



2004-009: Draft Annex to ISPM 27:2006 – *Erwinia amylovora* (Burrill)

| Comm no. | Para no. | Comment type | Comment | Explanation | Country |
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| 1. | G | Substantive | I support the document as it is and I have no comments | | Singapore, Lao People's Democratic Republic, Georgia, Jamaica, Thailand, United States of America, New Zealand, Barbados, Dominica, Mexico, Canada, Ghana, Korea, Republic of, OIRSA, Malawi, Burundi, Belize, Gabon, Australia |
| 2. | G | Substantive | | Insert some pictures of disease symptoms on typical host plants. These pictures would be useful to identify <i>Erwinia amylovora</i> . | Japan |
| 3. | 5 | Editorial | 1. Pest Information | The number of the section is missing. | EPPO, European Union, Georgia, Serbia |
| 4. | 5 | Editorial | 1. Pest Information | Editorial correction | Japan |
| 5. | 6 | Editorial | <i>Erwinia amylovora</i> is the causal agent of fire blight disease, which affects most species of the subfamily Maloideae of the family Rosaceae (Spiraeoideae). It was the first bacterium described as the causal agent of a plant disease (Burrill, 1883). <i>E. amylovora</i> is considered to be native to North America and was first detected outside North America in New Zealand in 1920. Fire blight was reported in England in 1957 and since then the bacterium has been detected in most areas of Europe where susceptible hosts are cultivated. <i>E. amylovora</i> is now present in more than 40 countries. It has not been recorded in South America and most African and Asian | Cf. paragraphs [8] and [121] (second reference). | EPPO, European Union, Georgia, Serbia |

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| | | | <p>countries (with the exception of countries surrounding the Mediterranean Sea), and it has been eradicated in Australia after one report there (van der Zwet, 2004). It represents a threat to the pome fruit industry of all these countries (Bonn and van der Zwet, 2000). Details on geographic distribution can be found in the EPPO Plant Quarantine Data Retrieval System (EPPO, 2012³).</p> <p>The most important host plants from both economic and epidemiological viewpoints are in the genera <i>Chaenomeles</i>, <i>Cotoneaster</i>, <i>Crataegus</i>, <i>Cydonia</i>, <i>Eriobotrya</i>, <i>Malus</i>, <i>Mespilus</i>, <i>Pyracantha</i>, <i>Pyrus</i>, <i>Sorbus</i> and <i>Stranvaesia</i> (Bradbury, 1986). The <i>E. amylovora</i> strains isolated from <i>Rubus</i> sp. in the United States is distinct from the strains on other hosts (Powney <i>et al.</i>, 2011b; Starr <i>et al.</i>, 1951).</p> <p>Fire blight is probably the most serious bacterial disease affecting <i>Pyrus communis</i> (pear) and <i>Malus domestica</i> (apple) cultivars in many countries. Epidemics are sporadic and are dependent on a number of factors, including favourable environmental conditions, sufficient inoculum level present in the orchard, and host susceptibility. The development of fire blight symptoms follows the seasonal growth development of the host plant. The disease begins in spring with the production of the primary inoculum from bacteria overwintering in cankers (Thomson, 2000) causing blossom infection, continuing into summer with shoot and fruit infection, and ending in winter with the development of cankers throughout the dormant period of the host (Thomson, 2000; van der Zwet and Beer, 1995).</p> | | |
| 6. | 6 | Substantive | <p><i>Erwinia amylovora</i> is the causal agent of fire blight disease, which affects most species of the subfamily Maloideae of the family Rosaceae (Spiraeoideae). It was the first bacterium described as the causal agent of a plant disease (Burrill, 1883). <i>E. amylovora</i> is considered to be native to North America and was first detected outside North America in New Zealand in 1920. Fire blight was reported in England in 1957 and since then the bacterium has been detected in most areas of Europe where susceptible hosts are cultivated. <i>E. amylovora</i> is now present in more than 40 countries. It has not been recorded in South America and most African and Asian countries (with the exception of countries surrounding the Mediterranean Sea), and it has been eradicated in Australia after one report there (van der Zwet, 2004). It represents a threat to the pome fruit industry of all these countries (Bonn and van der Zwet, 2000). Details on geographic distribution can be found in the EPPO Plant Quarantine Data Retrieval System (EPPO, 2012).</p> <p>The most important host plants from both economic and epidemiological viewpoints are in the genera <i>Chaenomeles</i>, <i>Cotoneaster</i>, <i>Crataegus</i>, <i>Cydonia</i>, <i>Eriobotrya</i>,</p> | According to the data sheet on <i>E. amylovora</i> in the Crop Protection Compendium, <i>E. amylovora</i> is easily dispersed by several agents (e.g., rain, wind, insects, birds, etc.) and widely imparts serious damage to nursery fields. | Japan |

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| | | | <p><i>Malus</i>, <i>Mespilus</i>, <i>Pyracantha</i>, <i>Pyrus</i>, <i>Sorbus</i> and <i>Stranvaesia</i> (Bradbury, 1986). The <i>E. amylovora</i> strains isolated from <i>Rubus</i> sp. in the United States is distinct from the strains on other hosts (Powney <i>et al.</i>, 2011b; Starr <i>et al.</i>, 1951).</p> <p>Fire blight is probably the most serious bacterial disease affecting <i>Pyrus communis</i> (pear) and <i>Malus domestica</i> (apple) cultivars in many countries. Epidemics are sporadic and are dependent on a number of factors, including favourable environmental conditions, sufficient inoculum level present in the orchard, and host susceptibility. The development of fire blight symptoms follows the seasonal growth development of the host plant. The disease begins in spring with the production of the primary inoculum from bacteria overwintering in cankers (Thomson, 2000) causing blossom infection, continuing into summer with shoot and fruit infection, and ending in winter with the development of cankers throughout the dormant period of the host (Thomson, 2000; van der Zwet and Beer, 1995). In this infection cycle, the disease is easily dispersed by birds, insects, rain or wind in this(Thomson, 2000).</p> | | |
| 7. | 7 | Editorial | 2. Taxonomic Information | The number of the section is missing. | EPPO, European Union, Georgia, Serbia |
| 8. | 7 | Editorial | 2. Taxonomic Information | Editorial correction | Japan |
| 9. | 8 | Editorial | <p>Name: <i>Erwinia amylovora</i> (Burrill 1882) Winslow <i>et al.</i>, 1920</p> <p>Synonyms: <i>Micrococcus amylovorus</i> Burrill 1882</p> <p><i>Bacillus amylovorus</i> (Burrill 1882) Trevisan 1889</p> <p>"<i>Bacterium amylovorus</i>" [sic] (Burrill 1882) Chester 1897</p> <p><i>Erwinia amylovora</i> f.sp. <i>rubi</i> Starr, Cardona and Falson (Starr <i>et al.</i>, 1951)</p> <p>Taxonomic position: Proteobacteria, Y subdivision, Enterobacteriales, Enterobacteriaceae</p> <p>Common names: Fire blight (EPPO, 2013)</p> | 1) For <i>Bacillus amylovorus</i> , the year is missing after "Burrill". 2) For <i>Erwinia amylovora</i> f.sp. <i>rubi</i> , prevention of a repetition. NB: it should be "Folsom" not "Falson". | EPPO, European Union, Georgia, Serbia |

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| 10. | 9 | Editorial | 3. Detection | The number of the section is missing. | EPPO, European Union, Georgia, Serbia |
| 11. | 9 | Editorial | 3. Detection | Editorial correction | Japan |
| 12. | 11 | Editorial | 3.1 Detection in plants with symptoms | It's more clarity. | China |
| 13. | 13 | Editorial | <p>The recommended procedures are indicated in the Figure1.</p> <p>Symptoms of fire blight on the most common hosts such as <i>Pyrus communis</i> (pear), <i>Malus domestica</i>, (apple), <i>Cydonia</i> spp. (quince), <i>Eriobotrya japonica</i> (loquat), <i>Cotoneaster</i> spp. (cotoneaster), <i>Pyracantha</i> spp. (pyracantha) and <i>Crataegus</i> spp. (hawthorn) are similar and easily recognized. The name of the disease is descriptive of its major characteristic: the brownish, necrotic appearance of twigs, flowers and leaves, as though they had been burned by fire. The typical symptoms are the brown to black colour of leaves on affected branches, the production of exudates, and the characteristic “shepherd’s crook” of terminal shoots. Depending on the affected plant part, the disease produces blossom blight, shoot/twig blight, leaf blight, fruit blight, limb/trunk blight or collar/rootstock blight (van der Zwet and Beer, 1995; van der Zwet and Keil, 1979).</p> <p>In apple and pear trees the first symptoms usually appear in early spring when average temperatures rise above 15 °C, during humid weather. Infected blossoms become soaked with water, then wilt, shrivel, and turn orange or brown to black. Peduncles may also appear water-soaked, become dark green and finally brown or black, sometimes oozing droplets of sticky bacterial exudate. Infected leaves wilt and shrivel, and entire spurs turn brown in apples and dark brown to black in pears, but remain attached to the tree for some time. Upon infection young fruitlets turn brown but also remain attached to the tree. Immature fruit lesions appear oily or water-soaked, becoming brown to black and often exuding droplets of bacterial ooze. Characteristic reddish-brown streaks are often found in the subcortical tissues when the bark is peeled from infected limbs or twigs (Thomson, 2000; van der Zwet and Keil, 1979). Brown to black slightly depressed cankers form in the bark of twigs, branches or the trunk of infected trees. These cankers later become defined by cracks near the margin of diseased and healthy tissue (Thomson, 2000).</p> <p>Confusion may occur between fire blight and blight- or blast-like symptoms –</p> | To prevent confusion between Pyrus and Pyracantha and Malus and Mespilus. | EPPO, European Union, Georgia, Serbia |

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| | | | <p>especially in blossoms and buds – caused by other pathogenic bacteria and fungi, insect damage or physiological disorders. Other bacteria that cause fire blight-like symptoms include <i>Erwinia pyrifoliae</i>, the causal agent of bacterial shoot blight of <i>Pyrus pyrifolia</i> (Asian pear) (Kim <i>et al.</i>, 1999); <i>Erwinia piriflorinigrans</i>, isolated from necrotic pear blossoms in Spain (López <i>et al.</i>, 2011); <i>Erwinia uzenensis</i>, recently described in Japan (Matsuura <i>et al.</i>, 2012); other <i>Erwinia</i> spp. reported in Japan that cause bacterial shoot blight (Kim <i>et al.</i>, 2001a, 2001b; Palacio-Bielsa <i>et al.</i>, 2012; Tanii <i>et al.</i>, 1981); and <i>Pseudomonas syringae</i> pv. <i>syringae</i>, the causal agent of blossom blast. A definitive diagnosis of fire blight should always be obtained through laboratory analysis.</p> | | |
| 14. | 15 | Technical | <p>Plant material should be analysed as soon as possible after collection, but may be stored at 4–8 °C for up to two <u>one</u> weeks until processing. Precautions to avoid cross-contamination should be taken when collecting samples, during transport and processing, and especially while isolating the bacterium or extracting DNA.</p> <p>The samples should be processed with a general procedure valid for isolation, serological tests and polymerase chain reaction (PCR) analysis, before or after enrichment. The use of freshly prepared antioxidant maceration buffer (polyvinylpyrrolidone (PVP)-10, 20 g; mannitol, 10 g; ascorbic acid, 1.76 g; reduced glutathione, 3 g; phosphate-buffered saline (PBS), 10 mM, 1 litre; pH 7.2; sterilized by filtration) is required for successful enrichment, as indicated by Gorris <i>et al.</i> (1996). The samples can be processed also in <u>sterile water or in</u> PBS, pH 7.2 (NaCl, 8 g; KCl, 0.2 g; Na₂HPO₄·12H₂O, 2.9 g; KH₂PO₄, 0.2 g; distilled water, 1 litre) for direct isolation, immunofluorescence (IF) or PCR.</p> <p>Plant parts (flowers, shoots, twigs, leaves or fruit) showing the most typical symptoms, and with bacterial exudate if possible, are carefully selected. Material for processing is selected from the leading edge of disease lesions. The plant tissue is cut into pieces of approximately 0.1–1.0 g, lightly crushed in antioxidant maceration buffer, <u>PBS or sterile water</u> (described in the previous paragraph) at 1:50 (w/v), left to stand for at least 5 min, and placed on ice for a few minutes. Triplicate samples (1 ml each) of each macerate are transferred to sterile micro centrifuge tubes, with one tube stored at –20 °C for subsequent analysis by PCR and another tube's contents adjusted to 30% glycerol and stored at –80 °C for confirmation testing, if necessary. The third tube is kept on ice for performing enrichment before enzyme-linked immunosorbent assay (ELISA) or PCR, and isolation on selective media (Figure 1). If IF is to be performed (i.e. IF analysis is optional), the slides are prepared and fixed on the same day that the samples are macerated. The PCR</p> | <p>1. First paragraph, first sentence: "two weeks" is too long. The standard practice seems to be a storage for up to one week before processing and up to two weeks after.</p> <p>2. Second paragraph: the end of the first sentence 'before or after enrichment' implies that isolation without enrichment is not possible. This is not consistent with the flow diagram where both isolation and enrichment isolation are mentioned. Both options should be possible. Suggest deletion of the end of the sentence.</p> <p>3. Second paragraph, last sentence: replace "in PBS" by "in sterile water or in PBS"; both options should be given.</p> <p>4. Third paragraph, third sentence: replace "in antioxidant maceration buffer" by "in antioxidant maceration buffer, PBS or sterile water".</p> | EPPO, European Union, Georgia, Serbia |

