



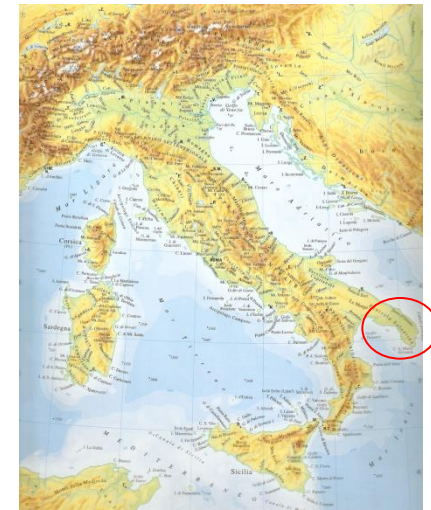
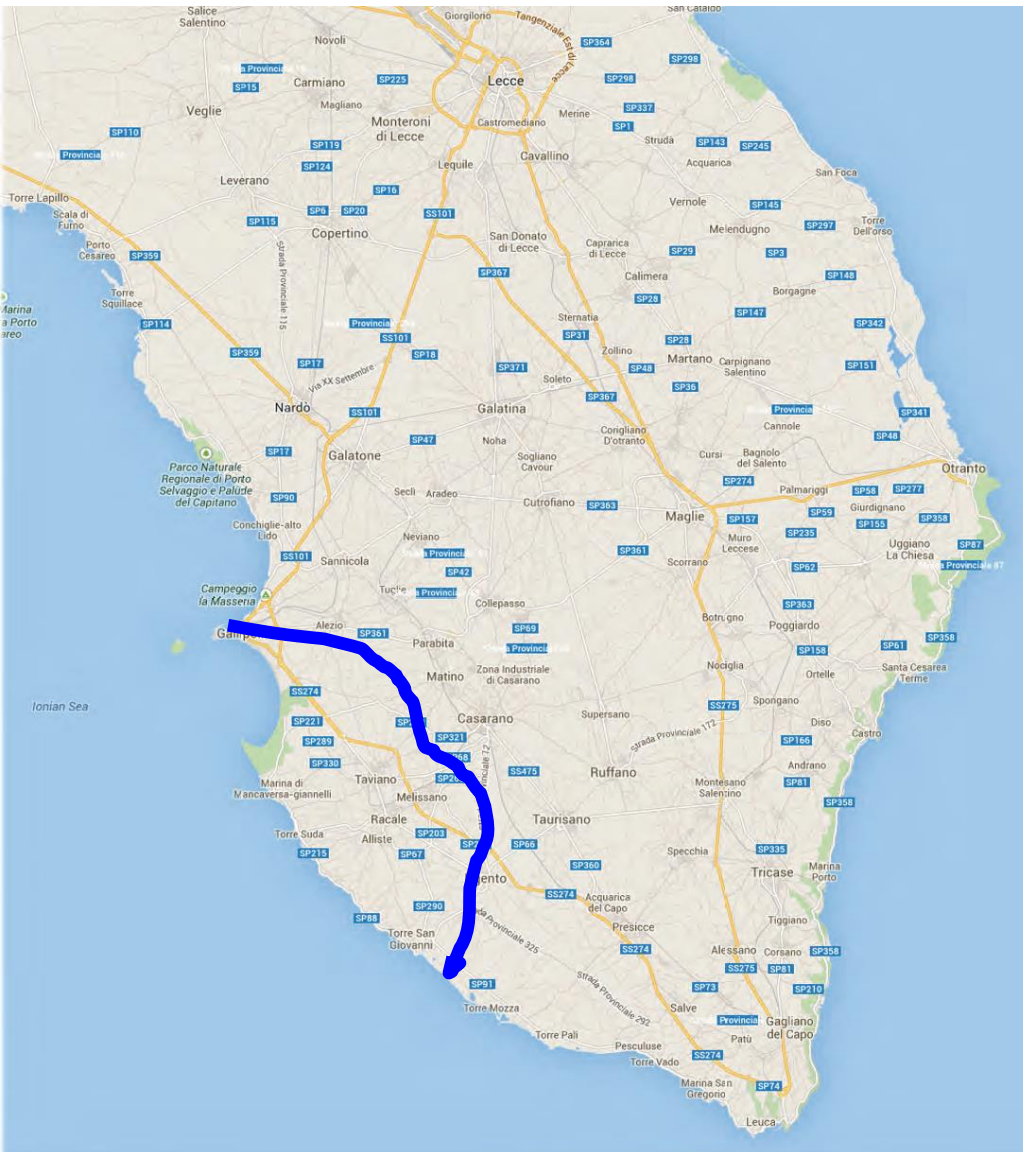
UNIVERSITÀ
DEGLI STUDI DI BARI
ALDO MORO



Consiglio Nazionale delle Ricerche
Istituto per la Protezione Sostenibile
delle Piante
Bari

Current situation on the outbreak of *Xylella fastidiosa* in Southern Italy

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Institute for Sustainable Plant Protection
Former Institute of Plant Virology,
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Bari - Italy



The "Quick decline syndrome" is a severe disease of olive that appeared a few years ago in a restricted area near Gallipoli (province of Lecce, Salento peninsula, southern Italy). It soon expanded in epidemic form in the area delimited by a **blue line** which, in 2013, amounted to *ca.* 8,000 ha

Olives trees in 2010



The same trees in 2013



Symptoms of the disease consist in the appearance of leaf scorching and dessication of small peripheral branches distributed at random on the canopy which, with time, extend to the rest of it



In the groves heavily affected by "quick decline" all plants are symptomatic



Final stage of a "quick decline" attack. Olives have been heavily pruned in the hope to stimulate new sprouting



The canopy was totally dessicated but the plants are not dead. They are pushing suckers from the crown







THE PUTATIVE ACTORS

Quick decline that affects aged olive trees is a complex disease in whose aetiology the following agents seem to be implicated:

- (i) The leopard moth (*Zeuzera pyrina*)
- (ii) A set of xylem-inhabiting fungi, especially of the genus *Phaeoacremonium*, among which *P. parasiticum* prevails. These fungi are frequent inhabitants of olive wood and the precursors of a common wood condition of old trees known as wood decay
- (iii) The xylem-limited bacterium *Xylella fastidiosa*

The Leopard moth (*Zeuzera pyrina*)



