

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.

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| Laboratory contact details | EUPH DNA Barcoding , , EUPHRESCO |
| Short description of the test | DNA Barcoding - Optimizing and validating DNA barcoding protocols for arthropods |
| Date, reference of the validation report | 2016-06-30 - Final_report_DNA barcoding.doc |
| 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? | Euphresco |
| If yes, please specify | EUPHRESCO DNA Barcoding - Optimizing and validating DNA barcoding protocols for plant pests |
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| Description of the test | |
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| Organism(s) | Anoplophora glabripennis (ANOLGL) Bemisia tabaci (BEMITA) Vespa crabro (VESPCC) Spodoptera eridania (PRODER) Liriomyza huidobrensis (LIRIHU) |
| Detection / identification | identification |
| Method(s) | Molecular other |
| Method: Molecular other | |
| Reference of the test description | |
| As or adapted from an EPPO diagnostic protocol | yes |
| EPPO Diagnostic Protocol name | PM 7/129 DNA barcoding as an identification tool for a number of regulated pests (version 1) |
| As or adapted from an IPPC diagnostic protocol | no |
| Is the test modified compared to the reference test | no |
| Kit | |
| Is a kit used | no |
| Other information | |

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| Other details on the test | BIO-X-ACT™ Short Mix (Bioline) |
| Are the performance characteristics included in the EPPO diagnostic protocol? | yes |
| Performance Criteria : | |
| Organism 1.: | Anoplophora glabripennis(ANOLGL) |
| <u>Analytical sensitivity</u> | |
| What is smallest amount of target that can be detected reliably? | 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. |
| <u>Diagnostic sensitivity</u> | |
| Proportion of infected/infested samples tested positive compared to results from the standard test, see appendix 2 of PM 7/98 | 100% |
| Organism 2.: | Bemisia tabaci(BEMITA) |
| <u>Analytical sensitivity</u> | |
| What is smallest amount of target that can be detected reliably? | 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. |
| <u>Diagnostic sensitivity</u> | |
| Proportion of infected/infested samples tested positive compared to results from the standard test, see appendix 2 of PM 7/98 | 100% |
| Organism 3.: | Vespa crabro(VESPCC) |
| <u>Analytical sensitivity</u> | |
| What is smallest amount of target that can be detected reliably? | 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. |
| <u>Diagnostic sensitivity</u> | |
| Proportion of infected/infested samples tested positive compared to results from the standard test, see appendix 2 of PM 7/98 | 100% |
| Organism 4.: | Spodoptera eridania(RODER) |
| <u>Analytical sensitivity</u> | |
| What is smallest amount of target that can be detected reliably? | 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. |
| <u>Diagnostic sensitivity</u> | |
| Proportion of infected/infested samples tested positive compared to results from the standard test, see appendix 2 of PM 7/98 | 91% (100% after re-analysing the consensus sequence data provided by TPS participants) |
| Organism 5.: | Liriomyza huidobrensis(LIRIHU) |

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| Analytical sensitivity | |
| What is smallest amount of target that can be detected reliably? | 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. |
| Diagnostic sensitivity | |
| Proportion of infected/infested samples tested positive compared to results from the standard test, see appendix 2 of PM 7/98 | 100% |
| Test performance study | |
| Test performance study? | yes |
| Brief details of the test performance study and its output. It available, link to published article/report | Test performance study involving 13 laboratories (11 of which provided data). See report for more information. |
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| The following complementary files are available online: | <ul style="list-style-type: none"> • EUPHRESCO DNA Barcoding - report |

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