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| | | | analysis should be performed as soon as is convenient, using the macerated sample stored at –20 °C. | | |
| 15. | 18 | Editorial | <p>When symptoms are very advanced or the environmental conditions after infection are not favourable for bacterial multiplication, the number of culturable <i>E. amylovora</i> cells can be very low. Isolation under these conditions can result in plates with few cells of the pathogen and that can be overcrowded with saprophytic and antagonistic bacteria. If this is suspected, the sample should be re-tested and/or enriched before isolation. The induction of the reversible and viable but non-culturable state has been described for <i>E. amylovora in vitro</i> using copper treatments and in fruits (Ordax <i>et al.</i>, 2009), and it can be the cause of false negative isolation results.</p> <p>In general, plating on three media is advised for maximum likelihood of recovery of <i>E. amylovora</i>, especially when samples are not in good condition. Depending on the number and microbial composition of the sample, each medium can be more or less efficient. Three media (CCT, King's B and Levan) have been validated in two ring tests, with Levan having the highest plating efficiency.</p> | The first paragraph should be put after the second paragraph. | EPPO, European Union, Georgia, Serbia |
| 16. | 22 | Editorial | <p>Cycloheximide is added at 0.05 g/litre to King's B and Levan media when fungi are expected in the isolations. Dilutions of 1:10 and 1:100 of each macerate are prepared in PBS (NaCl, 8 g; KCl, 0.2 g; Na₂HPO₄·12H₂O, 2.9 g; KH₂PO₄, 0.2 g; distilled water, 1 litre).</p> <p>Preferably 100 µl of the macerates and their dilutions are spread, by triple streaking in 130 mm plates or fifty microliters spread in standard 90 mm Petri dishes. Plates are incubated at 25 °C for up to 4 days. The final reading is usually taken at 72 h. Colonies of <i>E. amylovora</i> on CCT medium are pale violet, circular, high convex to domed, smooth and mucoid and they grow more slowly than on King's B or Levan media. Colonies on King's B medium are creamy white, circular and non-fluorescent under ultraviolet (UV) light at 366 nm. Colonies on Levan medium are white, circular, domed, smooth and mucoid. Levan-negative colonies of <i>E. amylovora</i> have been reported (Bereswill <i>et al.</i>, 1997).</p> <p>Pure cultures are obtained from individual suspect colonies of each sample by dilution and streaking onto King's B medium. Presumptive colonies of <i>E. amylovora</i> are identified preferably by inoculating any available <i>E. amylovora</i> host to test pathogenicity; by double antibody sandwich indirect (DASI)-ELISA, PCR or by other appropriate tests (e.g. biochemical tests, IF, fatty acid profile), <u>or by inoculating any available <i>E. amylovora</i> host to test pathogenicity</u>, as indicated in section 4.</p> | 1. Second paragraph, last but one sentence: one comma to delete after "circular". 2. Third paragraph, second sentence: "by inoculating..." to be transferred to the end of the sentence after the other identification tests are positive. | EPPO, European Union, Georgia, Serbia |

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| | | | <p>When analysing symptomatic samples, good correlation is expected between isolation, IF, enrichment DASI-ELISA (see section 3.1.4.1) and PCR.</p> <p>In the 2003 and 2010 ring tests, the accuracy of isolation was 0.88 and 0.81 for King's B, 0.92 and 0.89 for Levan, and 0.92 and 0.95 for CCT media, respectively (López <i>et al.</i>, 2006; Lopez, IVIA, Spain, personal communication, 2012). In the 2009 ring test, accuracy of isolation was 0.96 for CCT (Dreo <i>et al.</i>, 2009).</p> | | |
| 17. | 24 | Editorial | <p>Enrichment is used to multiply the initial population of culturable <i>E. amylovora</i> in a sample and to perform enrichment DASI-ELISA <u>or PCR</u>. It should be carried out before isolation (even for symptomatic samples) when a low number of culturable <i>E. amylovora</i> cells is expected to be present (e.g. for copper-treated samples, samples with old symptoms, samples collected during unfavourable weather conditions for fire blight such as the winter season). The enrichment step greatly increases the sensitivity of DASI-ELISA. The use of two validated liquid media for enrichment – one non-selective (King's B) and one semi-selective (CCT) – is advised because the composition and population size of the microbiota is unknown.</p> <p>The tissue sample is macerated as described in section 3.1.2 and 0.9 ml is immediately dispensed into each of two sterile 10–15 ml tubes (to ensure sufficient aeration) containing 0.9 ml of each liquid enrichment medium (King's B without agar, and CCT made with nutrient broth instead of nutrient agar). The tubes are incubated at 25 °C for 48–72 h without shaking. A longer incubation is recommended when processing plant samples collected in winter. Both enrichment broths and the 1:10 and 1:100 dilutions prepared in PBS are spread onto CCT plates, by triple streaking, to obtain isolated colonies. Plates are incubated at 25 °C for 72–96 h. Final reading of the CCT plates is at 72 h and must be followed by purification of colonies and identification.</p> <p>The use of semi-selective medium for plating and dilution is advised because the enrichment step will permit growth of the pathogen but will also allow abundant multiplication of other bacteria. The accuracy of the enrichment isolation on King's B and CCT was 0.97 in the 2010 ring test.</p> | <p>First paragraph, first sentence: "and to perform enrichment DASI-ELISA": Paragraph about enrichment should provide an introduction to all methods post-enrichment. So need to add PCR.</p> | EPPO, European Union, Georgia, Serbia |
| 18. | 27 | Editorial | <p>A kit for enrichment DASI-ELISA has been validated in two ring tests and is available commercially from Plant Print Diagnostics SL, Valencia, Spain. It is based on the mixture of two specific monoclonal antibodies described in Gorris <i>et al.</i> (1996) and requires prior enrichment of the samples, as previously described. The following protocol must be followed strictly for maximum accuracy. Before ELISA, the required amount of the enriched extracts and controls is treated by incubating in a water bath</p> | <p>The reference for the ring test 2010 is López et al 2010</p> | EPPO, European Union, Georgia, Serbia |

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| | | | <p>at 100 °C for 10 min. This treatment is necessary for optimum specificity. The boiled samples are processed (at room temperature) by ELISA on the same day (or stored at –20 °C for subsequent analysis) following the instructions provided by the manufacturer of the commercial kit.</p> <p>The ELISA is negative if the average optical density (OD) reading from duplicate sample wells is <2x the OD in the negative sample extract control wells (providing the OD for the positive control wells are above 1.0 after 90 min incubation and are greater than twice the OD obtained for negative sample extracts). The ELISA is positive if the average OD reading from duplicate sample wells is >2x the OD in the negative sample extract control wells (providing all negative control wells are lower than 2x the average OD reading of the positive control wells).</p> <p>Negative ELISA readings in positive control wells indicate that the test has not been performed correctly and/or the reagents were not well prepared. Positive ELISA readings in negative control wells indicate cross-contamination or non-specific antibody binding. In both cases, the test should be repeated or a second test based on a different biological principle, such as PCR, should be performed.</p> <p>The accuracy of the DASI-ELISA was 0.79 and 0.82 for enrichment in King's B medium (King's B-DASI-ELISA), and 0.83 and 0.77 for enrichment in CCT medium (CCT-DASI-ELISA) in the 2003 and 2010 ring tests, respectively (López <i>et al.</i>, 2006; 2010 López 2013, personnel communication unpublished).</p> | | |
| 19. | 27 | Substantive | <p>A kit for enrichment DASI-ELISA has been validated in two ring tests and is available commercially from Plant Print Diagnòstics SL, Valencia, Spain. It is based on the mixture of two specific monoclonal antibodies described in Gorris <i>et al.</i> (1996) and requires prior enrichment of the samples, as previously described. The following protocol must be followed strictly for maximum accuracy. Before ELISA, the required amount of the enriched extracts and controls is treated by incubating in a water bath at 100 °C for 10 min. This treatment is necessary for optimum specificity. The boiled samples are processed (at room temperature) by ELISA on the same day (or stored at –20 °C for subsequent analysis) following the instructions provided by the manufacturer of the commercial kit.</p> <p>The ELISA is negative if the average optical density (OD) reading from duplicate sample wells is <2x the OD in the negative sample extract control wells (providing the OD for the positive control wells are above 1.0 after 90 min incubation and are greater than twice the OD obtained for negative sample extracts). The ELISA is</p> | Print Diagnòstics SL, Valencia, Spain - It is not appropriate to use commercial names. It should be mentioned in a correspondent footnote or deleted. | COSAVE, Uruguay, Chile, Brazil, Peru, Argentina |

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| | | | <p>positive if the average OD reading from duplicate sample wells is >2x the OD in the negative sample extract control wells (providing all negative control wells are lower than 2x the average OD reading of the positive control wells).</p> <p>Negative ELISA readings in positive control wells indicate that the test has not been performed correctly and/or the reagents were not well prepared. Positive ELISA readings in negative control wells indicate cross-contamination or non-specific antibody binding. In both cases, the test should be repeated or a second test based on a different biological principle, such as PCR, should be performed.</p> <p>The accuracy of the DASI-ELISA was 0.79 and 0.82 for enrichment in King's B medium (King's B-DASI-ELISA), and 0.83 and 0.77 for enrichment in CCT medium (CCT-DASI-ELISA) in the 2003 and 2010 ring tests, respectively (López <i>et al.</i>, 2006; López 2013, personnel communication unpublished).</p> | | |
| 20. | 29 | Technical | <p>To make tissue prints, freshly cut plant sections are pressed carefully against a nitrocellulose membrane. Prints are prepared for positive and negative controls. Printed membranes can be kept for several months in a dry place at room temperature. A kit is available commercially from Plant Print Diagnostics. A validated source of antibodies to <i>E. amylovora</i> should be used. To develop prints, the manufacturer's instructions should be followed. The prints are observed under low power magnification (x10 or x20). The test is positive when purple–violet precipitates appear in the sections of plant material that is printed on the membrane and not in the plant material print of the negative control. If exudates or colonies are printed they should appear violet when positive. The test is negative when no purple–violet precipitates appear, as in the negative control.</p> | <p>Add: "A validated source of antibodies to <i>E. amylovora</i> should be used."</p> | <p>EPPO, European Union, Georgia, Serbia</p> |
| 21. | 31 | Substantive | <p>Immunofluorescence (IF) is a recommended alternative serological method, and it is easy to follow the standard protocol (Anonymous, 1998). A validated source of antibodies to <i>E. amylovora</i> should be used. Two commercial antibodies have been validated in one ring test: one monoclonal antibody is available through Plant Print Diagnostics SL and one polyclonal antibody from Loewe Biochemicals (Sauerlach, Germany).</p> <p>The IF should be performed on fresh sample extracts fixed onto slides. Undiluted macerates and dilutions of 1:10 and 1:100 in PBS are used to spot windows of the IF slides. The monoclonal or polyclonal antibodies are used at the appropriate dilutions in PBS. The appropriate fluorescein isothiocyanate (FITC) conjugates are diluted in PBS: goat anti-mouse for monoclonal antibodies (GAM-FITC), and goat</p> | <p>(Loewe Biochemicals (Sauerlach, Germany)) - We consider that it is not necessary to include commercial kit names because it is already mentioned that is important to use validated source of antibodies. Commercial names should be mentioned in a correspondent footnote or deleted.</p> | <p>COSAVE, Uruguay, Chile, Brazil, Peru, Argentina</p> |

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| | | | <p>anti-rabbit (GAR-FITC) or anti-goat for polyclonal antibodies.</p> <p>The test on a sample is negative if green fluorescing cells with morphology typical of <i>E. amylovora</i> are observed in the positive controls, but not in the sample windows. The test on a sample is positive if green fluorescing cells with typical morphology are observed in the positive controls and in the sample windows, but not in the negative control windows. As a population of 10^3 cells/ml is considered the limit for reliable detection by IF, for samples with $>10^3$ cells/ml, the IF test is considered positive. For samples with $<10^3$ cells/ml, or weakly fluorescing cells, the result of the IF may be considered uncertain.</p> <p>The accuracy of IF in the 2003 ring test was 0.70 for the Plant Print Diagnostics SL monoclonal antibody, and 0.72 for the Loewe Biochemicals polyclonal antiserum, confirming that the sensitivity of the technique is around 10^3 c.f.u./ml.</p> | | |
| 22. | 33 | Substantive | Two lateral flow devices are available commercially for rapid analysis of plant material: Ea AgriStrip (Bioreba, Reinach, Switzerland) and Pocket Diagnostics (Forsite Diagnostics, York, UK). Following the manufacturers' instructions their accuracy in the ring tests was 0.66 and 0.55 for Ea AgriStrip and 0.64 and 0.56 for Pocket Diagnostics in the 2009 and 2010 ring tests, respectively. These results were obtained for the detection of <i>E. amylovora</i> in samples from 1 –to 10^6 c.f.u./g, but the accuracy was approximately 1.0 when analysing samples with 10^5 – 10^6 c.f.u./g, the minimum number expected in symptomatic samples (López <i>et al.</i> , 2010). The kits are recommended for use only with symptomatic samples. | To add a Footnote 1: The use of the brand Bioreba and Forsite Diagnostics for the lateral flow devices in this diagnostic protocol implies no approval of them to the exclusion of others that may also be suitable. This information is given for the convenience of users of this protocol and does not constitute an endorsement by the CPM of the chemical, reagent and/or equipment named. Equivalent products may be used if they can be shown to lead to the same results | COSAVE, Uruguay, Chile, Brazil, Peru, Argentina |
| 23. | 37 | Editorial | <p>For the test result obtained to be regarded as reliable the following controls should be considered for each series of nucleic acid isolations and amplification of the target nucleic acid. Controls used will depend on the test used and the level of certainty required. For PCR, a positive nucleic acid control, an internal control and a negative amplification control (no template control) are the minimum controls that should be used.</p> <p>Positive nucleic acid control</p> <p>This control is used to monitor the efficiency of the PCR amplification of the target nucleic acid (but does not test the nucleic acid extraction procedure). Pre-prepared (stored) nucleic acid, whole genome amplified DNA or a synthetic control (e.g.</p> | In the section "Positive nucleic acid control", a full stop is missing at the end of the first sentence. | EPPO, European Union, Georgia, Serbia |