Adult



Larva



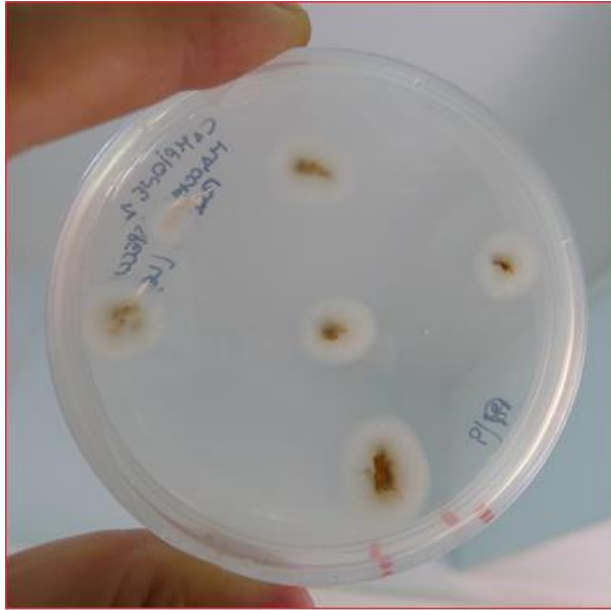
Entrance of the galleries drilled in the branches



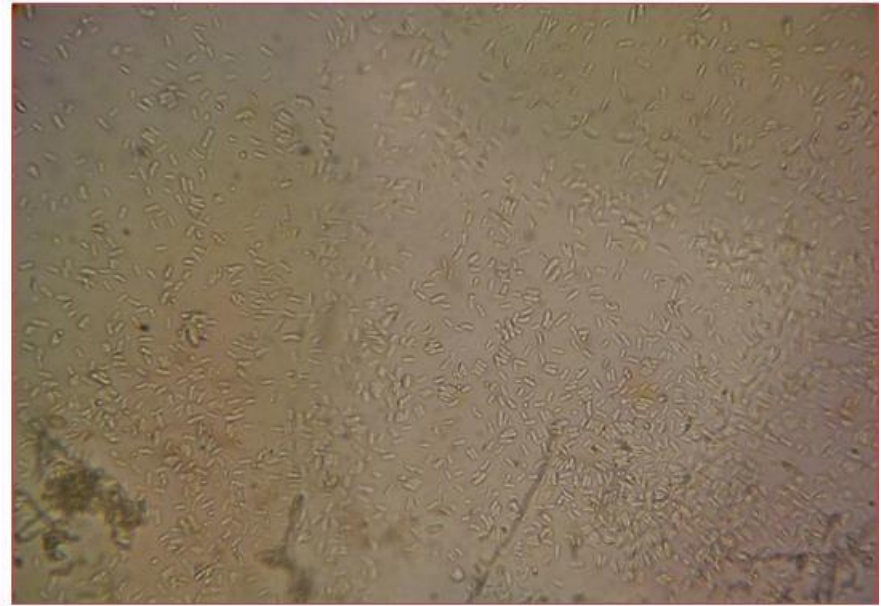
Galleries in a trunk



The fungus (*Phaeoacremonium parasiticum*)



Colonies



Conidia

Journal of Plant Pathology (2013), 95 (3), 659-668

DISEASE NOTE

FUNGAL SPECIES ASSOCIATED WITH A SEVERE DECLINE OF OLIVE IN SOUTHERN ITALY

F. Nigro¹, D. Boscia², I. Antelmi¹ and A. Ippolito¹

Phaeoacremonium parasiticum and the olive



Sectioned olive branch showing extensive necrosis of the sapwood colonized by *P. parasiticum*



Necroses start from *Zeuzera* galleries



Xylella fastidiosa

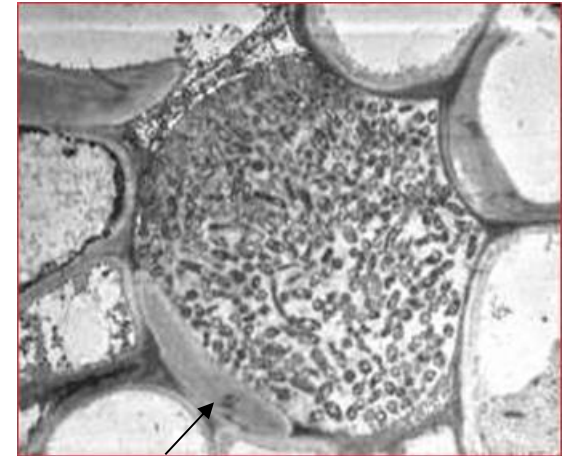
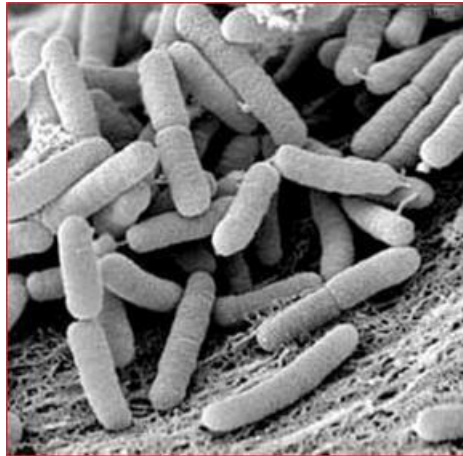
Gram-negative rod-shaped bacterium with a thick rippled cell wall that localizes in the xylem vessels and can be grown in axenic culture with difficulty. It is transmitted by xylem-feeding hemiptera (cicadellidae and cercopidae) and has a wide natural host range, i.e. more than 150 herbaceous and woody species, some of which are of great economic importance (grapevine, coffee, citrus, stone fruits). Infection to olive is a rare event. Up to 2013 reported only from California

Journal of Plant Pathology (2013), 95 (3), 659-668

DISEASE NOTE

**IDENTIFICATION OF DNA SEQUENCES
RELATED TO *XYLELLA FASTIDIOSA* IN
OLEANDER, ALMOND AND OLIVE TREES
EXHIBITING LEAF SCORCH SYMPTOMS
IN APULIA (SOUTHERN ITALY)**

