable consensus sequences.
<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>	100%
<b>Organism 3.:</b>	<b>Vespa crabro(VESPCC)</b>
<b>Analytical sensitivity</b>	
<b>What is smallest amount of target that can be detected reliably?</b>	Genomic DNA extract dilutions tested ranged from 0.3 ng/µl to 3.9 ng/µl for the arthropod samples and resulted in amplicons, high quality trace data and identifiable consensus sequences.
<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>	100%
<b>Organism 4.:</b>	<b>Spodoptera eridania(PRODER)</b>
<b>Analytical sensitivity</b>	
<b>What is smallest amount of target that can be detected reliably?</b>	Genomic DNA extract dilutions tested ranged from 0.3 ng/µl to 3.9 ng/µl for the arthropod samples and resulted in amplicons, high quality trace data and identifiable consensus sequences.
<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>	91% (100% after re-analysing the consensus sequence data provided by TPS participants)
<b>Organism 5.:</b>	<b>Liriomyza huidobrensis(LIRIHU)</b>
<b>Analytical sensitivity</b>	
<b>What is smallest amount of target that can be detected reliably?</b>	Genomic DNA extract dilutions tested ranged from 0.3 ng/µl to 3.9 ng/µl for the arthropod samples and resulted in amplicons, high quality trace data and identifiable consensus sequences.
<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>	100%
<b>Test performance study</b>	
<b>Test performance study?</b>	yes
<b>Brief details of the test performance study and its output. If available, link to published article/report</b>	Test performance study involving 13 laboratories (11 of which provided data). See report for more information.
The following complementary files are available online:	
<ul style="list-style-type: none"> <li>• <a href="#">EUPHRESCO DNA Barcoding - report</a></li> </ul>	

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