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| | | | <p>cloned PCR product) may be used.</p> <p>Internal control</p> <p>For conventional and real-time PCR, plant internal controls (House Keeper Genes (HKG) such as COX (Weller <i>et al.</i>, 2000) or 16S rDNA (Weisberg <i>et al.</i>, 1991) could be incorporated into the PCR protocols to eliminate the possibility of PCR false negatives due to nucleic acid extraction failure or degradation or the presence of PCR inhibitors.</p> <p>Negative amplification control (no template control).</p> <p>This is a necessary control for conventional and real-time PCR to rule out false positives due to contamination during preparation of the reaction mixture. PCR-grade water that was used to prepare the reaction mixture is added at the amplification stage.</p> <p>Positive extraction control</p> <p>This control is used to ensure that nucleic acid from the target is of sufficient quantity and quality and that the target is detected. Nucleic acid is extracted from infected host tissue or healthy plant tissue that has been spiked with the target.</p> <p>The positive control should be approximately one-tenth of the amount of leaf tissue used per plant for the DNA extraction. For PCR, care needs to be taken to avoid cross-contamination due to aerosols from the positive control or from positive samples. If required, the positive control used in the laboratory should be sequenced so that this sequence can be readily compared with sequences obtained from PCR amplicons of the correct size. Alternatively, synthetic positive controls can be made with a known sequence, which, again, can be compared to PCR amplicons of the correct size.</p> <p>Negative extraction control</p> <p>This control is used to monitor contamination during nucleic acid extraction as well as cross-reactions with the host tissue. The control requires nucleic acid that is</p> | | |

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| | | | extracted from uninfected host tissue and subsequently amplified. It is recommended that multiple controls are included when large numbers of positives are expected. | | |
| 24. | 37 | Technical | <p>For the test result obtained to be regarded as reliable the following controls should be considered for each series of nucleic acid isolations and amplification of the target nucleic acid. Controls used will depend on the test used and the level of certainty required. For PCR, a positive nucleic acid control, an internal control and a negative amplification control (no template control) are the minimum controls that should be used.</p> <p>Positive nucleic acid control</p> <p>This control is used to monitor the efficiency of the PCR amplification of the target nucleic acid (but does not test the nucleic acid extraction procedure) Pre-prepared (stored) nucleic acid, whole genome amplified DNA or a synthetic control (e.g. cloned PCR product) may be used.</p> <p>Internal control</p> <p>For conventional and real-time PCR, plant internal controls (House Keeper Genes (HKG) such as COX (Weller <i>et al.</i>, 2000) or 16S rDNA (Weisberg <i>et al.</i>, 1991) could be may be incorporated into the PCR protocols to eliminate the possibility of PCR false negatives due to nucleic acid extraction failure or degradation or the presence of PCR inhibitors.</p> <p>Negative amplification control (no template control).</p> <p>This is a necessary control for conventional and real-time PCR to rule out false positives due to contamination during preparation of the reaction mixture. PCR-grade water that was used to prepare the reaction mixture is added at the amplification stage.</p> <p>Positive extraction control</p> <p>This control is used to ensure that nucleic acid from the target is of sufficient quantity and quality and that the target is detected. Nucleic acid is extracted from infected</p> | In the section "Internal control", "could be" should be replaced by "may be". | EPPO, European Union, Georgia, Serbia |

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| | | | <p>host tissue or healthy plant tissue that has been spiked with the target.</p> <p>The positive control should be approximately one-tenth of the amount of leaf tissue used per plant for the DNA extraction. For PCR, care needs to be taken to avoid cross-contamination due to aerosols from the positive control or from positive samples. If required, the positive control used in the laboratory should be sequenced so that this sequence can be readily compared with sequences obtained from PCR amplicons of the correct size. Alternatively, synthetic positive controls can be made with a known sequence, which, again, can be compared to PCR amplicons of the correct size.</p> <p>Negative extraction control</p> <p>This control is used to monitor contamination during nucleic acid extraction as well as cross-reactions with the host tissue. The control requires nucleic acid that is extracted from uninfected host tissue and subsequently amplified. It is recommended that multiple controls are included when large numbers of positives are expected.</p> | | |
| 25. | 39 | Substantive | <p>Three DNA extraction methods – Llop <i>et al.</i> (1999), Taylor <i>et al.</i> (2001) and the REDExtract-N-AmpPlant PCR Kit (Sigma-Aldrich, USA) – were evaluated in the 2009 ring test (Dreo <i>et al.</i>, 2009), with four PCR protocols with accuracies ranging from 0.67 to 0.76. The methods showed comparable results in the 2010 ring test (Lopez <i>et al.</i>, 2010), as indicated below in the accuracies given for the different PCR methods. Their efficiencies did not improve after diluting the extracts 1:10, suggesting that few or no inhibitors were present. Based on these findings, the Llop <i>et al.</i> (1999) extraction method is recommended as it has been extensively tested in a number of countries and is cheap and easy to set up in the laboratory.</p> <p>DNA extraction according to Llop <i>et al.</i>(1999)</p> <p>One millilitre of a sample macerate prepared according to section 3.1.2 and/or 1 ml enriched macerate is centrifuged at 10 000 g for 5 min at room temperature. The supernatant is discarded, and the pellet is resuspended in 500 µl extraction buffer</p> | <p>To add a footnote 2: The use of the brand Sigma for the REDExtract-N-AmpPlant PCR kit in this diagnostic protocol implies no approval of them to the exclusion of others that may also be suitable. This information is given for the convenience of users of this protocol and does not constitute an endorsement by the CPM of the chemical, reagent and/or equipment named. Equivalent products may be used if they can be shown to lead to the same results</p> | <p>COSAVE, Uruguay, Chile, Brazil, Peru, Argentina</p> |

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| | | | (Tris-HCl pH 7.5, 24.2 g; NaCl, 14.6 g; ethylenediaminetetraacetic acid (EDTA), 9.3 g; sodium dodecyl sulphate (SDS), 5 g; PVP-10, 20 g; distilled water, 1 litre; sterilized by filtration) and incubated for 1 h at room temperature before centrifugation at 4 000 g for 5 min. Approximately 450 µl supernatant is mixed with an equal volume of isopropanol, inverted, and left at room temperature for 30 min to 1 h. The precipitated nucleic acid is centrifuged at 10 000 g for 5 min, the supernatant is discarded and the pellet is air-dried. If there is still a coloured precipitate (brown or green) at the bottom of the tube, this is carefully removed while discarding the supernatant, thus obtaining a cleaner DNA pellet. The pellet is resuspended in 200 µl water. It should be used for PCR immediately or stored at – 20 °C. | | |
| 26. | 41 | Editorial | <p>There are many PCR primers and protocols described for <i>E. amylovora</i> detection and some have shown specificity problems (Powney <i>et al.</i>, 2011a; Roselló <i>et al.</i>, 2006). The primers and protocols validated in ring tests were those of Bereswill <i>et al.</i> (1992) and Llop <i>et al.</i> (2000), with or without previous enrichment, in 2003; and those of Taylor <i>et al.</i> (2001), Stöger <i>et al.</i> (2006) and Obradovic <i>et al.</i> (2007) in 2009 and 2010. The discovery of fully virulent <i>E. amylovora</i> strains without the pEA29 plasmid (Llop <i>et al.</i>, 2006) and experiences from different countries (Powney <i>et al.</i>, 2011a) indicate that two PCR protocols should be used: one with primers based on pEA29 sequences, and another targeting unique chromosomal sequences. PCR can be carried out using the primers and conditions validated in the ring tests, although amplification conditions should be optimized for different thermo cyclers.</p> <p>PCR according to Bereswill <i>et al.</i> (1992)</p> <p>The primers are: Forward primer: A: 5'-CGG TTT TTA ACG CTG GG-3'</p> <p>Reverse primer: B: 5'-GGG CAA ATA CTC GGA TT-3'.</p> <p>The targeted sequences are in the plasmid pEA29. The PCR mixture is composed of: ultrapure water, 17.4 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 1.5 µl; dNTPs 10 mM, 0.5 µl; primer A 10 pmol/µl, 0.25 µl; primer B 10 pmol/µl, 0.25 µl; Taq DNA polymerase 5 U/µl, 0.1 µl. The extracted DNA sample volume is 2.5 µl, and should be added to 22.5 µl of the PCR mix. The cycling parameters are: a denaturation step of 93 °C for 5 min followed by 40 cycles of 93 °C for 30 s, 52 °C for 30 s and 72 °C for 1 min 15 s, with a final elongation step at 72 °C for 10 min. The amplicon size is 900 base pairs (bp) according to Bereswill <i>et al.</i> (1992), although variations in size</p> | 1) Section "PCR according to Stöger <i>et al.</i> (2006)", last paragraph: clearer. 2) Section "PCR according to Gottsberger...(2007)", last paragraph: clearer. 3) Section "Nested PCR according to Llop <i>et al.</i> (2000): a blank is missing in the expression "Forward primer: PEANT1". | EPPO, European Union, Georgia, Serbia |

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| | | | <p>can occur between 900 and 1 100 bp depending on the number of 8 bp repeats within the amplified fragment (Jones and Geider, 2001).</p> <p>The accuracy in the 2004 ring test was 0.51 but increased to 0.74 and 0.78 after enrichment of the samples in King's B and CCT media, respectively (López <i>et al.</i>, 2006).</p> <p>PCR according to Taylor <i>et al.</i> (2001)</p> <p>The primers are:</p> <p>G1-F: 5'-CCT GCA TAA ATC ACC GCT GAC AGC TCA ATG-3'</p> <p>G2-R: 5'-GCT ACC ACT GAT CGC TCG AAT CAA ATC GGC-3'.</p> <p>The targeted sequences are chromosomal. The PCR mixture is composed of: ultrapure water, 14.3 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 0.75 µl; dNTPs 10 mM, 0.25 µl; G1-F 10pmol/µl, 1 µl; G2-R 10pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.2 µl. An extracted DNA sample of 5 µl is added to 45 µl PCR mix. The cycling parameters are: 95 °C for 3 min followed by 40 cycles of 94 °C for 30 s, 60 °C for 30 s and 72 °C for 1 min, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The amplicon size is 187 bp.</p> <p>The accuracy in the 2010 ring test was 0.77 using the Llop <i>et al.</i> (1999) DNA extraction procedure.</p> <p>PCR according to Stöger <i>et al.</i> (2006)</p> <p>The primers (from Llop <i>et al.</i>, 2000) are:</p> <p>Forward primer: PEANT 1-F: 5'-TAT CCC TAA AAA CCT CAG TGC-3'</p> <p>Reverse primer: PEANT 2-R: 5'-GCA ACC TTG TGC CCT TTA-3'.</p> <p>The targeted sequences are in the plasmid pEA29. Stöger <i>et al.</i> (2006)</p> | | |

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| | | | <p>recommended this method be used with DNA extracted using the REExtract-N-AmpT Kit (Sigma-Aldrich). The PCR mixture is composed of: ultrapure water, 5 µl; REExtract-N-Amp PCR Ready Mix, 10 µl; PEANT 1-F 10 pmol/µl, 0.5 µl; PEANT 2-R 10 pmol/µl, 0.5 µl; extracted DNA, 4 µl. The cycling parameters are: 95 °C for 5 min followed by 35 cycles of 95 °C for 15 s, 58 °C for 30 s, and 72 °C for 45 s, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The expected amplicon is 391 bp.</p> <p>The accuracy <u>was 0.76</u> in the 2009 ring test was 0.76 and <u>0.72</u> in the 2010 ring test 0.72.</p> <p>PCR according to Gottsberger adapted from Obradovic <i>et al.</i> (2007)</p> <p>The primers are:</p> <p>FER1-F: 5'-AGC AGC AAT TAA TGG CAA GTA TAG TCA-3'</p> <p>rgER2-R: 5'-AAA AGA GAC ATC TGG ATT CAG ACA AT-3'.</p> <p>The targeted sequences are chromosomal. The mixture is composed of: ultrapure water, 14.3 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 0.75 µl; dNTPs 10 mM, 0.25 µl; FER1-F 10 pmol/µl, 1 µl; rgER2-R 10 pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.2 µl; extracted DNA, 5 µl. The cycling parameters are: 94 °C for 3 min followed by 41 cycles of 94 °C for 10 s, 60 °C for 10 s and 72 °C for 30 s, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The expected amplicon size is 458 bp.</p> <p>The accuracy <u>was 0.76</u> in the 2009 ring test was 0.76, and <u>0.68</u> in the 2010 ring test 0.68, using the DNA extraction method described by Llop <i>et al.</i> (1999).</p> <p>Nested PCR according to Llop <i>et al.</i> (2000)</p> <p>The nested-PCR of Llop <i>et al.</i> (2000) uses two sets of primers, which are combined in a single reaction tube. Because of the different annealing temperatures the two PCRs are run consecutively. The external primers are those designed by McManus and Jones (1995) and are based on sequences of the pEA29 plasmid. The internal</p> | | |

