

Emergence of Xylella fastidiosa in Spain: current situation

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Process for oficial diagnosis of Xylella fastidiosa in Spain



Official Plant Health Laboratory from Balearic islands









Autonomous community region level

> **Official Plant Health Laboratory from Comunitat Valenciana**









Laboratory of Biology & Ecology of Soil Microbiota IAS-CSIC

National level

- **Confirmation of Diagnosis**
- **Training and support**

National level

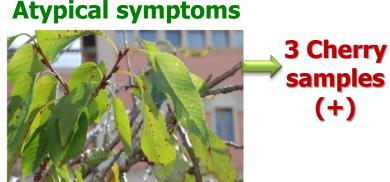
Subspecies characterization

- Official surveys and monitoring
- **Contingency Plan Implementation**

Bulletin OEPP/EPPO Bulletin (2016) 0 (0), 1-38 ISSN 0250-8052. DOI: 10.1111/epp.12327 European and Mediterranean Plant Protection Organization Organisation Européenne et Méditerranéenne pour la Protection des Plantes PM 7/24 (2) Diagnostics Diagnostic PM 7/24 (2) Xylella fastidiosa

First outbreak: October 2016 in a garden center near Manacor city

Official surveys Decision EU 2015/789



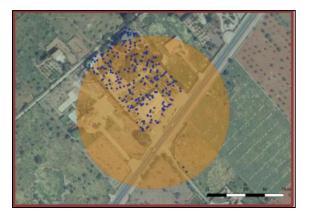




Typical symptoms



Infected zone



- > 258 samples analyzed
- > 1921 host plants eradicated

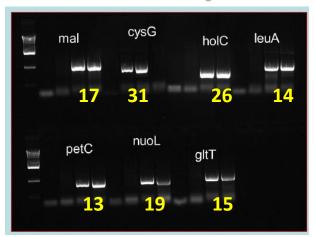




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> Identification of X. fastidiosa subspecies and STs in 1st outbreak

MLST analysis

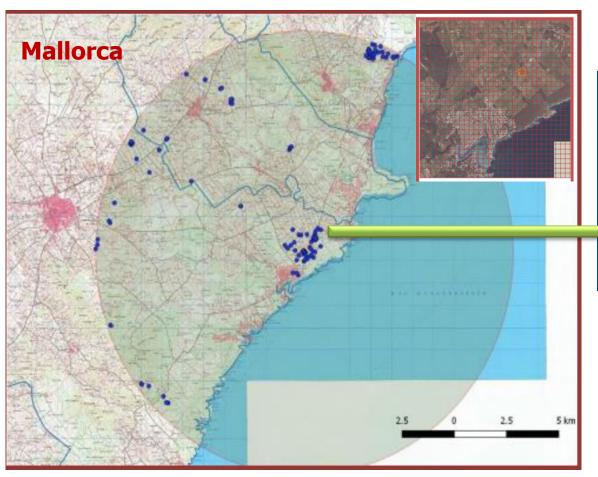


pilU and rpoD genes

ST	leuA	petC	malF	cysG	holC	nuoL	gitT
1	1	1	1	1	1	1	1

Host	#plants	Subspecies/ST
Cherry	3	Xf subsp. fastidiosa ST1
Polygala myrtifolia	3	Xf subsp. fastidiosa ST1