M. Saponari¹, D. Boscia¹, F. Nigro² and G.P. Martelli^{1,2}



Role of *Xylella* in the genesis of quick decline



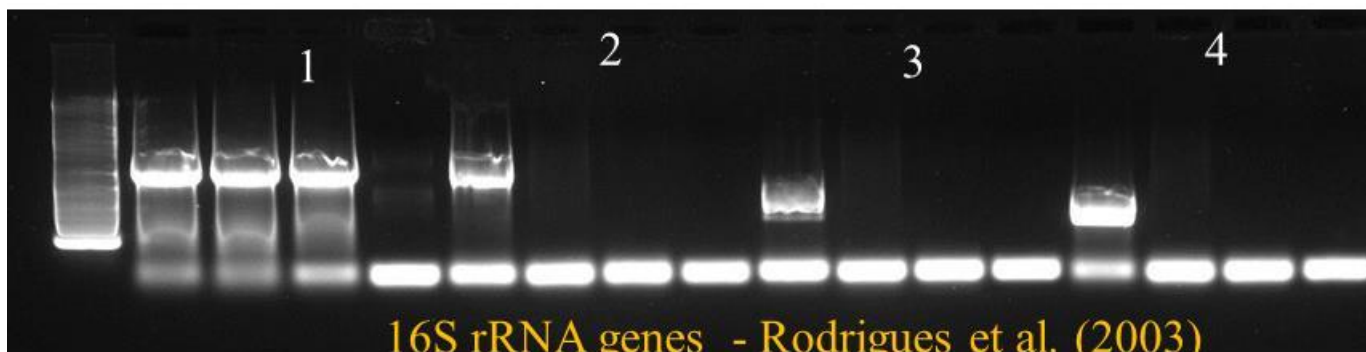
To know it, **one must await the results of pathogenicity tests** (graft-transmission and infection with pure inocula from bacterial cultures).

It is possible, however, to hypothesize for *Xylella* the role of aggravator.

In fact, it is conceivable that the invasion and clogging of xylem vessels that have escaped fungal colonization would impair water uptake with consequent collapse of the tree

Identification of *Xylella* in Salentinian olives: PCR

The type of symptoms and the velocity with which quick decline had spread from the first infection focus had suggested, at the end of September 2013, the analysis of symptomatic olives for the presence of *Xylella fastidiosa*. PCR tests using primers for the amplification of the 16S rRNA gene gave positive results that were later confirmed by thousands of additional tests.

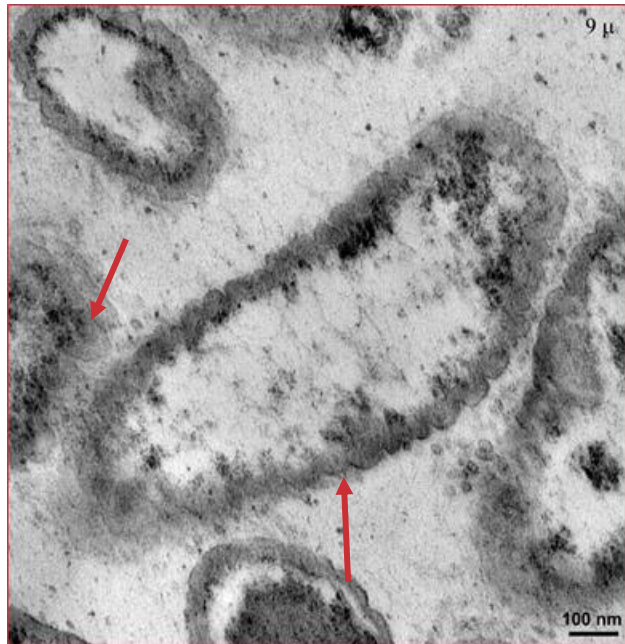


1. 16S Universal for bacteria
2. 16S *Xylella*-specific (0067-1439)
3. 16S *Xylella*-specific (0067-0838A)
4. 16S *Xylella*-specific (08385-1439)