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| | | | <p>primers are those described by Llop <i>et al.</i> (2000). The sequences are the following:</p> <p>External primers:</p> <p>Forward primer: AJ75: 5'-CGT ATT CAC GGC TTC GCA GAT-3'</p> <p>Reverse primer: AJ76: 5'-ACC CGC CAG GAT AGT CGC ATA-3'</p> <p>Internal primers:</p> <p>Forward primer: PEANT1: 5'-TAT CCC TAA AAA CCT CAG TGC-3'</p> <p>Reverse primer: PEANT2: 5'-GCA ACC TTG TGC CCT TTA-3'</p> <p>The PCR mixture is composed of: ultrapure water, 36.25 µl; buffer 10x, 5 µl; MgCl₂ 50 mM, 3 µl; dNTPs 10 mM, 0.5 µl; AJ75 0.1 pmol/µl, 0.32 µl; AJ76 0.1 pmol/µl, 0.32 µl; PEANT1 10 pmol/µl, 1 µl; primer PEANT2 10 pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.6 µl. A DNA sample volume of 2 µl should be added to 48 µl PCR mix. The cycling parameters are: a denaturation step of 94 °C for 4 min followed by 25 cycles of 94 °C for 60 s and 72 °C for 90 s. This first round PCR is followed in the same thermocycler by a second denaturation step of 94 °C for 4 min and 40 cycles of 94 °C for 60 s, 56 °C for 60 s, and 72 °C for 60 s, with a final elongation step at 72 °C for 10 min. The expected amplicon size is 391 bp, although variations in size can occur.</p> <p>The accuracy was 0.69 and 0.72 in the 2003 and 2010 ring tests, respectively, but increased after enrichment up to 0.84 (King's B medium) and 0.86 (CCT medium) in the 2003 ring test, and to 0.79 (King's B) and 0.88 (CCT) in 2010.</p> | | |
| 27. | 41 | Substantive | <p>There are many PCR primers and protocols described for <i>E. amylovora</i> detection and some have shown specificity problems (Powney <i>et al.</i>, 2011a; Roselló <i>et al.</i>, 2006). The primers and protocols validated in ring tests were those of Bereswill <i>et al.</i> (1992) and Llop <i>et al.</i> (2000), with or without previous enrichment, in 2003; and those of Taylor <i>et al.</i> (2001), Stöger <i>et al.</i> (2006) and Obradovic <i>et al.</i> (2007) in 2009 and 2010. The discovery of fully virulent <i>E. amylovora</i> strains without the pEA29 plasmid (Llop <i>et al.</i>, 2006) and experiences from different countries (Powney <i>et al.</i>, 2011a) indicate that two PCR protocols should be used: one with</p> | REExtract-N-AmpT Kit (Sigma-Aldrich). - Related to footnote 2. | COSAVE, Uruguay, Chile, Brazil, Peru, Argentina |

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| | | | <p>primers based on pEA29 sequences, and another targeting unique chromosomal sequences. PCR can be carried out using the primers and conditions validated in the ring tests, although amplification conditions should be optimized for different thermo cyclers.</p> <p>PCR according to Bereswill <i>et al.</i> (1992)</p> <p>The primers are: Forward primer: A: 5'-CGG TTT TTA ACG CTG GG-3'</p> <p>Reverse primer: B: 5'-GGG CAA ATA CTC GGA TT-3'.</p> <p>The targeted sequences are in the plasmid pEA29. The PCR mixture is composed of: ultrapure water, 17.4 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 1.5 µl; dNTPs 10 mM, 0.5 µl; primer A 10 pmol/µl, 0.25 µl; primer B 10 pmol/µl, 0.25 µl; Taq DNA polymerase 5 U/µl, 0.1 µl. The extracted DNA sample volume is 2.5 µl, and should be added to 22.5 µl of the PCR mix. The cycling parameters are: a denaturation step of 93 °C for 5 min followed by 40 cycles of 93 °C for 30 s, 52 °C for 30 s and 72 °C for 1 min 15 s, with a final elongation step at 72 °C for 10 min. The amplicon size is 900 base pairs (bp) according to Bereswill <i>et al.</i> (1992), although variations in size can occur between 900 and 1 100 bp depending on the number of 8 bp repeats within the amplified fragment (Jones and Geider, 2001).</p> <p>The accuracy in the 2004 ring test was 0.51 but increased to 0.74 and 0.78 after enrichment of the samples in King's B and CCT media, respectively (López <i>et al.</i>, 2006).</p> <p>PCR according to Taylor <i>et al.</i> (2001)</p> <p>The primers are:</p> <p>G1-F: 5'-CCT GCA TAA ATC ACC GCT GAC AGC TCA ATG-3'</p> <p>G2-R: 5'-GCT ACC ACT GAT CGC TCG AAT CAA ATC GGC-3'.</p> <p>The targeted sequences are chromosomal. The PCR mixture is composed of: ultrapure water, 14.3 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 0.75 µl; dNTPs 10 mM,</p> | | |

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| | | | <p>0.25 µl; G1-F 10pmol/µl, 1 µl; G2-R 10pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.2 µl. An extracted DNA sample of 5 µl is added to 45 µl PCR mix. The cycling parameters are: 95 °C for 3 min followed by 40 cycles of 94 °C for 30 s, 60 °C for 30 s and 72 °C for 1 min, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The amplicon size is 187 bp.</p> <p>The accuracy in the 2010 ring test was 0.77 using the Llop <i>et al.</i> (1999) DNA extraction procedure.</p> <p>PCR according to Stöger <i>et al.</i> (2006)</p> <p>The primers (from Llop <i>et al.</i>, 2000) are:</p> <p>Forward primer: PEANT 1-F: 5'-TAT CCC TAA AAA CCT CAG TGC-3'</p> <p>Reverse primer: PEANT 2-R: 5'-GCA ACC TTG TGC CCT TTA-3'.</p> <p>The targeted sequences are in the plasmid pEA29. Stöger <i>et al.</i> (2006) recommended this method be used with DNA extracted using the REExtract-N-AmpT Kit (Sigma-Aldrich). The PCR mixture is composed of: ultrapure water, 5 µl; REExtract-N-Amp PCR Ready Mix, 10 µl; PEANT 1-F 10 pmol/µl, 0.5 µl; PEANT 2-R 10 pmol/µl, 0.5 µl; extracted DNA, 4 µl. The cycling parameters are: 95 °C for 5 min followed by 35 cycles of 95 °C for 15 s, 58 °C for 30 s, and 72 °C for 45 s, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The expected amplicon is 391 bp.-</p> <p>The accuracy in the 2009 ring test was 0.76 and in the 2010 ring test 0.72.</p> <p>PCR according to Gottsberger adapted from Obradovic <i>et al.</i> (2007)</p> <p>The primers are:</p> <p>FER1-F: 5'-AGC AGC AAT TAA TGG CAA GTA TAG TCA-3'</p> | | |

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| | | | <p>rgER2-R: 5'-AAA AGA GAC ATC TGG ATT CAG ACA AT-3'.</p> <p>The targeted sequences are chromosomal. The mixture is composed of: ultrapure water, 14.3 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 0.75 µl; dNTPs 10 mM, 0.25 µl; FER1-F 10 pmol/µl, 1 µl; rgER2-R 10 pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.2 µl; extracted DNA, 5 µl. The cycling parameters are: 94 °C for 3 min followed by 41 cycles of 94 °C for 10 s, 60 °C for 10 s and 72 °C for 30 s, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The expected amplicon size is 458 bp.</p> <p>The accuracy in the 2009 ring test was 0.76, and in the 2010 ring test 0.68, using the DNA extraction method described by Llop <i>et al.</i> (1999).</p> <p>Nested PCR according to Llop <i>et al.</i> (2000)</p> <p>The nested-PCR of Llop <i>et al.</i> (2000) uses two sets of primers, which are combined in a single reaction tube. Because of the different annealing temperatures the two PCRs are run consecutively. The external primers are those designed by McManus and Jones (1995) and are based on sequences of the pEA29 plasmid. The internal primers are those described by Llop <i>et al.</i> (2000). The sequences are the following:</p> <p>External primers:</p> <p>Forward primer: AJ75: 5'-CGT ATT CAC GGC TTC GCA GAT-3'</p> <p>Reverse primer: AJ76: 5'-ACC CGC CAG GAT AGT CGC ATA-3'</p> <p>Internal primers:</p> <p>Forward primer: PEANT1: 5'-TAT CCC TAA AAA CCT CAG TGC-3'</p> <p>Reverse primer: PEANT2: 5'-GCA ACC TTG TGC CCT TTA-3'.</p> <p>The PCR mixture is composed of: ultrapure water, 36.25 µl; buffer 10x, 5 µl; MgCl₂ 50 mM, 3 µl; dNTPs 10 mM, 0.5 µl; AJ75 0.1 pmol/µl, 0.32 µl; AJ76 0.1 pmol/µl, 0.32 µl; PEANT1 10 pmol/µl, 1 µl; primer PEANT2 10 pmol/µl, 1 µl; Taq DNA</p> | | |

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| | | | <p>polymerase 5 U/μl, 0.6 μl. A DNA sample volume of 2 μl should be added to 48 μl PCR mix. The cycling parameters are: a denaturation step of 94 °C for 4 min followed by 25 cycles of 94 °C for 60 s and 72 °C for 90 s. This first round PCR is followed in the same thermocycler by a second denaturation step of 94 °C for 4 min and 40 cycles of 94 °C for 60 s, 56 °C for 60 s, and 72 °C for 60 s, with a final elongation step at 72 °C for 10 min. The expected amplicon size is 391 bp, although variations in size can occur.</p> <p>The accuracy was 0.69 and 0.72 in the 2003 and 2010 ring tests, respectively, but increased after enrichment up to 0.84 (King's B medium) and 0.86 (CCT medium) in the 2003 ring test, and to 0.79 (King's B) and 0.88 (CCT) in 2010.</p> | | |
| 28. | 41 | Technical | <p>There are many PCR primers and protocols described for <i>E. amylovora</i> detection and some have shown specificity problems (Powney <i>et al.</i>, 2011a; Roselló <i>et al.</i>, 2006). The primers and protocols validated in ring tests were those of Bereswill <i>et al.</i> (1992) and Llop <i>et al.</i> (2000), with or without previous enrichment, in 2003; and those of Taylor <i>et al.</i> (2001), Stöger <i>et al.</i> (2006) and Obradovic <i>et al.</i> (2007) in 2009 and 2010. The discovery of fully virulent <i>E. amylovora</i> strains without the pEA29 plasmid (Llop <i>et al.</i>, 2006) and experiences from different countries (Powney <i>et al.</i>, 2011a) indicate that two PCR protocols should be used: one with primers based on pEA29 sequences, and another targeting unique chromosomal sequences. PCR can be carried out using the primers and conditions validated in the ring tests, although amplification conditions should be optimized for different thermo cyclers.</p> <p>PCR according to Bereswill <i>et al.</i> (1992)The primers are: Forward primer: A: 5'-CGG TTT TTA ACG CTG GG-3'Reverse primer: B: 5'-GGG CAA ATA CTC GGA TT-3'</p> <p>The targeted sequences are in the plasmid pEA29. The PCR mixture is composed of: ultrapure water, 17.4 μl; buffer 10x, 2.5 μl; MgCl₂ 50 mM, 1.5 μl; dNTPs 10 mM, 0.5 μl; primer A 10 pmol/μl, 0.25 μl; primer B 10 pmol/μl, 0.25 μl; Taq DNA polymerase 5 U/μl, 0.1 μl. The extracted DNA sample volume is 2.5 μl, and should be added to 22.5 μl of the PCR mix. The cycling parameters are: a denaturation step of 93 °C for 5 min followed by 40 cycles of 93 °C for 30 s, 52 °C for 30 s and 72 °C for 1 min 15 s, with a final elongation step at 72 °C for 10 min. The amplicon size is 900 base pairs (bp) according to Bereswill <i>et al.</i> (1992), although variations in size can occur between 900 and 1 100 bp depending on the number of 8 bp repeats</p> | Some strains of <i>Erwinia amylovora</i> without the pEA29 plasmid still remain toxicity. So, there is false negative in the detection methods for Pea29 . | China |

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| | | | <p>within the amplified fragment (Jones and Geider, 2001).</p> <p>The accuracy in the 2004 ring test was 0.51 but increased to 0.74 and 0.78 after enrichment of the samples in King's B and CCT media, respectively (López <i>et al.</i>, 2006).</p> <p>PCR according to Taylor <i>et al.</i> (2001)</p> <p>The primers are:</p> <p>G1-F: 5'-CCT GCA TAA ATC ACC GCT GAC AGC TCA ATG-3'</p> <p>G2-R: 5'-GCT ACC ACT GAT CGC TCG AAT CAA ATC GGC-3'.</p> <p>The targeted sequences are chromosomal. The PCR mixture is composed of: ultrapure water, 14.3 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 0.75 µl; dNTPs 10 mM, 0.25 µl; G1-F 10pmol/µl, 1 µl; G2-R 10pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.2 µl. An extracted DNA sample of 5 µl is added to 45 µl PCR mix. The cycling parameters are: 95 °C for 3 min followed by 40 cycles of 94 °C for 30 s, 60 °C for 30 s and 72 °C for 1 min, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The amplicon size is 187 bp.</p> <p>The accuracy in the 2010 ring test was 0.77 using the Llop <i>et al.</i> (1999) DNA extraction procedure.</p> <p>PCR according to Stöger <i>et al.</i> (2006)The primers (from Llop <i>et al.</i>, 2000) are:Forward primer: PEANT 1-F: 5'-TAT CCC TAA AAA CCT CAG TGC-3'</p> <p>Reverse primer: PEANT 2-R: 5'-GCA ACC TTG TGC CCT TTA-3'.The targeted sequences are in the plasmid pEA29. Stöger <i>et al.</i> (2006) recommended this method be used with DNA extracted using the REDExtract-N-AmpT Kit (Sigma-Aldrich). The PCR mixture is composed of: ultrapure water, 5 µl; REDExtract-N-Amp PCR Ready Mix, 10 µl; PEANT 1-F 10 pmol/µl, 0.5 µl; PEANT 2-R 10 pmol/µl, 0.5 µl; extracted DNA, 4 µl. The cycling parameters are: 95 °C for 5 min followed by 35 cycles of 95 °C for 15 s, 58 °C for 30 s, and 72 °C for 45 s, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The expected amplicon is 391 bp. The</p> | | |