Contingency plan: Monitoring of the territory in the buffer zone



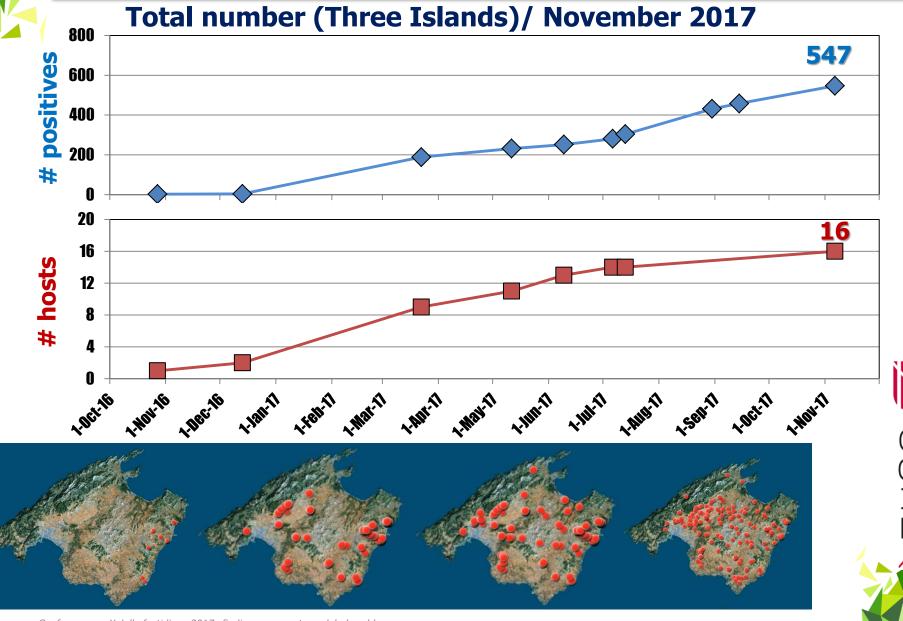
More Xf-positives appeared:

- In different host
- In different islands





Buffer zone samples taken in the demarcated area



Positives appeared spread throughout the three main islands

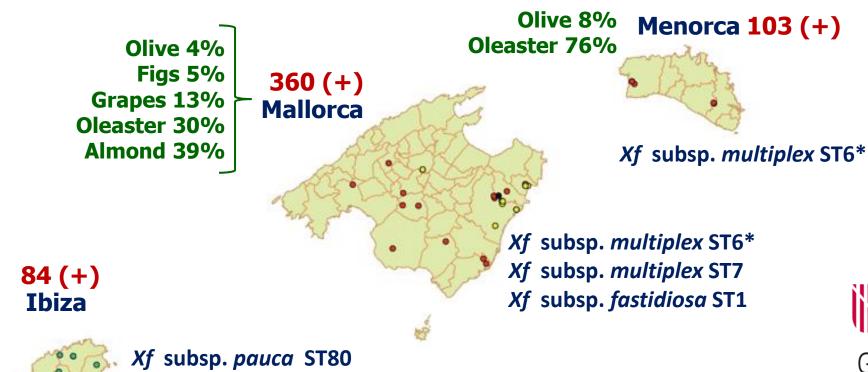


ST	leuA	petC	malF	cysG	holC	nuoL	gitT	Subsp
1	1	1	1	1	1	1	1	fastidiosa
6*	3	3	3	3*	3	3	3	multiplex 느
<u>7</u>	3	3	3	7	3	3	3	multiplex
73*	7	6	8*	27*	10	16	8*	pauca

Pending new
numbering for allele G
and ST assignation

New alleles and ST described

Three subspecies of Xylella fastidiosa and four Sequence types (ST)





Olive 33% Oleaster 50%

Subspecie

- Multiplex ST 7*
- Fastidiosa ST 1
- Multiplex ST 6*
- Pauca ST 80

Total positives = 547 (Nov 2017)

> Host plants: 16 species of crops, ornamentals and natural vegetation

Grapes *fastidiosa* ST1



Olive



Almond *fastidiosa* ST1



Oleaster *multiplex* ST6*/*pauca* ST80 *multiplex* ST6*/*pauca* ST80



Cherry *fastidiosa* ST1



Fig *multiplex* ST6*



Plum Subsp. ?



Walnut Subsp. P



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> Host plants: 16 species of crops, ornamentals and natural vegetation

Polygala myrtifolia **AII ST 1, ST6*, ST7, ST80**



Fraxinus angustifolia **Subsp.** *multiplex* **ST6***



Lavandula dentata *multiplex* ST6*/ *pauca* ST80



Acacia sp. *multiplex* ST6*/ *pauca* ST80



Rosmarius officinalis **Subsp.** *multiplex***ST6***



Nerium oleander Subsp. ?