Identification of *Xylella* in Salentinian olives: electron microscopy

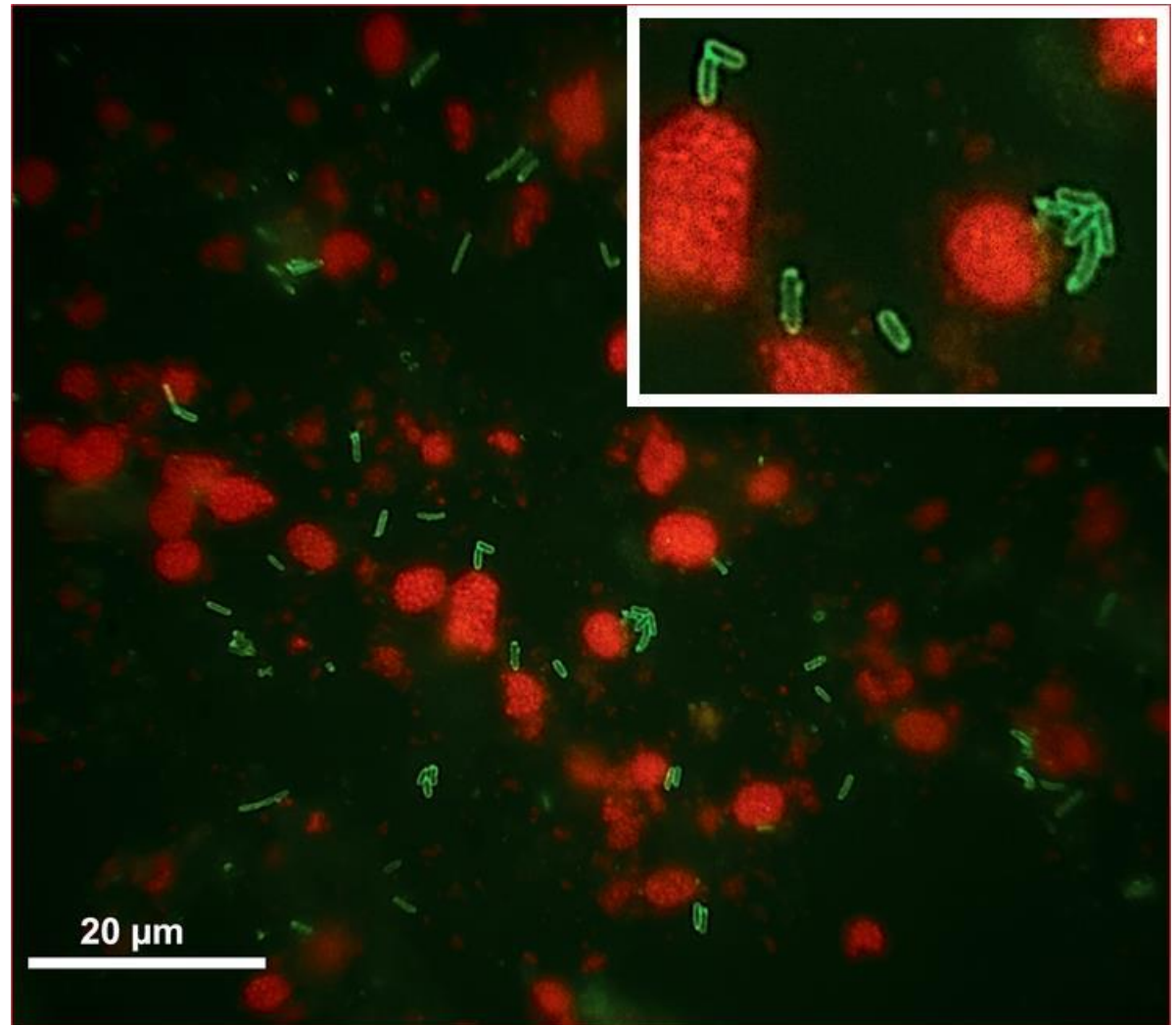
Bacterial cells within a xylem vessel

Bacterial cell showing
the rippled cell wall
typical of *X. fastidiosa*



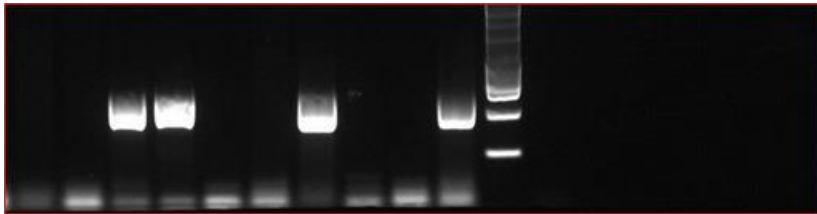
Identification of *Xylella* in Salentinian olives: immunofluorescence

Extract from the
petiole of an olive
leaf exposed to an
immunofluorescent
kit (Loewe)

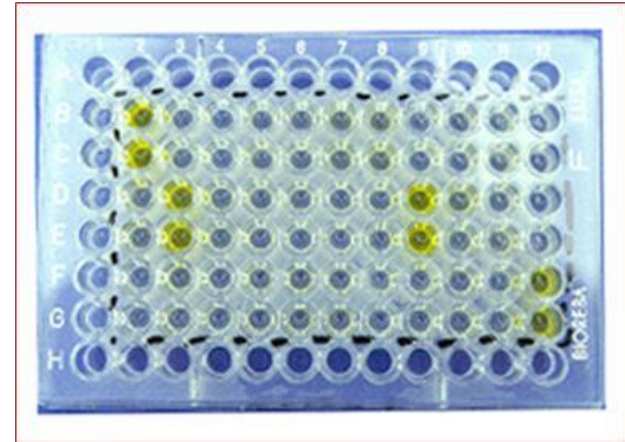


Diagnosis

The first tests were carried out with PCR using four different *Xylella*-specific primers



ELISA was adopted following the results of a ring-test in which three accredited laboratories participated. Doubtful results have always been checked by PCR

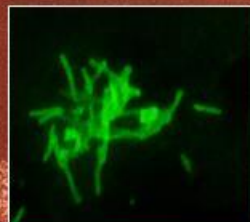


LETTER TO THE EDITOR

DETECTION OF *XYLELLA FASTIDIOSA* IN OLIVE TREES BY MOLECULAR AND SEROLOGICAL METHODS

G. Loconsole¹, O. Potere², D. Boscia¹, G. Altamura³, K. Djelouah⁴, T. Elbeaino⁴, D. Frasheri⁴, D. Lorusso⁴, F. Palmisano³, P. Pollastro³, M.R. Silletti³, N. Trisciuzzi³, F. Valentini⁴, V. Savino² and M. Saponari¹