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| | | | <p>accuracy in the 2009 ring test was 0.76 and in the 2010 ring test 0.72.</p> <p>PCR according to Gottsberger adapted from Obradovic <i>et al.</i> (2007)</p> <p>The primers are:</p> <p>FER1-F: 5'-AGC AGC AAT TAA TGG CAA GTA TAG TCA-3'</p> <p>rgER2-R: 5'-AAA AGA GAC ATC TGG ATT CAG ACA AT-3'.</p> <p>The targeted sequences are chromosomal. The mixture is composed of: ultrapure water, 14.3 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 0.75 µl; dNTPs 10 mM, 0.25 µl; FER1-F 10 pmol/µl, 1 µl; rgER2-R 10 pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.2 µl; extracted DNA, 5 µl. The cycling parameters are: 94 °C for 3 min followed by 41 cycles of 94 °C for 10 s, 60 °C for 10 s and 72 °C for 30 s, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The expected amplicon size is 458 bp.</p> <p>The accuracy in the 2009 ring test was 0.76, and in the 2010 ring test 0.68, using the DNA extraction method described by Llop <i>et al.</i> (1999).</p> <p>Nested PCR according to Llop <i>et al.</i> (2000)The nested-PCR of Llop <i>et al.</i> (2000) uses two sets of primers, which are combined in a single reaction tube. Because of the different annealing temperatures the two PCRs are run consecutively. The external primers are those designed by McManus and Jones (1995) and are based on sequences of the pEA29 plasmid. The internal primers are those described by Llop <i>et al.</i> (2000). The sequences are the following:</p> <p>External primers:Forward primer: AJ75: 5'-CGT ATT CAC GGC TTC GCA GAT-3'Reverse primer: AJ76: 5'-ACC CGC CAG GAT AGT CGC ATA-3'</p> <p>Internal primers:Forward primer:PEANT1: 5'-TAT CCC TAA AAA CCT CAG TGC-3'Reverse primer: PEANT2: 5'-GCA ACC TTG TGC CCT TTA-3'</p> <p>The PCR mixture is composed of: ultrapure water, 36.25 µl; buffer 10x, 5 µl; MgCl₂ 50 mM, 3 µl; dNTPs 10 mM, 0.5 µl; AJ75 0.1 pmol/µl, 0.32 µl; AJ76 0.1 pmol/µl, 0.32 µl; PEANT1 10 pmol/µl, 1 µl; primer PEANT2 10 pmol/µl, 1 µl; Taq DNA</p> | | |

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| | | | <p>polymerase 5 U/μl, 0.6 μl. A DNA sample volume of 2 μl should be added to 48 μl PCR mix. The cycling parameters are: a denaturation step of 94 °C for 4 min followed by 25 cycles of 94 °C for 60 s and 72 °C for 90 s. This first round PCR is followed in the same thermocycler by a second denaturation step of 94 °C for 4 min and 40 cycles of 94 °C for 60 s, 56 °C for 60 s, and 72 °C for 60 s, with a final elongation step at 72 °C for 10 min. The expected amplicon size is 391 bp, although variations in size can occur.</p> <p>The accuracy was 0.69 and 0.72 in the 2003 and 2010 ring tests, respectively, but increased after enrichment up to 0.84 (King's B medium) and 0.86 (CCT medium) in the 2003 ring test, and to 0.79 (King's B) and 0.88 (CCT) in 2010.</p> | | |
| 29. | 41 | Technical | <p>There are many PCR primers and protocols described for <i>E. amylovora</i> detection and some have shown specificity problems (Powney <i>et al.</i>, 2011a; Roselló <i>et al.</i>, 2006). The primers and protocols validated in ring tests were those of Bereswill <i>et al.</i> (1992) and Llop <i>et al.</i> (2000), with or without previous enrichment, in 2003; and those of Taylor <i>et al.</i> (2001), Stöger <i>et al.</i> (2006) and Obradovic <i>et al.</i> (2007) in 2009 and 2010. The discovery of fully virulent <i>E. amylovora</i> strains without the pEA29 plasmid (Llop <i>et al.</i>, 2006) and experiences from different countries (Powney <i>et al.</i>, 2011a) indicate that two PCR protocols should be used: one with primers based on pEA29 sequences, and another targeting unique chromosomal sequences. PCR can be carried out using the primers and conditions validated in the ring tests, although amplification conditions should be optimized for different thermo cyclers.</p> <p>PCR according to Bereswill <i>et al.</i> (1992)</p> <p>The primers are: Forward primer: A: 5'-CGG TTT TTA ACG CTG GG-3'</p> <p>Reverse primer: B: 5'-GGG CAA ATA CTC GGA TT-3'.</p> <p>The targeted sequences are in the plasmid pEA29. The PCR mixture is composed of: ultrapure water, 17.4 μl; buffer 10x, 2.5 μl; MgCl₂ 50 mM, 1.5 μl; dNTPs 10 mM, 0.5 μl; primer A 10 pmol/μl, 0.25 μl; primer B 10 pmol/μl, 0.25 μl; Taq DNA polymerase 5 U/μl, 0.1 μl. The extracted DNA sample volume is 2.5 μl, and should be added to 22.5 μl of the PCR mix. The cycling parameters are: a denaturation step of 93 °C for 5 min followed by 40 cycles of 93 °C for 30 s, 52 °C for 30 s and 72 °C for 1 min 15 s, with a final elongation step at 72 °C for 10 min. The amplicon size is</p> | Correct the reference to the ring test, it is 2003 not 2004. | EPPO, European Union, Georgia, Serbia |

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| | | | <p>900 base pairs (bp) according to Bereswill <i>et al.</i> (1992), although variations in size can occur between 900 and 1 100 bp depending on the number of 8 bp repeats within the amplified fragment (Jones and Geider, 2001).</p> <p>The accuracy in the 2003⁴ ring test was 0.51 but increased to 0.74 and 0.78 after enrichment of the samples in King's B and CCT media, respectively (López <i>et al.</i>, 2006).</p> <p>PCR according to Taylor <i>et al.</i> (2001)</p> <p>The primers are:</p> <p>G1-F: 5'-CCT GCA TAA ATC ACC GCT GAC AGC TCA ATG-3'</p> <p>G2-R: 5'-GCT ACC ACT GAT CGC TCG AAT CAA ATC GGC-3'</p> <p>The targeted sequences are chromosomal. The PCR mixture is composed of: ultrapure water, 14.3 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 0.75 µl; dNTPs 10 mM, 0.25 µl; G1-F 10pmol/µl, 1 µl; G2-R 10pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.2 µl. An extracted DNA sample of 5 µl is added to 45 µl PCR mix. The cycling parameters are: 95 °C for 3 min followed by 40 cycles of 94 °C for 30 s, 60 °C for 30 s and 72 °C for 1 min, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The amplicon size is 187 bp.</p> <p>The accuracy in the 2010 ring test was 0.77 using the Llop <i>et al.</i> (1999) DNA extraction procedure.</p> <p>PCR according to Stöger <i>et al.</i> (2006)</p> <p>The primers (from Llop <i>et al.</i>, 2000) are:</p> <p>Forward primer: PEANT 1-F: 5'-TAT CCC TAA AAA CCT CAG TGC-3'</p> <p>Reverse primer: PEANT 2-R: 5'-GCA ACC TTG TGC CCT TTA-3'.</p> | | |

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| | | | <p>The targeted sequences are in the plasmid pEA29. Stöger <i>et al.</i> (2006) recommended this method be used with DNA extracted using the REDExtract-N-AmpT Kit (Sigma-Aldrich). The PCR mixture is composed of: ultrapure water, 5 µl; REDExtract-N-Amp PCR Ready Mix, 10 µl; PEANT 1-F 10 pmol/µl, 0.5 µl; PEANT 2-R 10 pmol/µl, 0.5 µl; extracted DNA, 4 µl. The cycling parameters are: 95 °C for 5 min followed by 35 cycles of 95 °C for 15 s, 58 °C for 30 s, and 72 °C for 45 s, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The expected amplicon is 391 bp.</p> <p>The accuracy in the 2009 ring test was 0.76 and in the 2010 ring test 0.72.</p> <p>PCR according to Gottsberger adapted from Obradovic <i>et al.</i> (2007)</p> <p>The primers are:</p> <p>FER1-F: 5'-AGC AGC AAT TAA TGG CAA GTA TAG TCA-3'</p> <p>rgER2-R: 5'-AAA AGA GAC ATC TGG ATT CAG ACA AT-3'.</p> <p>The targeted sequences are chromosomal. The mixture is composed of: ultrapure water, 14.3 µl; buffer 10x, 2.5 µl; MgCl₂ 50 mM, 0.75 µl; dNTPs 10 mM, 0.25 µl; FER1-F 10 pmol/µl, 1 µl; rgER2-R 10 pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.2 µl; extracted DNA, 5 µl. The cycling parameters are: 94 °C for 3 min followed by 41 cycles of 94 °C for 10 s, 60 °C for 10 s and 72 °C for 30 s, with a final extension step at 72 °C for 5 min, and cooling at 15 °C. The expected amplicon size is 458 bp.</p> <p>The accuracy in the 2009 ring test was 0.76, and in the 2010 ring test 0.68, using the DNA extraction method described by Llop <i>et al.</i> (1999).</p> <p>Nested PCR according to Llop <i>et al.</i> (2000)</p> <p>The nested-PCR of Llop <i>et al.</i> (2000) uses two sets of primers, which are combined in a single reaction tube. Because of the different annealing temperatures the two PCRs are run consecutively. The external primers are those designed by McManus and Jones (1995) and are based on sequences of the pEA29 plasmid. The internal</p> | | |

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| | | | <p>primers are those described by Llop <i>et al.</i> (2000). The sequences are the following:</p> <p>External primers:</p> <p>Forward primer: AJ75: 5'-CGT ATT CAC GGC TTC GCA GAT-3'</p> <p>Reverse primer: AJ76: 5'-ACC CGC CAG GAT AGT CGC ATA-3'</p> <p>Internal primers:</p> <p>Forward primer: PEANT1: 5'-TAT CCC TAA AAA CCT CAG TGC-3'</p> <p>Reverse primer: PEANT2: 5'-GCA ACC TTG TGC CCT TTA-3'.</p> <p>The PCR mixture is composed of: ultrapure water, 36.25 µl; buffer 10×, 5 µl; MgCl₂ 50 mM, 3 µl; dNTPs 10 mM, 0.5 µl; AJ75 0.1 pmol/µl, 0.32 µl; AJ76 0.1 pmol/µl, 0.32 µl; PEANT1 10 pmol/µl, 1 µl; primer PEANT2 10 pmol/µl, 1 µl; Taq DNA polymerase 5 U/µl, 0.6 µl. A DNA sample volume of 2 µl should be added to 48 µl PCR mix. The cycling parameters are: a denaturation step of 94 °C for 4 min followed by 25 cycles of 94 °C for 60 s and 72 °C for 90 s. This first round PCR is followed in the same thermocycler by a second denaturation step of 94 °C for 4 min and 40 cycles of 94 °C for 60 s, 56 °C for 60 s, and 72 °C for 60 s, with a final elongation step at 72 °C for 10 min. The expected amplicon size is 391 bp, although variations in size can occur.</p> <p>The accuracy was 0.69 and 0.72 in the 2003 and 2010 ring tests, respectively, but increased after enrichment up to 0.84 (King's B medium) and 0.86 (CCT medium) in the 2003 ring test, and to 0.79 (King's B) and 0.88 (CCT) in 2010.</p> | | |
| 30. | 45 | Substantive | <p>Based on an evaluation of real-time PCR protocols in the ring tests in 2009 and 2010 (Dreo <i>et al.</i>, 2009; Lopez <i>et al.</i>, 2010) the protocol described by Pirc <i>et al.</i> (2009), which targets chromosomal sequences, was recommended. A duplex real-time PCR based on chromosomal sequences is also available but has not been ring tested (Lehman <i>et al.</i>, 2008).</p> | <p>To add footnote 3: The use of the brand Applied Biosystems for the TaqMan Fast Universal PCR Master Mix and analysers 7900HT and 7900HT Fast in this diagnostic protocol implies no approval of them to the exclusion of others that may also be suitable. This information is given for the convenience of users of this protocol and does not</p> | <p>COSAVE, Uruguay, Chile, Brazil, Peru, Argentina</p> |

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| | | | <p>Real-time PCR according to Pirc <i>et al.</i> (2009)</p> <p>The following oligonucleotides are used:</p> <p>Ams116F primer: 5'-TCC CAC ATA CTG TGA ATC CA-3'</p> <p>Ams189R primer: 5'-GGG TAT TTG CGC TAA TTT TAT TCG-3'</p> <p>Ams141T probe: FAM-CCA GAA TCT GGC CCG CGT ATA CCG-TAMRA.</p> <p>The reaction is carried out in a final volume of 25 µl. The PCR mixture is composed of: ultrapure water, 2.5 µl; 2x TaqMan Fast Universal PCR Master Mix (Applied Biosystems), 12.5 µl; Ams116F 10 pmol/µl, 2.25 µl; Ams189R 10 pmol/µl, 2.25 µl; FAM-labelled Ams141T 10 pmol/µl, 0.5 µl; 5 µl DNA extract added to 20 µl PCR mix. The cycling parameters are: 2 min at 50 °C; 10 min at 95 °C; and 40 cycles of 15 s at 95 °C and 1 min at 60 °C. The standard mode for temperature ramping rates on analysers 7900HT and 7900HT Fast (Applied Biosystems)¹ are: 1.6 °C/s up and 1.6 °C/s down. It is possible to run reactions at slower ramp rates, but with faster ramp rates (up and down at approximately 3.5 °C/s) the results were not acceptable. The expected amplicon size is 74 bp.</p> <p>For analysis, there are usually different options available for setting the signal and noise limits, automatic or manual. The instructions for analysis of the real-time PCR results of the appropriate software should be followed; the baseline should be set automatically, and the threshold should be set manually crossing the exponential phase of the control amplification curves. The accuracy in the 2010 ring test was 0.80, 0.85 and 0.76 with the DNA extraction method of Llop <i>et al.</i> (1999), REExtract-N-Amp Plant PCR Kit and Taylor <i>et al.</i> (2001), respectively.</p> <p>Real-time PCR according to Gottesberger (2010)</p> <p>The following oligonucleotides that target the <i>E. amylovora</i> chromosome are used:</p> <p>hpEaF primer: 5'-CCG TGG AGA CCG ATC TTT TA-3'</p> | <p>constitute an endorsement by the CPM of the chemical, reagent and/or equipment named. Equivalent products may be used if they can be shown to lead to the same results TaqMan Universal PCR Master Mix (Applied Biosystems), - related to footnote 3.</p> | |