Cistus monspeliensis **Subsp.** *fastidiosa* **ST1**



Rhamnus alaternus



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Olive (*Olea europea var. europaea*) *Xylella fastidiosa* subsp. *multiplex*

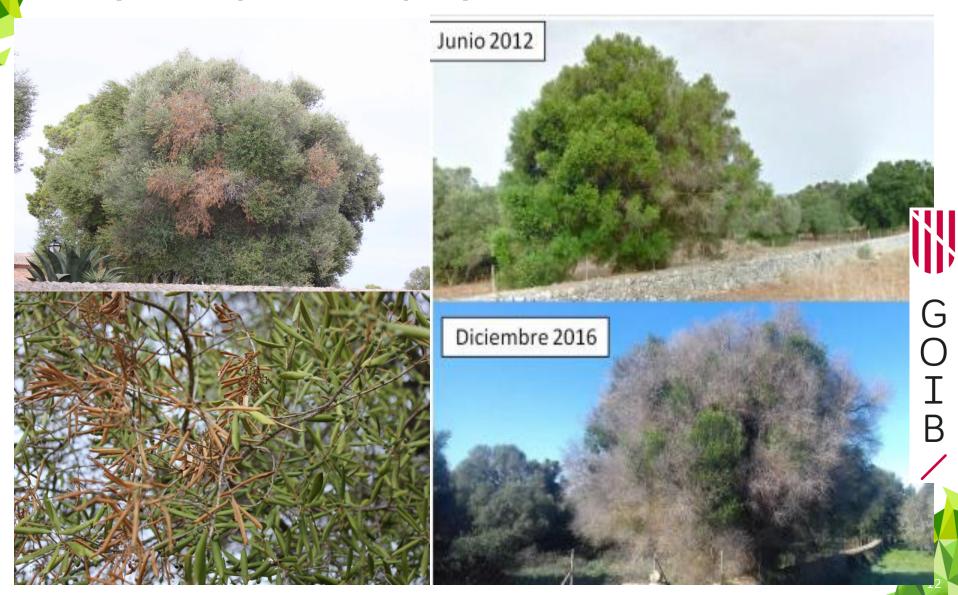








Olive (*Olea europea v*ar. *europaea*) *Xylella fastidiosa* subsp. *multiplex*



Olive (*Olea europea var. europaea*) *Xylella fastidiosa* subsp. *multiplex*



GOIB/

Almond (*Prunus dulcis*) Xylella fastidiosa subsp. fastidiosa









Almond (Prunus dulcis) Xylella fastidiosa subsp. fastidiosa





Grapes (Vitis vinifera)

Xylella fastidiosa subsp. fastidiosa









> From middle 2017 a Contention Plan proposed: Eradication of positives

About 60% of samples eliminated

Island	Number of Positives	Positive eliminated	TOTAL eliminated
Mallorca	360	171	2.263
Ibiza	103	88	318
Menorca	84	64	98
Formentera	0	0	1
TOTAL	547	323	2680