Xylella was successfully isolated in
culture
from several infected species



Journal of Plant Pathology (2014), 96 (3), 1-5



Edizioni ETS Pisa, 2014

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doi: 10.4454/JPP.V96I2.024

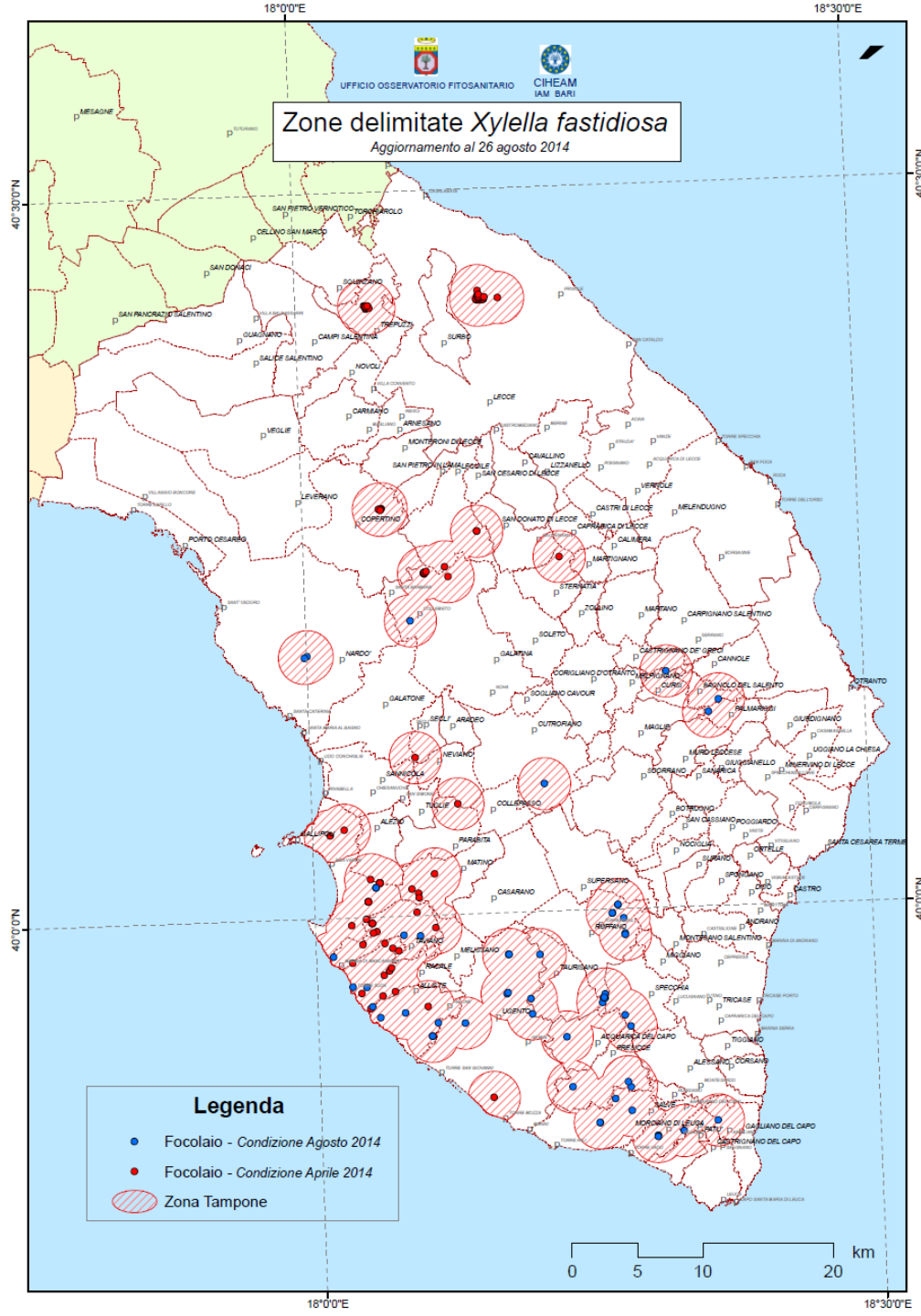
SHORT COMMUNICATION

ISOLATION OF A *XYLELLA FASTIDIOSA* STRAIN INFECTING OLIVE AND OLEANDER IN APULIA, ITALY

C. Cariddi¹, M. Saponari², D. Boscia², A. De Stradis², G. Loconsole², F. Nigro¹, F. Porcelli¹, O. Potere¹
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Host range

Oleander





Almond





Cherry







Polygala myrtifolia



*Polygala
mirtifolia*



Westringia fruticosa



*Westringia
fruticosa*



Acacia saligna



Spartium junceum



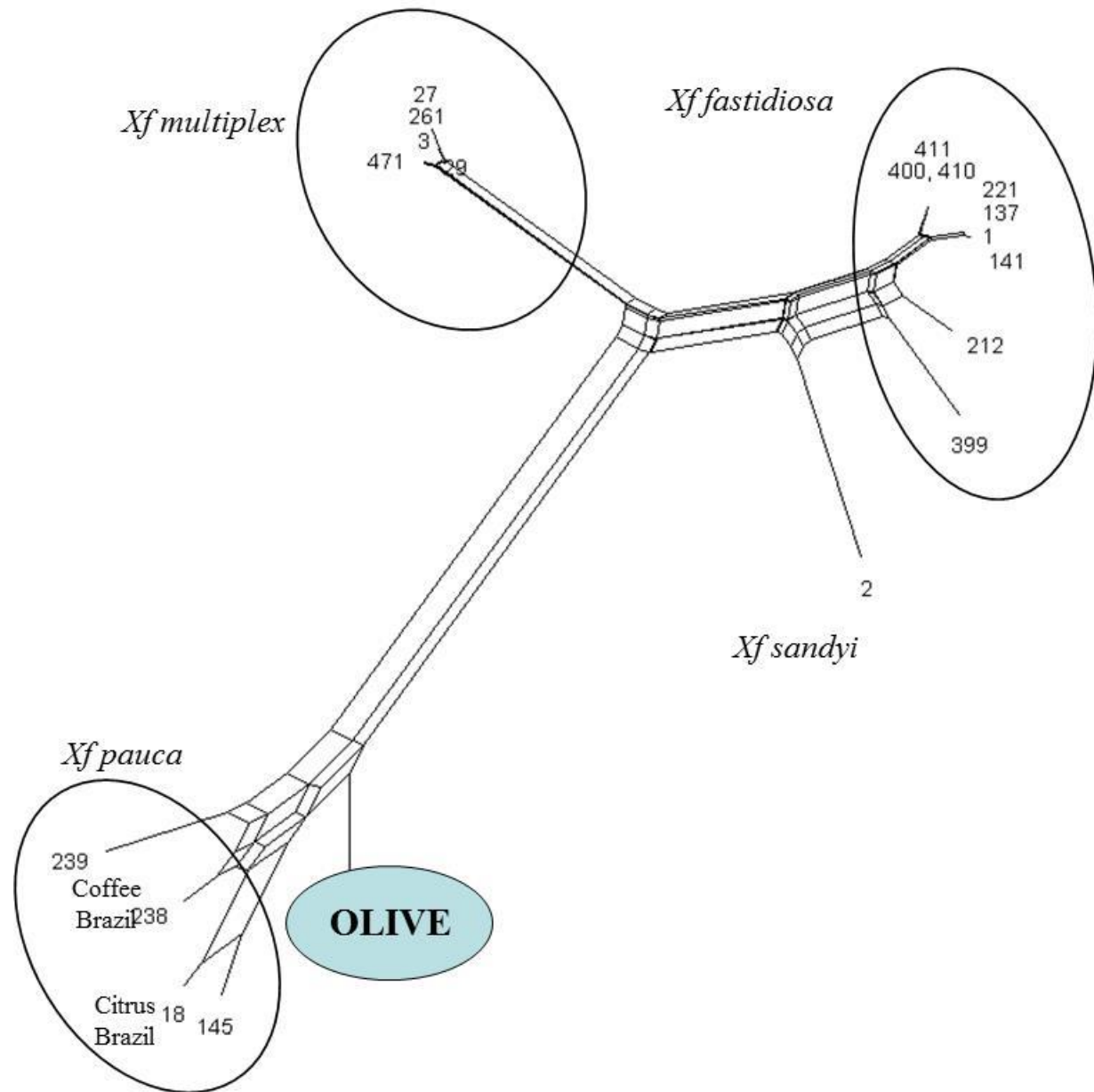
The monthly monitoring (since december 2013) of weeds (over 100 species) did not identify any further host of *Xylella fastidiosa*