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| | | | <p>hpEaR primer: 5'-AAG TTT CTC CGC CCT ACG AT-3'</p> <p>hpEaP probe: FAM-TCG TCG AAT GCT GCC TCT CT-MGB.</p> <p>The reaction is carried out in a final volume of 20 µl. The PCR mixture is composed of: ultrapure water, 6 µl; 2x TaqMan Universal PCR Master Mix (Applied Biosystems), 10 µl; hpEaF 10 pmol/µl, 1 µl; hpEaR 10 pmol/µl, 1 µl; hpEaP 10 pmol/µl, 1 µl; 1 µl DNA extract added to 19 µl PCR mix. The cycling parameters are: 2 min at 50 °C, 10 min at 95°C, and 50 cycles of 15 s at 95 °C and 1 min at 60°C. The expected amplicon size is 138 bp.</p> <p>For analysis, there are usually different options available for setting the signal and noise limits, automatic or manual. The instructions for analysis of the real-time PCR results of the appropriate software should be followed; the baseline should be set automatically, and the threshold should be set manually crossing the exponential phase of the control amplification curves. The accuracy of this real-time PCR could not be tested in the 2010 ring test; however, it was tested in parallel with the real-time PCR described in Pirc <i>et al.</i> (2009) by one laboratory and gave the same qualitative results with the DNA extraction from Llop <i>et al.</i> (1999).</p> | | |
| 31. | 45 | Technical | <p>Based on an evaluation of real-time PCR protocols in the ring tests in 2009 and 2010 (Dreo <i>et al.</i>, 2009; Lopez <i>et al.</i>, 2010) the protocol described by Pirc <i>et al.</i> (2009), which targets chromosomal sequences, was recommended. A duplex real-time PCR based on chromosomal sequences is also available but has not been ring tested (Lehman <i>et al.</i>, 2008).</p> <p>Real-time PCR according to Pirc <i>et al.</i> (2009)</p> <p>The following oligonucleotides are used:</p> <p>Ams116F primer: 5'-TCC CAC ATA CTG TGA ATC CA-3'</p> <p>Ams189R primer: 5'-GGG TAT TTG CGC TAA TTT TAT TCG-3'</p> <p>Ams141T probe: FAM-CCA GAA TCT GGC CCG CGT ATA CCG-TAMRA.</p> <p>The reaction is carried out in a final volume of 25 µl. The PCR mixture is composed</p> | Change in the spelling of a name, modification of the concentration in the solution of hpEaP from 10pmol/µl to 1pmol/µl. | EPPO, European Union, Georgia, Serbia |

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| | | | <p>of: ultrapure water, 2.5 µl; 2x TaqMan Fast Universal PCR Master Mix (Applied Biosystems), 12.5 µl; Ams116F 10 pmol/µl, 2.25 µl; Ams189R 10 pmol/µl, 2.25 µl; FAM-labelled Ams141T 10 pmol/µl, 0.5 µl; 5 µl DNA extract added to 20 µl PCR mix. The cycling parameters are: 2 min at 50 °C; 10 min at 95 °C; and 40 cycles of 15 s at 95 °C and 1 min at 60 °C. The standard mode for temperature ramping rates on analysers 7900HT and 7900HT Fast (Applied Biosystems)¹ are: 1.6 °C/s up and 1.6 °C/s down. It is possible to run reactions at slower ramp rates, but with faster ramp rates (up and down at approximately 3.5 °C/s) the results were not acceptable. The expected amplicon size is 74 bp.</p> <p>For analysis, there are usually different options available for setting the signal and noise limits, automatic or manual. The instructions for analysis of the real-time PCR results of the appropriate software should be followed; the baseline should be set automatically, and the threshold should be set manually crossing the exponential phase of the control amplification curves. The accuracy in the 2010 ring test was 0.80, 0.85 and 0.76 with the DNA extraction method of Llop <i>et al.</i> (1999), REDExtract-N-Amp Plant PCR Kit and Taylor <i>et al.</i> (2001), respectively.</p> <p>Real-time PCR according to Gottesberger (2010)</p> <p>The following oligonucleotides that target the <i>E. amylovora</i> chromosome are used:</p> <p>hpEaF primer: 5'-CCG TGG AGA CCG ATC TTT TA-3'</p> <p>hpEaR primer: 5'-AAG TTT CTC CGC CCT ACG AT-3'</p> <p>hpEaP probe: FAM-TCG TCG AAT GCT GCC TCT CT-MGB.</p> <p>The reaction is carried out in a final volume of 20 µl. The PCR mixture is composed of: ultrapure water, 6 µl; 2x TaqMan Universal PCR Master Mix (Applied Biosystems), 10 µl; hpEaF 10 pmol/µl, 1 µl; hpEaR 10 pmol/µl, 1 µl; hpEaP 10 pmol/µl, 1 µl; 1 µl DNA extract added to 19 µl PCR mix. The cycling parameters are: 2 min at 50 °C, 10 min at 95°C, and 50 cycles of 15 s at 95 °C and 1 min at 60°C. The expected amplicon size is 138 bp.</p> <p>For analysis, there are usually different options available for setting the signal and</p> | | |

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| | | | noise limits, automatic or manual. The instructions for analysis of the real-time PCR results of the appropriate software should be followed; the baseline should be set automatically, and the threshold should be set manually crossing the exponential phase of the control amplification curves. The accuracy of this real-time PCR could not be tested in the 2010 ring test; however, it was tested in parallel with the real-time PCR described in Pirc <i>et al.</i> (2009) by one laboratory and gave the same qualitative results with the DNA extraction from Llop <i>et al.</i> (1999). | | |
| 32. | 55 | Substantive | <p>The LAMP protocol was developed and described by Temple <i>et al.</i> (2008) and Temple and Johnson (2011). It was evaluated in the 2010 ring test because it was considered appropriate for laboratories not equipped for PCR and it is easy to perform. In the ring test, the LAMP protocol using primers to detect the chromosomal gene <i>amsL</i> of <i>E. amylovora</i> and was found to lack appropriate sensitivity for analysis of samples with low bacterial populations. Consequently, the LAMP protocol described below to detect chromosomal <i>amsL</i> is recommended only for the analysis of symptomatic samples with more than 10^5–10^6 c.f.u./ml. The protocol from Temple and Johnson (2011) using primers to detect pEA29 was not evaluated in the ring test.</p> <p>LAMP primers to detect <i>amsL</i> B are:</p> <p>ALB Fip: 5'-CTG CCT GAG TAC GCA GCT GAT TGC ACG TTT TAC AGC TCG CT-3'</p> <p>ALB Bip: 5'-TCG TCG GTA AAG TGA TGG GTG CCC AGC TTA AGG GGC TGA AG-3'</p> <p>ALB F: 5'-GCC CAC ATT CGA ATT TGA CC-3'</p> <p>ALB B: 5'-CGG TTA ATC ACC GGT GTC A-3'.</p> <p>Primers Fip and Bip were used at 2.4 μM and primers F and B at 0.2 μM final concentrations. Melting temperatures for primers were between 58 and 60 °C. The LAMP reaction mixture is composed of: 10x ThermoPol buffer (New England Biolabs), 5 μl; dNTPs 10 mM, 5 μl; MgSO₄ 100 mM, 2 μl; bovine serum albumin (BSA) 10 mg/ml, 2 μl; ALB FIP 100 μM, 1.2 μl; ALB BIP 100 μM, 1.2 μl; ALB F 10 μM, 1 μl; ALB B 10 μM, 1 μl; <i>Bst</i> DNA polymerase 8 U/μl, 2 μl; sample, 5 μl; ultrapure water, 24.6 μl. Note that the <i>Bst</i> DNA polymerase, water and template</p> | to add footnote 4: The use of the brand New England Biolabs for the ThermoPol buffer in this diagnostic protocol implies no approval of them to the exclusion of others that may also be suitable. This information is given for the convenience of users of this protocol and does not constitute an endorsement by the CPM of the chemical, reagent and/or equipment named. Equivalent products may be used if they can be shown to lead to the same results | COSAVE, Uruguay, Chile, Brazil, Peru, Argentina |

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| | | | <p>DNA are not added to the master mix, but are added separately after aliquoting the master mix. Before starting the LAMP reaction, a water bath or a thermal cycler is set at 65 °C. The mix is prepared and 18.4 µl of the master mix is pipetted into each individual 0.2 ml PCR tube. The <i>Bst</i> DNA polymerase, template DNA and ultrapure water are then pipetted separately into each tube with master mix. The tubes are spun down in a plate spinner (1000 r.p.m. for 30 s) and then are placed in the water bath (65 °C) in a holder so the reaction end is submerged, or in the thermocycler (65 °C) for 55 min. The tubes are removed and allowed to cool for 10 s.</p> <p>The test on a sample is positive if the presence of precipitate as cloudiness in the tube or the presence of a solid white magnesium pyrophosphate precipitate at the bottom of the tube is observed, as for the positive control. A clear solution indicates a negative test result, as should be observed for the negative control.</p> <p>The accuracy in the 2010 ring test was 0.64, but for samples with 10⁵–10⁶ c.f.u./ml the accuracy was 0.80. For this reason LAMP is recommended only for the analysis of symptomatic samples.</p> | | |
| 33. | 58 | Editorial | Asymptomatic samples can be processed individually (preferred) or in groups of up to 100 (EPPO/OEPP, 2013 1992). Precautions to avoid cross-contamination should be taken when collecting the samples and during the extraction process. Sampling and sample preparation can be performed following one of these protocols: | Cf. second reference of paragraph [121]. EPPO 1992 was withdrawn in 2012. Appendix I of PM 7/ 020 gives guidance on preparation of asymptomatic material. Therefore replace the reference to EPPO (2013). | EPPO, European Union, Georgia, Serbia |
| 34. | 63 | Editorial | <p>Direct analysis of asymptomatic samples is normally negative for <i>E. amylovora</i> because of the low bacterial population. Consequently, when analysing asymptomatic material, it is an absolute requirement to perform enrichment from samples prepared in the antioxidant buffer (described above; Gorris <i>et al.</i>, 1996a) for 72 h at approximately 25 °C. The recommended screening tests are indicated in the flow diagram in Figure 2.</p> <p>It is advisable to perform at least two of these screening tests based on different biological principles:</p> <p><u>Move the last sentence "The recommended screening tests are indicated in the flow diagram in Figure 2." to the front of Para..57.</u></p> | It's more orderly. | China |

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| 35. | 63 | Editorial | <p>Direct analysis of asymptomatic samples is normally negative for <i>E. amylovora</i> because of the low bacterial population. Consequently, when analysing asymptomatic material, it is an absolute requirement to perform enrichment from samples prepared in the antioxidant buffer (described above; Gorris <i>et al.</i>, 1996^a) for 72 h at approximately 25 °C. The recommended screening tests are indicated in the flow diagram in Figure 2.</p> <p>It is advisable to perform at least two of these screening tests based on different biological principles:</p> | This is no "Gorris <i>et al.</i> , 1996b" so the "a" is not needed (cf. [122]). | EPPO, European Union, Georgia, Serbia |
| 36. | 64 | Substantive | <p>1. Enrichment-isolation. Follow the procedure for symptomatic samples (section 3.1.3.2).</p> | Shouldn't immunofluoresce be added? Because it has been mentioned in the point 3.1.2. | COSAVE, Uruguay, Chile, Brazil, Peru, Argentina |
| 37. | 67 | Technical | If any of the screening tests are positive, isolation of the pathogen from the extract stored at –80 °C with glycerol or from the enriched samples should be attempted. When three tests or more are positive and the isolation is negative, it is reasonable to strongly suspect the presence of <i>E. amylovora</i> in the sample, but identification requires isolation of the pathogen from new samples and subsequent identification of the bacterium. | Second sentence "it is reasonable to strongly suspect the presence": We agree. But this is not consistent with the Flow chart Figure 2 where the diagnosis is "not detected". Consider whether there is a need to refer to strong suspicion and need for isolation from new samples in the flow diagram. For example as a footnote. | EPPO, European Union, Georgia, Serbia |
| 38. | 68 | Editorial | 4. Identification | The number of the section is missing. | EPPO, European Union, Georgia, Serbia |
| 39. | 68 | Editorial | 4. Identification | Editorial correction | Japan |
| 40. | 69 | Editorial | Identification should be based on results obtained from several techniques because other species of <i>Erwinia</i> such as <i>E. piriflorinigrans</i> (López <i>et al.</i> , 2011), <i>E. pyrifoliae</i> (Kim <i>et al.</i> , 1999; Rhim <i>et al.</i> , 1999), <i>E. uzenensis</i> (Matsuura <i>et al.</i> , 2012) and other <i>Erwinia</i> spp. (Kim <i>et al.</i> , 2001a, 2001b; Palacio-Bielsa <i>et al.</i> , 2012) that have relatively similar morphology to that of <i>E. amylovora</i> share some serological and molecular characteristics. Differentiation of <i>E. amylovora</i> from these closely related <i>Erwinia</i> species (that can be found in similarly symptomatic tissues in some hosts) can be achieved with a combination of three techniques based on different biological principles: (1) PCR based on chromosomal DNA (sections 3.1.5.2 and 4.3.1); (2) DAS-ELISA using specific monoclonal antibodies as described for detection (see | <p>1. Third paragraph: current name of institute.</p> <p>2. Codes are not used for "France", "Belgium", "New Zealand" and "United States". 3. A comma is missing after "(INRA)".</p> | EPPO, European Union, Georgia, Serbia |