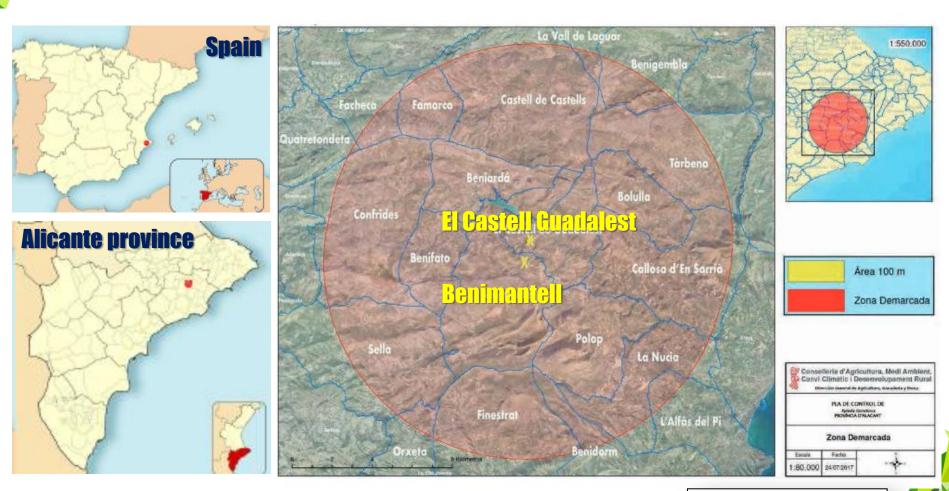




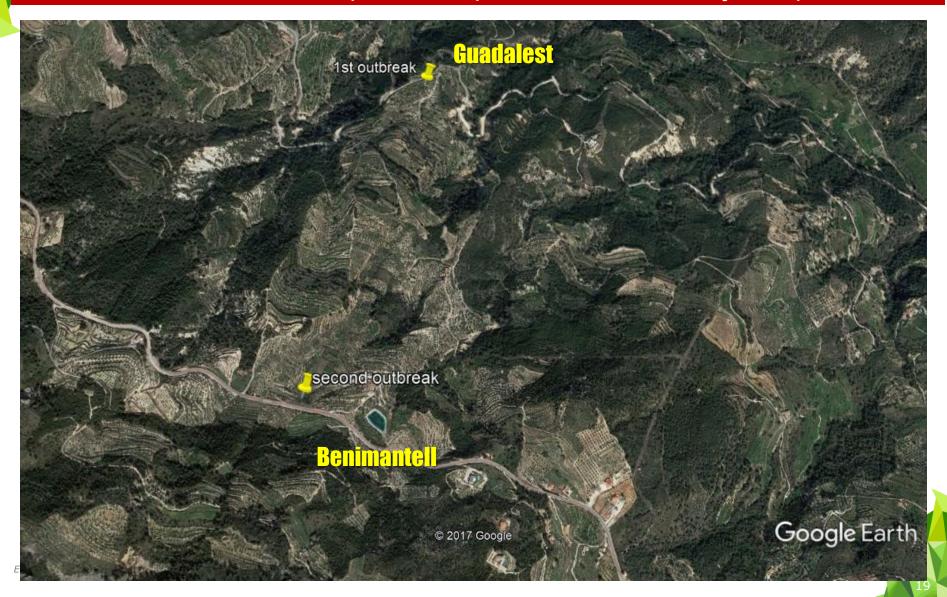


- **▶ 1**st outbreak: 29 June 2017, 1 orchard
- > 2nd Outbreak: 25 July 2017, 1 orchard

(34.383 has. demarcated)



> 1st outbreak: 29 June 2017, 1 orchard; 2nd Outbreak: 25 July 2017, 1 orchard



1st outbreak: 29 june 2017, 1 orchard; 2nd Outbreak: 25 July 2017, 1 orchard



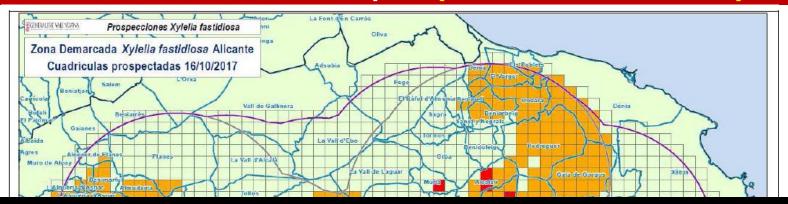
3rd outbreak: 31 August 2017, 26 almond plots (110.945 has demarcated)



Xylella fastidiosa subsp. multiplex ST pending



4th outbreak: 20 October 58 almond plots (139.789 has. demarcated)