Grapevine and Citrus were
never found infected

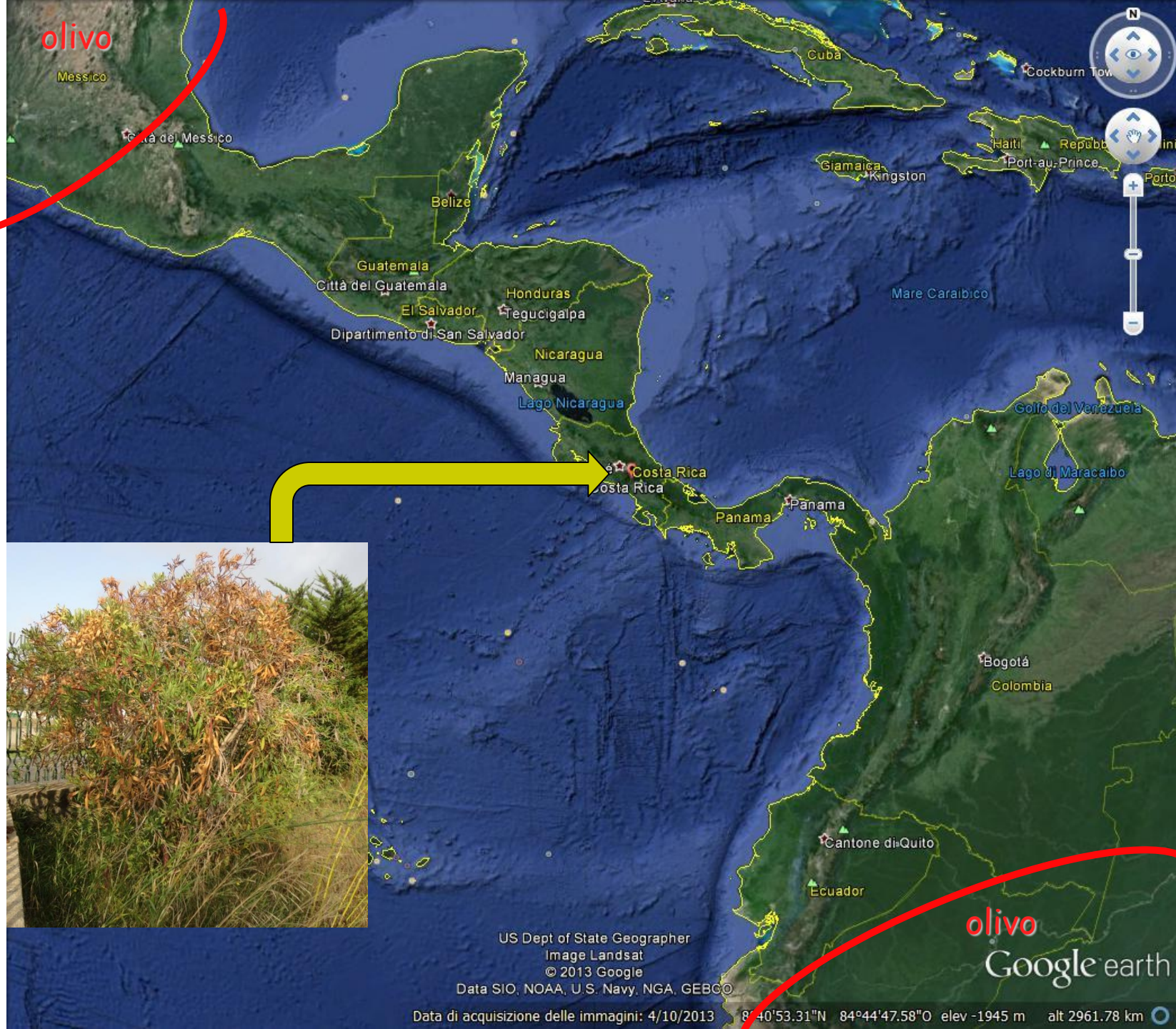


Concatenated
sequences of
the seven genes
showed that the
Salentinian strain
is a divergent
variant of
Xf pauca.

This variant
proved to be
**identical to a
variant infecting
oleander in
Costa Rica**



olivo



US Dept of State Geographer
Image Landsat
© 2013 Google

Data SIO, NOAA, U.S. Navy, NGA, GEBCO

Data di acquisizione delle immagini: 4/10/2013 8°40'53.31"N 84°44'47.58"O elev -1945 m alt 2961.78 km

olivo
Google earth

Epidemiology

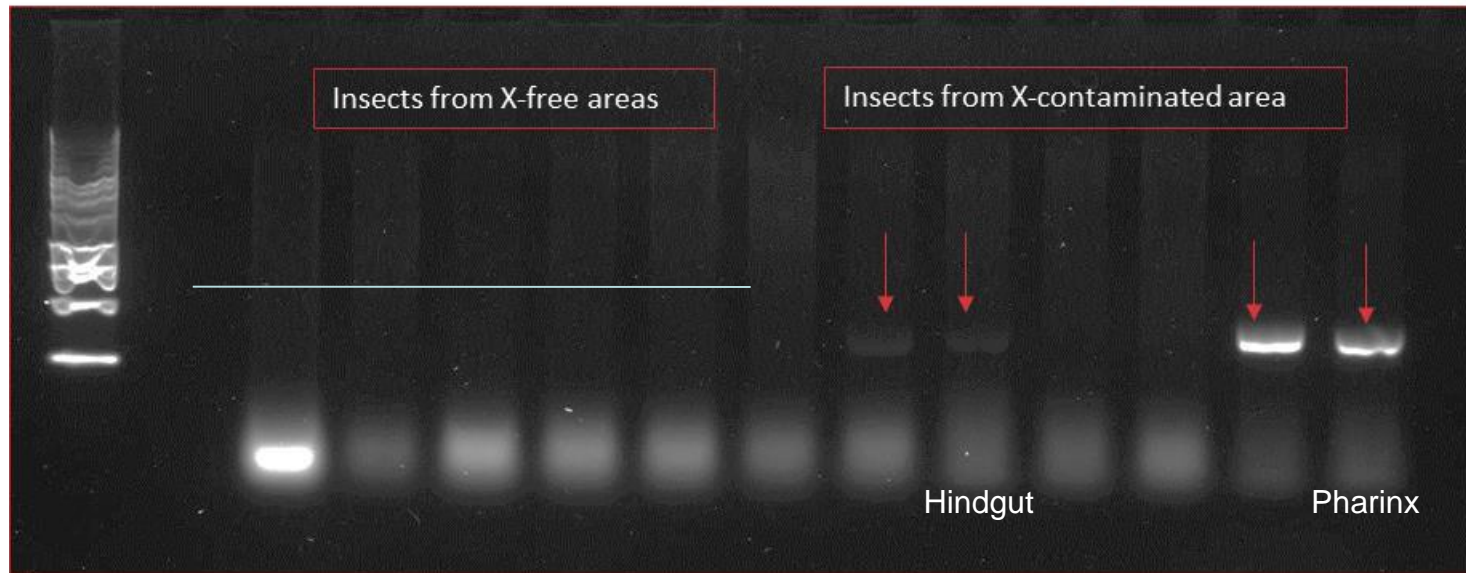
Epidemiological studies underway follow four different routes:

- (i) Identification of natural sources of inoculum (natural flora)
- (i) Capture of leafhoppers thriving on the natural flora, their identification and analysis for the presence of *Xylella*
- (iii) Transmission trials using *Xylella*-positive insects
- (iv) Placement of bait plants in infected olive groves



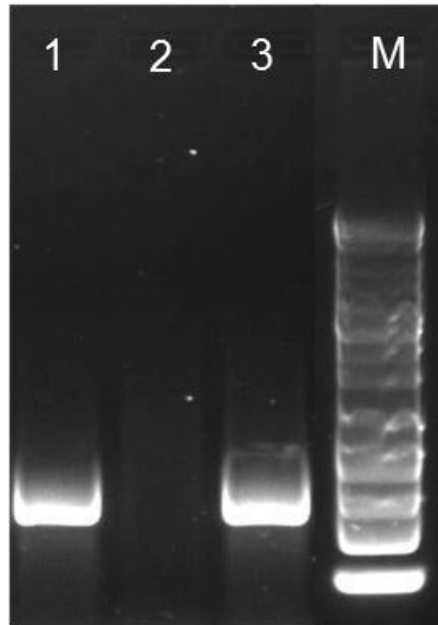
Vectors

Xylella was identified in several individuals of *Philaenus spumarius* captured in diseased olive groves but in none of those found in *Xylella*-free areas

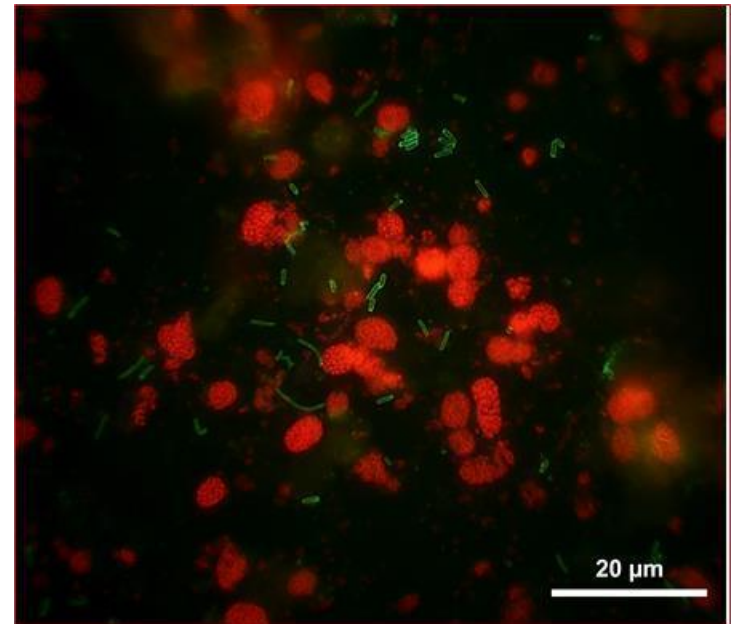


Transmission trials

Xylella-positive adults of *P. spumarius* that were allowed to feed on *Vinca rosea* seedlings were able to transmit the bacterium to two of five seedlings. Ascertained by PCR and immunofluorescence



1, Control(+); 2, Control(-);
3, *Vinca* exposed to *P. spumarius*



ARTHROPODS IN RELATION TO PLANT DISEASE

Infectivity and Transmission of *Xylella fastidiosa* by *Philaenus spumarius* (Hemiptera: Aphrophoridae) in Apulia, Italy