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| | | | <p>section 3.1.4.1, excluding the enrichment step); and (3) inoculation into fire blight hosts to fulfil the requirements of Koch's postulates, including re-isolation of the inoculated pathogen (section 4.4).</p> <p>For identification of colonies, at least two of these three techniques are recommended to be performed. Other tests can also be used depending on the experience of the laboratory and these are described below. When required, the final confirmation of a culture's identification should include a pathogenicity test.</p> <p>The <i>E. amylovora</i> isolates recommended for use as positive controls are NCPPB683 and CFBP 1430. The following collections, among others, can provide different <i>E. amylovora</i> reference strains: National Collection of Plant Pathogenic Bacteria (NCPBP), Fera Central Science Laboratory, York, United Kingdom; Collection Française de Bactéries Phytopathogènes (CFBP), French National Institute for Agricultural Research (INRA), Station Phytobactériologie, Angers, France; Belgian Co-ordinated Collection of Microorganisms BCCM/LMG Bacteria Collection, Gent, Belgium; The International Collection of Microorganisms from Plants, Manaaki Whenua Landcare Research, Auckland, New Zealand; and The American Type Culture Collection (ATTC), Manassas, VA, United States. The authenticity of the strains can be guaranteed only if directly obtained from the culture collections.</p> | | |
| 41. | 69 | Substantive | <p>Identification should be based on results obtained from several techniques because other species of <i>Erwinia</i> such as <i>E. piriflorinigra</i>s (López <i>et al.</i>, 2011), <i>E. pyrifoliae</i> (Kim <i>et al.</i>, 1999; Rhim <i>et al.</i>, 1999), <i>E. uzenensis</i> (Matsuura <i>et al.</i>, 2012) and other <i>Erwinia</i> spp. (Kim <i>et al.</i>, 2001a, 2001b; Palacio-Bielsa <i>et al.</i>, 2012) that have relatively similar morphology to that of <i>E. amylovora</i> share some serological and molecular characteristics. Differentiation of <i>E. amylovora</i> from these closely related <i>Erwinia</i> species (that can be found in similarly symptomatic tissues in some hosts) can be achieved with a combination of three techniques based on different biological principles: (1) Nutritional and enzymic identification (section 4.1) (42) PCR based on chromosomal DNA (sections 3.1.5.2 and 4.3.1); (23) DASI-ELISA using specific monoclonal antibodies as described for detection (see section 3.1.4.1, excluding the enrichment step); and (3) inoculation into fire blight hosts to fulfil the requirements of Koch's postulates, including re-isolation of the inoculated pathogen (section 4.4).</p> <p>For identification of colonies, at least two of these three techniques are recommended to be performed. Other tests can also be used depending on the experience of the laboratory and these are described below. When required, the</p> | <p>Three techniques indicate nutritional and enzymic identification, molecular identification and serological identification. The pathogenicity test in section 4.4 is an independent phase in the Fig.1 so it is not included among the three techniques.</p> | Japan |

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| | | | <p>final confirmation of a culture's identification should include a pathogenicity test.</p> <p>The <i>E. amylovora</i> isolates recommended for use as positive controls are NCPPB683 and CFBP 1430. The following collections, among others, can provide different <i>E. amylovora</i> reference strains: National Collection of Plant Pathogenic Bacteria (NCPPB), Central Science Laboratory, York, UK; Collection Française de Bactéries Phytopathogènes (CFBP), French National Institute for Agricultural Research (INRA) Station Phytobactériologie, Angers, France; Belgian Co-ordinated Collection of Microorganisms BCCM/LMG Bacteria Collection, Gent, Belgium; The International Collection of Microorganisms from Plants, Manaaki Whenua Landcare Research, Auckland, New Zealand; and The American Type Culture Collection (ATTC), Manassas, VA, United States. The authenticity of the strains can be guaranteed only if directly obtained from the culture collections.</p> | | |
| 42. | 80 | Editorial | <p>Identification of <i>E. amylovora</i> can be obtained biochemically by profiling on the API system 20 E and 50 CH strips (bioMérieux²¹).</p> <p>API 20 E. The manufacturer's instructions should be followed for preparing the suspension and inoculating the strip. The strip is incubated at 25–26°C. The readings after 48 h should be as indicated for a typical <i>E. amylovora</i> culture: the tests lysine decarboxylase (LDC), ornithine decarboxylase (ODC), citrate utilization (CIT), H₂S production (SH₂), urease (URE), tryptophan deaminase (TDA), indole production (IND), and rhamnose oxidation (RHA) should be negative, while sucrose oxidation (SAC) should be positive. Other tests may vary by strain, according to Donat <i>et al.</i> (2007).</p> <p>API 50 CH. A suspension of OD 1.0 (at 600 nm wavelength) is prepared in PBS. One millilitre of the suspension is added to 20 ml Ayers medium (NH₄H₂PO₄, 1 g; KCl, 0.2 g; MgSO₄, 0.2 g; bromothymol blue 0.2%, 75 ml; distilled water, 1 litre; pH 7; sterilized at 120 °C for 20 min) (Ayers <i>et al.</i>, 1919). The manufacturer's instructions should be followed for inoculation of the strip. The strip is incubated at 25–26 °C under aerobic conditions. Utilization of the different carbohydrates is observed by development of a yellow colour in the well. The reading after 72 h for a typical <i>E. amylovora</i> culture on API 50 CH should be: l-arabinose, ribose, d-glucose, d-fructose, mannitol, sorbitol, N-acetylglucosamine, sucrose, trehalose and β-gentiobiose positive. The remaining sugars are not utilized by <i>E. amylovora</i> in these conditions, but some strains can utilize glycerol and d-fucose, according to Donat <i>et al.</i> (2007).</p> | Note 1 says see footnote 10 which does not exist. Should it be deleted and note 2 should become note 1. | EPPO, European Union, Georgia, Serbia |

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|----------|----------|--------------|---|---|---|
| 43. | 80 | Substantive | <p>Identification of <i>E. amylovora</i> can be obtained biochemically by profiling on the API system 20 E and 50 CH strips (bioMérieux²).</p> <p>API 20 E. The manufacturer's instructions should be followed for preparing the suspension and inoculating the strip. The strip is incubated at 25–26°C. The readings after 48 h should be as indicated for a typical <i>E. amylovora</i> culture: the tests lysine decarboxylase (LDC), ornithine decarboxylase (ODC), citrate utilization (CIT), H₂S production (SH₂), urease (URE), tryptophan deaminase (TDA), indole production (IND), and rhamnose oxidation (RHA) should be negative, while sucrose oxidation (SAC) should be positive. Other tests may vary by strain, according to Donat <i>et al.</i> (2007).</p> <p>API 50 CH. A suspension of OD 1.0 (at 600 nm wavelength) is prepared in PBS. One millilitre of the suspension is added to 20 ml Ayers medium (NH₄H₂PO₄, 1 g; KCl, 0.2 g; MgSO₄, 0.2 g; bromothymol blue 0.2%, 75 ml; distilled water, 1 litre; pH 7; sterilized at 120 °C for 20 min) (Ayers <i>et al.</i>, 1919). The manufacturer's instructions should be followed for inoculation of the strip. The strip is incubated at 25–26 °C under aerobic conditions. Utilization of the different carbohydrates is observed by development of a yellow colour in the well. The reading after 72 h for a typical <i>E. amylovora</i> culture on API 50 CH should be: l-arabinose, ribose, d-glucose, d-fructose, mannitol, sorbitol, N-acetylglucosamine, sucrose, trehalose and β-gentiobiose positive. The remaining sugars are not utilized by <i>E. amylovora</i> in these conditions, but some strains can utilize glycerol and d-fucose, according to Donat <i>et al.</i> (2007).</p> | According to previous comments this would be footnote 5. In this point say: the citrate utilization (CIT) should be negative, but in 4.1 (Nutritional and enzymatic identification) say: citrate utilization (+). It should be clarified. | COSAVE, Uruguay, Chile, Brazil, Peru, Argentina |
| 44. | 87 | Editorial | Suspected <i>E. amylovora</i> colonies can be presumptively identified by slide agglutination. A dense suspension of cells is mixed with a drop of PBS and a drop of <i>E. amylovora</i> specific antiserum (undiluted, or at 1:5 to 1:10 dilution only) on a slide. Monoclonal antibodies can be used providing they agglutinate the reference strains. The specificity of the antibodies must be established in advance. | The second occurrence of "E. amylovora" should be in italics. | EPPO, European Union, Georgia, Serbia |
| 45. | 89 | Editorial | A suspension of approximately 10 ⁶ cells/ml is prepared in PBS from levan-positive, non-fluorescent colonies and the IF procedure described in section 3.1.4.3 is applied. The specificity of the antibodies must be established in advance. | Editorial correction | Japan |
| 46. | 102 | Editorial | Suspected <i>E. amylovora</i> colonies should be inoculated back into host plants to fulfil Koch's postulates and verify their pathogenicity. For plant inoculation, susceptible cultivars of pear, apple, loquat, <i>Crataegus</i> , <i>Cotoneaster</i> or <i>Pyracantha</i> spp. are used. Young shoots are inoculated by cutting across a young leaf through the central vein with scissors dipped in a 10 ⁹ c.f.u./ml suspension of each isolate prepared in PBS. The plants are maintained at 20–25 °C at approximately 80% | Second paragraph: the paragraph is about "detached immature fruits". | EPPO, European Union, Georgia, Serbia |

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| | | | <p>relative humidity for one to two weeks. Detached young shoots that have been surface-sterilized (treated with 70% ethanol for 30 s then washed three times with sterile distilled water) from greenhouse-grown plants can also be inoculated in the same way and kept in tubes with sterile 1% agar. The tubes should be kept at 20–25 °C with 16 h light per day.</p> <p>Inoculation can also be performed on detached immature fruits of susceptible cultivars of pear, apple and loquat by placing 10 µl of 10⁹ c.f.u./ml suspensions of the isolates in PBS into a fresh wound on the shoot or on the surface of disinfected fruits (treated with 70% commercial chlorine for 30 min then washed three times with sterile distilled water). They should be incubated in a humid chamber at 25 °C for 3–5 days.</p> <p><i>E. amylovora</i>-like colonies are re-isolated and characterized from inoculated organs showing typical fire blight symptoms. A positive test is evident by the oozing of bacteria and browning around the inoculation site after 2–7 days, as seen in the positive <i>E. amylovora</i> control, providing no lesions are or only a small necrotic lesion is observed at the wound site in the negative control.</p> <p>Other inoculation techniques are possible. Hypersensitive reactions in tobacco leaves may indicate expression of the <i>hrp</i> genes of <i>E. amylovora</i>, but this test may be positive for many other plant pathogenic bacteria. Tobacco plants of cv. Xanthi or Samsun with more than five to six leaves should be used. Bacterial suspensions of 10⁹ c.f.u./ml (OD at 600 nm, 1.0) are prepared and a needle and syringe used to inject the suspensions into the intracellular space of mature leaves. Complete collapse of the infiltrated tissue after 24-48 h at room temperature is recorded as positive, as observed in the positive <i>E. amylovora</i> control.</p> | | |
| 47. | 102 | Substantive | <p>Suspected <i>E. amylovora</i> colonies should be inoculated back into host plants to fulfil Koch's postulates and verify their pathogenicity. For plant inoculation, susceptible cultivars of pear, apple, loquat, <i>Crataegus</i>, <i>Cotoneaster</i> or <i>Pyracantha</i> spp. are used. Young shoots are inoculated by cutting across a young leaf through the central vein with scissors dipped in a 10⁹ c.f.u./ml suspension of each isolate prepared in PBS. The plants are maintained at 20–25 °C at approximately 80% relative humidity for one to two weeks. Detached young shoots that have been surface-sterilized (treated with 70% ethanol for 30 s then washed three times with sterile distilled water) from greenhouse-grown plants can also be inoculated in the same way and kept in tubes with sterile 1% agar. The tubes should be kept at 20–</p> | <p>Add the example of susceptible cultivars of each plant. This information is useful for countries without <i>Erwinia amylovora</i></p> | Japan |