106 positive samples (only almond)
86 plots

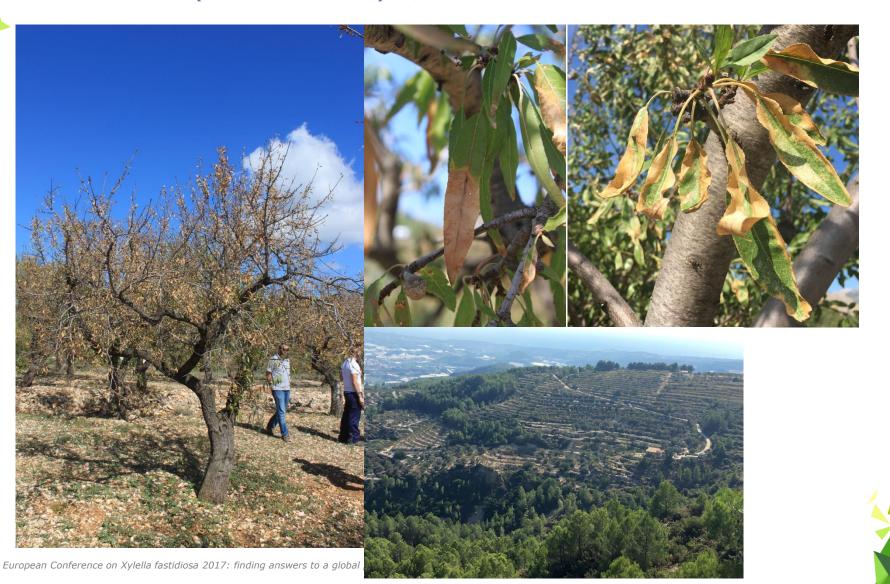
19 almond plots and 594 trees eradicated



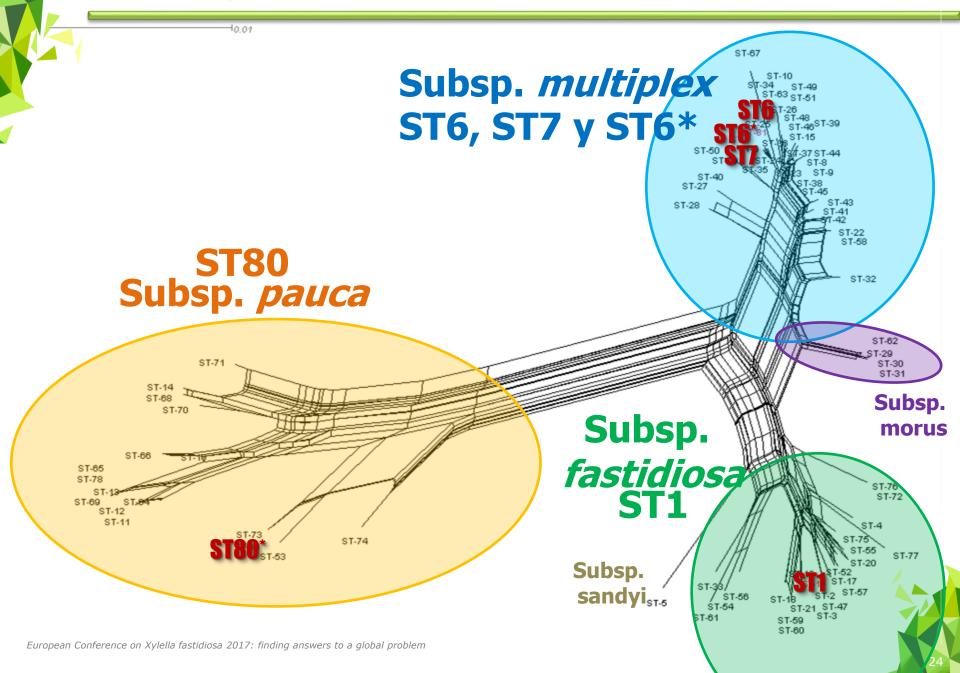
Xylella fastidiosa subsp. multiplex ST pending



