

**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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| <b>Laboratory contact details</b>                                      | Institute for Sustainable Plant Protection<br>via Amendola, 122/D, 70126 Bari, Italy                                |
| <b>Short description of the test</b>                                   | Detection of <i>Xylella fastidiosa</i> in perennial host species by Real time PCR (Harper et al. 2010 erratum 2013) |
| <b>Date, reference of the validation report</b>                        | 2015-10-22 - internal report  |
| <b>Validation process according to EPPO Standard PM7/98?</b>           | no  |
| <b>Is the lab accredited for this test?</b>                            | yes   |
| <b>Was the validated data generated in the framework of a project?</b> |   |
| <b>Description of the test</b>   |   |
| <b>Organism(s)</b>   | <i>Xylella fastidiosa</i> (XYLEFA)  |
| <b>Detection / identification</b>                                      | detection   |
| <b>Method(s)</b>   | Molecular Extraction DNA RNA<br>Molecular real time PCR   |
| <b>Method: Molecular Extraction DNA RNA</b>                            |   |
| <b>Reference of the test description</b>                               |   |
| <b>Kit</b>   |   |
| <b>Is a kit used</b>   | yes   |
| <b>Manufacturer name</b>   | QIAGEN  |
| <b>Specify the kit used</b>  | DNeasy mericon Food Kit   |
| Kit used following the manufacturer's instructions?                    |   |
| <b>Other information</b>   |   |
| <b>Other details on the test</b>                                       | "Dneasy mericon food kit" (QIAGEN) for total DNA extraction   |
| <b>Method: Molecular real time PCR</b>                                 |   |
| <b>Reference of the test description</b>                               |   |
| <b>Other information</b>   |   |
| <b>Reaction type</b>   | Probe   |
| <b>Other details on the test</b>                                       | Real time PCR with Taqman probe Harper S.J., Ward L.I., Clover G.R.G., 2010. Development of LAMP and                |

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|  | real-time PCR methods for the rapid detection of <i>Xylella fastidiosa</i> for quarantine and field applications. <i>Phytopathology</i> 100: 1282-1288.  |
| <b>Are the performance characteristics included in the EPPO diagnostic protocol?</b>   | <b>no</b>  |
| <b>Performance Criteria :</b>  |  |
| <b>Organism 1.:</b>  | <b><i>Xylella fastidiosa</i>(XYLEFA)</b>   |
| <b><u>Analytical sensitivity</u></b>   |  |
| <b>What is smallest amount of target that can be detected reliably?</b>  | up to 10 <sup>2</sup> cfu/ml (corresponding to 7 cfu/reaction) using dilutions ranging from 10 <sup>7</sup> to 10 CFU/ml, prepared by spiking the inactivated bacterial culture in total nucleic acids recovered from olive reference sources known to be not infected by <i>Xylella fastidiosa</i> .  |
| <b><u>Diagnostic sensitivity</u></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>Standard test(s)</b>  | 26 obtained positive samples/ 26 expected positive samples   |
| <b><u>Diagnostic Specificity</u></b>   |  |
| <b>Proportion of uninfected/uninfested samples (true negatives) testing negative compared to results from a standard test</b>        | 100%   |
| <b>Specify the test(s)</b>   | 10 obtained negative samples/ 10 expected negative samples   |
| <b><u>Repeatability</u></b>  |  |
| <b>Provide the calculated % of agreement for a given level of the pest (see PM 7/98)</b>   | 100%   |
| <b>Test performance study</b>  |  |
| <b>Test performance study?</b>   | no   |
| <b>Other information</b>   |  |
| <b>Any other information considered useful</b>   | This protocol is designed for the extraction of total DNA from a large-scale sample of raw or processed food material. The protocol can be performed manually or automated using a dedicated workstation starting from 0,5-0,8 g of fresh small pieces of midribs and petioles into extraction bags and homogenized adding 5ml of Food Lysis Buffer, using available equipments (Polytron, Homex, etc); 1 ml of sap is incubate for 30 min at 60°C and after on ice for several minutes, then centrifuged for 5 min at 2500 x g. From this step, total nucleic acids are purified following the manufacturer's instructions (Qiagen) and eluted in a final volume of 100 µl. |