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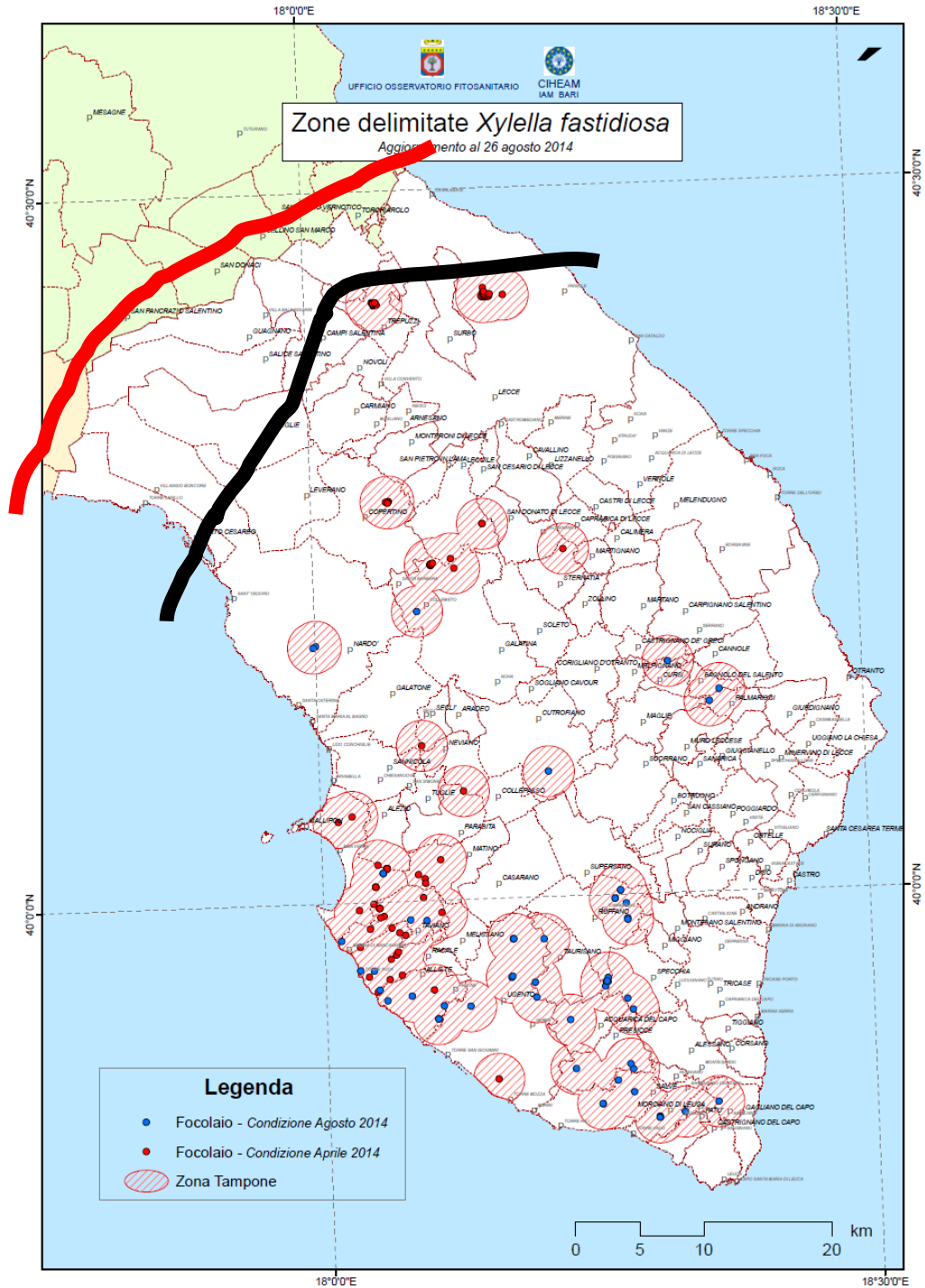
J. Econ. Entomol. 107(4): 1316–1319 (2014); DOI: <http://dx.doi.org/10.1603/EC14142>



The **control of vectors** is actually considered the most effective action to contain the spread of the epidemic

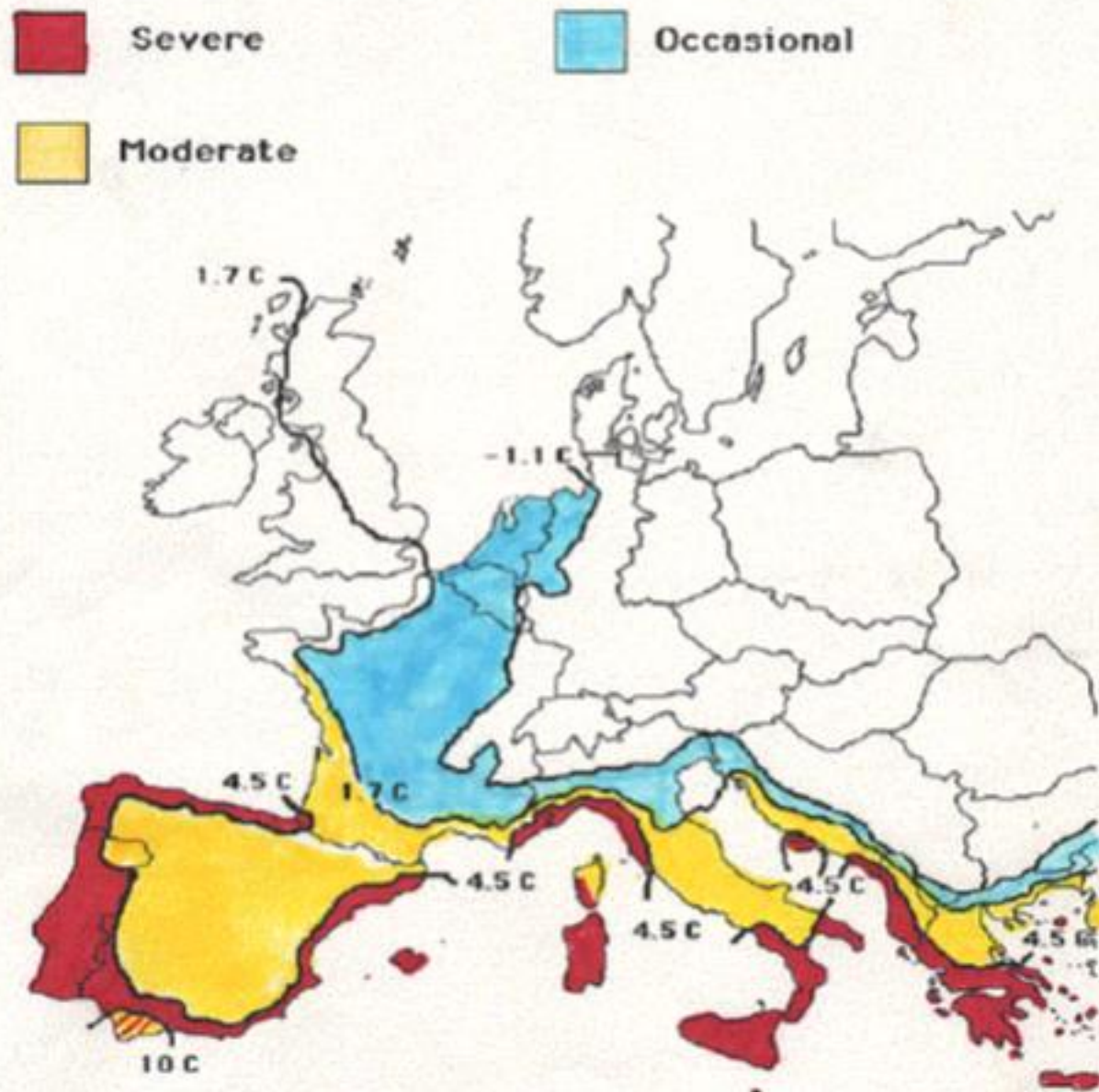
Buffer zone

Security line



The influence of the climate

Potential for Pierce's Disease: Estimates from Minimum January Temperatures



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Plant Protection Service of Apulia Region