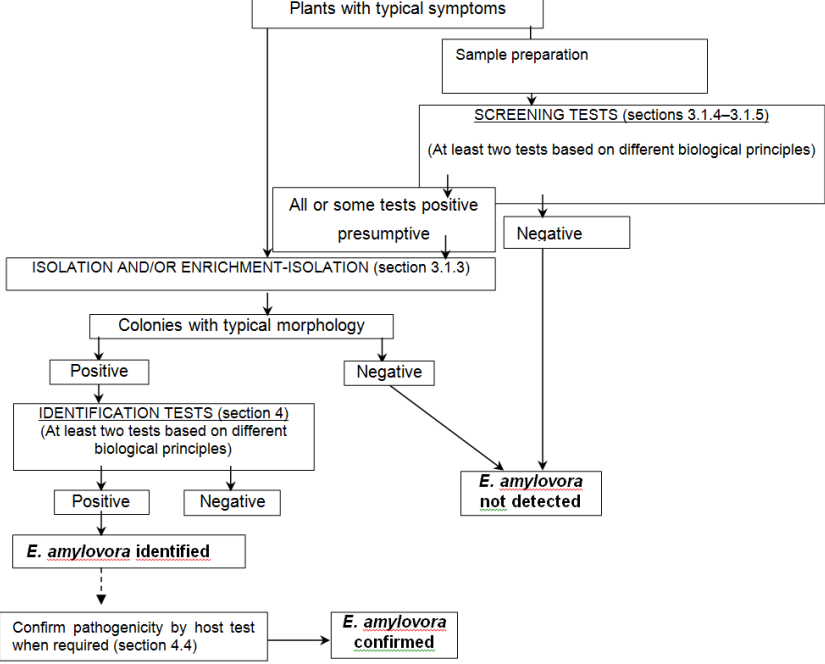
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| | | | <p>25 °C with 16 h light per day.</p> <p>Inoculation can also be performed on detached immature fruits of susceptible cultivars of pear, apple and loquat by placing 10 µl of 10⁹ c.f.u./ml suspensions of the isolates in PBS into a fresh wound on the shoot or on the surface of disinfected fruits (treated with 70% commercial chlorine for 30 min then washed three times with sterile distilled water). They should be incubated in a humid chamber at 25 °C for 3–5 days.</p> <p><i>E. amylovora</i>-like colonies are re-isolated and characterized from inoculated organs showing typical fire blight symptoms. A positive test is evident by the oozing of bacteria and browning around the inoculation site after 2–7 days, as seen in the positive <i>E. amylovora</i> control, providing no lesions are or only a small necrotic lesion is observed at the wound site in the negative control.</p> <p>Other inoculation techniques are possible. Hypersensitive reactions in tobacco leaves may indicate expression of the <i>hrp</i> genes of <i>E. amylovora</i>, but this test may be positive for many other plant pathogenic bacteria. Tobacco plants of cv. Xanthi or Samsun with more than five to six leaves should be used. Bacterial suspensions of 10⁹ c.f.u./ml (OD at 600 nm, 1.0) are prepared and a needle and syringe used to inject the suspensions into the intracellular space of mature leaves. Complete collapse of the infiltrated tissue after 24-48 h at room temperature is recorded as positive, as observed in the positive <i>E. amylovora</i> control.</p> | | |
| 48. | 103 | Editorial | 5. Records | The number of the section is missing. | EPPO, European Union, Georgia, Serbia |
| 49. | 105 | Editorial | 6. Contact Points for Further Information | The number of the section is missing. | EPPO, European Union, Georgia, Serbia |
| 50. | 108 | Editorial | Plant Health and Environment Laboratory, Investigation and Diagnostic Centres, Ministry for Primary Industries, 231 Morrin Road, St Johns, Auckland 1140, New Zealand (Robert Taylor; e-mail: Robert.Taylor@mpi.govt.nz ; tel.: +64 99093548; fax: +64 99095739). | The information is missing for Mr Rodney Roberts (cf. paragraph [2], section "Consultation on technical level", and paragraph [110]). | EPPO, European Union, Georgia, Serbia |

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| 51. | 109 | Editorial | 7. Acknowledgements | The number of the section is missing. | EPPO, European Union, Georgia, Serbia |
| 52. | 111 | Editorial | 8. References | The number of the section is missing. | EPPO, European Union, Georgia, Serbia |
| 53. | 121 | Editorial | <p>EPPO/OEPP (European and Mediterranean Plant Protection Organization). 2013. PM 7/20 (2) <i>Erwinia amylovora</i>. EPPO Bulletin, 43: 21–45. doi: 10.1111/epp.12019(1992) Quarantine procedure no.40. <i>Erwinia amylovora</i>. Sampling and test methods. Bull. OEPP 22: 225–231.</p> <p>EPPO (European and Mediterranean Plant Protection Organization). 2013. EPPO Plant Quarantine Data Retrieval System (Available online at http://www.eppo.org/DATABASES/pqr/pqr.htm)</p> | Two references are merged. The 1992 standard has been withdrawn. Current reference added. | EPPO, European Union, Georgia, Serbia |
| 54. | 123 | Editorial | <p>Gottsberger, R.A. (2010). Development and evaluation of a real-time PCR assay targeting chromosomal DNA of <i>Erwinia amylovora</i>. <i>Letters in Applied Microbiology</i>, 51 (2010): 285–292.</p> <p>Gottsberger, R.A. (2010). Development and evaluation of a real-time PCR assay targeting chromosomal DNA of <i>Erwinia amylovora</i>. <i>Letters in Applied Microbiology</i>, 51 (2010): 285–292.</p> | This reference should be corrected (bold, italics...). | EPPO, European Union, Georgia, Serbia |
| 55. | 134 | Editorial | López, M.M., Llop, P., Gorrís, M.T., Keck, M., Peñalver, J., Donat, V. & Cambra, M. 2006. European protocol for diagnosis of <i>Erwinia amylovora</i> . <i>Acta Horticulturae</i> , 704: 99–103. | Paragraphs [134], [135] and [136] should be after paragraph [139] (alphabetical order of the authors). | EPPO, European Union, Georgia, Serbia |
| 56. | 135 | Editorial | López, M.M., Peñalver, J., Arilla, A., Morente, C., Dreó, T., Pirc, M., Poliakoff, F., Dousset, C., Visage, M., Achbani, E., Bersgma-Vlami, M., Drenova, N., Duffy, B., Marín, M., Meeke, E., Moumni, M., Obradovic, A., Palomo, J., Taylor, R., Stockwell, V. & Reisenzein, H. 2010. Ring test evaluation of techniques for <i>Erwinia amylovora</i> diagnosis and detections. 12th International Workshop on Fire Blight. Warsaw, Poland. Abstract 18. | Paragraphs [134], [135] and [136] should be after paragraph [139] (alphabetical order of the authors). | EPPO, European Union, Georgia, Serbia |
| 57. | 136 | Editorial | López, M.M., Roselló, M.M., Llop, P., Ferrer, S., Christen, R. & Gardan, L. 2011. <i>Erwinia piriflorinigrans</i> sp. nov., a novel pathogen that causes necrosis of pear blossoms. <i>International Journal of Systemic and Evolutionary Microbiology</i> , 61: 561–567. | Paragraphs [134], [135] and [136] should be after paragraph [139] (alphabetical order of the authors). | EPPO, European Union, Georgia, Serbia |

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| 58. | 137 | Editorial | Llop, P., Caruso, P., Cubero, J., Morente, C. & López, M.M. 1999. A simple extraction procedure for efficient routine detection of pathogenic bacteria in plant material by polymerase chain reaction. <i>Journal of Microbiological Methods</i> , 37: 23–31. | 1) Paragraph [[137] should be after paragraph [138] (alphabetical order of the authors). 2) A full stop is missing after "1999". | EPPO, European Union, Georgia, Serbia |
| 59. | 142 | Editorial | Ordax, M., Biosca, E.G., Wimalajeewa, S.C., López, M.M. & Marco-Noales, E. 2009. Survival of <i>Erwinia amylovora</i> in mature apple fruit calyces through the viable but nonculturable (VBNC) state. <i>Journal of Applied Microbiology</i> , 107: 106–116. | Paragraph [142] should be after paragraph [143] (alphabetical order of the authors). | EPPO, European Union, Georgia, Serbia |
| 60. | 146 | Technical | Pirc, M., Ravnikar, M., & Tomlinson, J. & Dreo, T. 2009. Improved fireblight diagnostics using quantitative real-time PCR detection of <i>Erwinia amylovora</i> chromosomal DNA. <i>Plant Pathology</i> , 58: 872–881. | Correction of the list of authors. | EPPO, European Union, Georgia, Serbia |
| 61. | 156 | Editorial | Temple, T.N. & Johnson, K.B. 2011. Evaluation of loop-mediated isothermal amplification for rapid detection of <i>Erwinia amylovora</i> on pear and apple fruit flowers. <i>Plant Disease</i> , 95: 423– 430 43 . | The title of the reference shouldn't be underlined. Correction to page numbers. | EPPO, European Union, Georgia, Serbia |
| 62. | 158 | Editorial | Thomson, S.V. 2000. Epidemiology of fire blight. In J. Vanneste, ed. <i>Fire blight, the disease and its causative agent Erwinia amylovora</i> . Wallingford, UK, CAB International. Pgs pp. 9– 36 . | Consistency with the other references. | EPPO, European Union, Georgia, Serbia |
| 63. | 161 | Editorial | van der Zwet, T. & Keil, H.L. 1979. <i>Fire blight: A bacterial disease of rosaceous plants</i> . United States Department of Agriculture (USDA) Handbook 510. Washington DC, USDA. | New paragraph [161]bis: The reference "Weisberg et al., 1991" is missing (cf. paragraph [37], paragraph about "Internal control"). | EPPO, European Union, Georgia, Serbia |
| 64. | 165 | Editorial | Weisberg et al., 1991 9. Figures Figure 1. Flow chart for the diagnosis of fire blight in plants with symptoms. | 1. The number and name of the section is missing. 2. The legends of the figures should be put below the figures. | EPPO, European Union, Georgia, Serbia |

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| 65. | 166 | Editorial | <p>Figure 2. Flow chart for the identification of <i>Erwinia amylovora</i> in asymptomatic samples.</p> | An arrow seems to be missing in the 3rd last row between "Negative" and "E. amylovora not detected" | EPPO, European Union, Georgia, Serbia |

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| 66. | 166 | Substantive |  <p data-bbox="441 1050 1301 1204">Figure 2. Flow chart for the identification of <i>Erwinia amylovora</i> in asymptomatic samples.</p> | <p data-bbox="1370 330 1865 491">Insert "Other tests that may be recommended to be performed" following "IDENTIFICATION TESTS (section 4) is Negative" . In the case of a false negative, another test might be necessary for identification.</p> | Japan |

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| 67. | 167 | Editorial | <pre> graph TD A[Asymptomatic sample] --> B[Pathogen extraction and enrichment] B --> C[SCREENING TESTS (sections 3.1.4-3.1.5) (At least two tests based on different biological principles)] C --> D[All or some tests positive] C --> E[All tests negative] D --> F[ISOLATION AND ENRICHMENT-ISOLATION (section 3.1.3)] E --> G[E. amylovora not detected] F --> H[Colonies with typical morphology] H --> I[Negative] H --> J[Positive] I --> G J --> K[IDENTIFICATION TESTS (Section 4) (At least two tests based on different biological principles)] K --> L[Negative] K --> M[Positive] L --> G M --> N[E. amylovora identified] N --> O[Confirm pathogenicity by host test (section 4.4)] O --> P[E. amylovora confirmed] </pre> | <p>"Figure 2. Flow chart for the diagnosis of fire blight in asymptomatic plants." (cf. paragraph [56]): this missing legend should be put after figure 2.</p> | <p>EPPO, European Union, Georgia, Serbia</p> |

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| 68. | 167 | Technical | <pre> graph TD A[Asymptomatic sample] --> B[Pathogen extraction and enrichment] B --> C[SCREENING TESTS (sections 3.1.4–3.1.5) (At least two tests based on different biological principles)] C --> D[All or some tests positive] C --> E[All tests negative] D --> F[ISOLATION AND ENRICHMENT-ISOLATION (section 3.1.3)] F --> G[Colonies with typical morphology] G --> H[Negative] G --> I[Positive] H --> J[E. amylovora not detected] I --> K[IDENTIFICATION TESTS (Section 4) (At least two tests based on different biological principles)] K --> L[Negative] K --> M[Positive] L --> J M --> N[E. amylovora identified] N --> O[Confirm pathogenicity by host test (section 4.4)] O --> P[E. amylovora confirmed] </pre> | Isolation can also be performed as a screening test. Modify the flow diagram. Mention that if isolation is positive then there is no need to continue with another isolation. Consider whether there is a need in the flow diagram to refer to cases where there is a strong suspicion (3 positive assays, but no isolate) and therefore a need for isolation from new samples. This could be added as a footnote. | EPPO, European Union, Georgia, Serbia |
| 69. | 168 | Editorial | Footnote 1: See footnote 10. | This footnote is not understood and doesn't seem useful. | EPPO, European Union, Georgia, Serbia |
| 70. | 168 | Substantive | Footnote 1: See footnote 10. | It is lacking the other footnotes mentioned in the draft PD. | COSAVE, Uruguay, |

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| | | | | | Chile, Brazil, Peru, Argentina |
| 71. | 169 | Editorial | Footnote 21: The use of products of the brand bioMérieux in this diagnostic protocol implies no approval of them to the exclusion of others that may also be suitable. This information is given for the convenience of users of this protocol and does not constitute an endorsement by the CPM of the chemical, reagent and/or equipment named. Equivalent products may be used if they can be shown to lead to the same results. | Delete. Footnotes are not used for other brand names. Alternatively change the number of the footnote: cf. paragraph [80] and comment on paragraph [168]. | EPPO, European Union, Georgia, Serbia |
| 72. | 169 | Substantive | <p>Footnote 2: The use of products of the brand bioMérieux in this diagnostic protocol implies no approval of them to the exclusion of others that may also be suitable. This information is given for the convenience of users of this protocol and does not constitute an endorsement by the CPM of the chemical, reagent and/or equipment named. Equivalent products may be used if they can be shown to lead to the same results.</p> <p><u>Add the following explanation in the paragraph[170].</u></p> <p><u>Footnote3:When using LAMP on a regular basis in an area which has a patent system such as Japan (Patent Nos. 3,313,358, 3,974,441 and 4,139,424), the United States of America(US6,410,278, US6,974,670 and US7,494,790), the European Union (Nos. 1,020,534, 1,873,260, 2,045,337 and 2,287,338), China (ZL008818262), the Republic of Korea (Patent No. 10-0612551), Australia (No. 779160), and the Russian Federation (No. 2,252,964), it is necessary for users to receive a license from Eiken Chemical Co., Ltd. before use. To protect the intellectual property right.</u></p> | To protect the intellectual property right. | Japan |