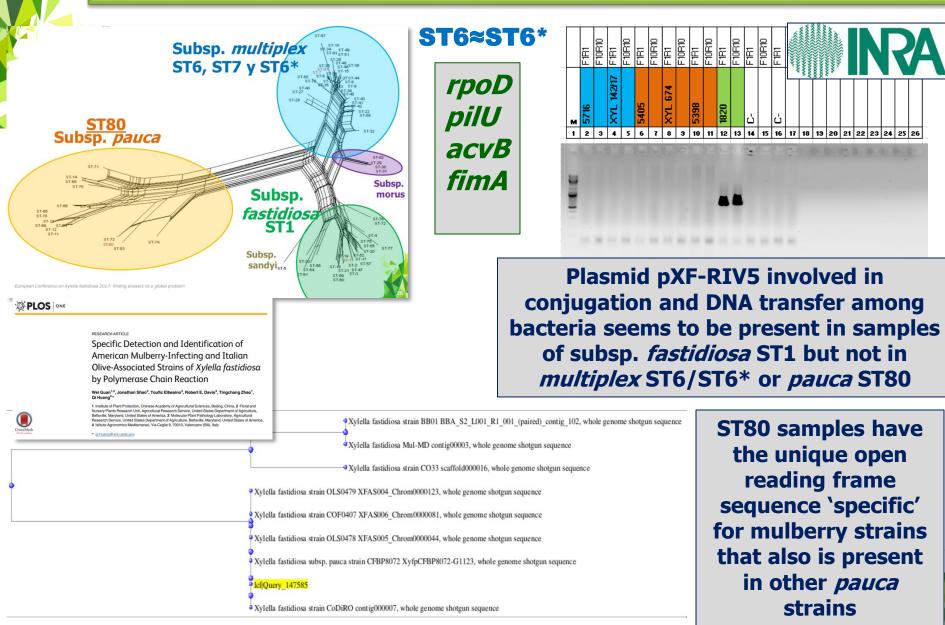
Almond (Prunus dulcis) Xylella fastidiosa subsp. multiplex ST 6



Current genetic diversity of Xylella fastidiosa in Spain



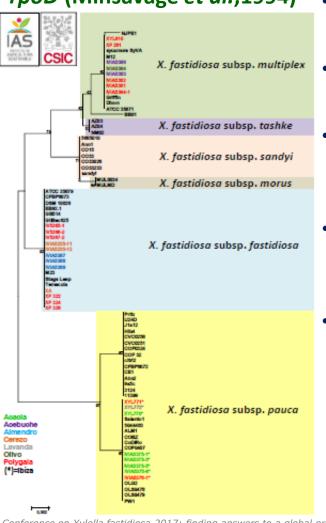
Current genetic diversity of Xylella fastidiosa in Spain



Current genetic diversity of Xylella fastidiosa in Spain

Difficult to identify the subspecies/ST for several positive samples

rpoD (Minsavage et al.,1994)



- **About 45% of samples identified to the** subspecies/ST level using the MLST approach
- DNA used from plant extracts (inhibition, low bacteria titer or Ct values high: >30)
- Most samples were initially processed in winter or early spring and many plants have been eradicated
- More than 70% of samples could be identified now at the subspecies level if following recommendations of revised EPPO protocol
- Strain isolation success was initially low, but now better -> sequencing of genomes

Revision of the EPPO protocol

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Situation of Xylella fastidiosa in Spain: Conclusions

- New host plants of *X. fastidiosa* have been described for Europe after outbreaks in Balearic islands and Alicante
 - Ficus carica, Fraxinus angustifolia, Juglans regia, Prunus domestica,
 Vitis vinifera
- ST6 widespread in France (Corsica) but not detected in almond (1 case only) whereas is the only host infected in Alicante so far
- > New subspecies and sequence types for Europe has been described
 - Subsp. fastidiosa ST1
 - Subsp. multiplex ST6* (A case in olive in California?)
 - Subsp. pauca ST80 (New for science)
- Several and independent introductions have occurred in Europe and Spain in the past and from different origins (probably EEUU and Central America)
- Sequencing of the genome of representative isolates can provide insights into their origin and biology



Acknowledgements















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