



Xylella fastidiosa : **state of the art**

genetic variability,

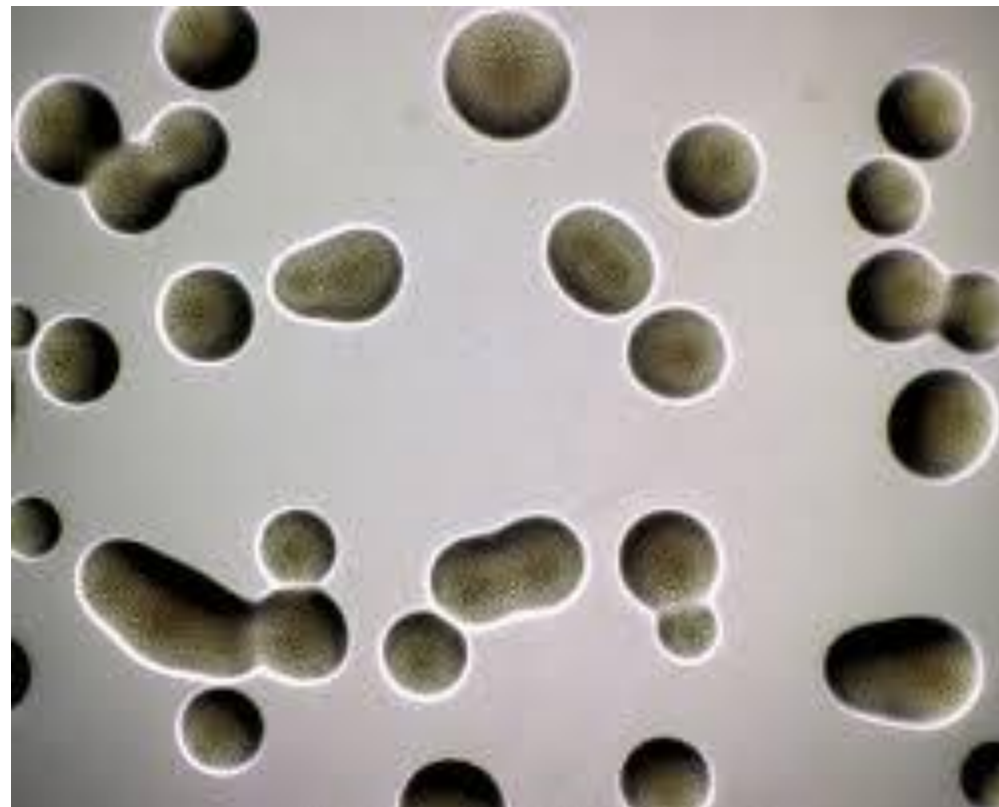
hosts, symptomatology, distribution

Donato Boscia

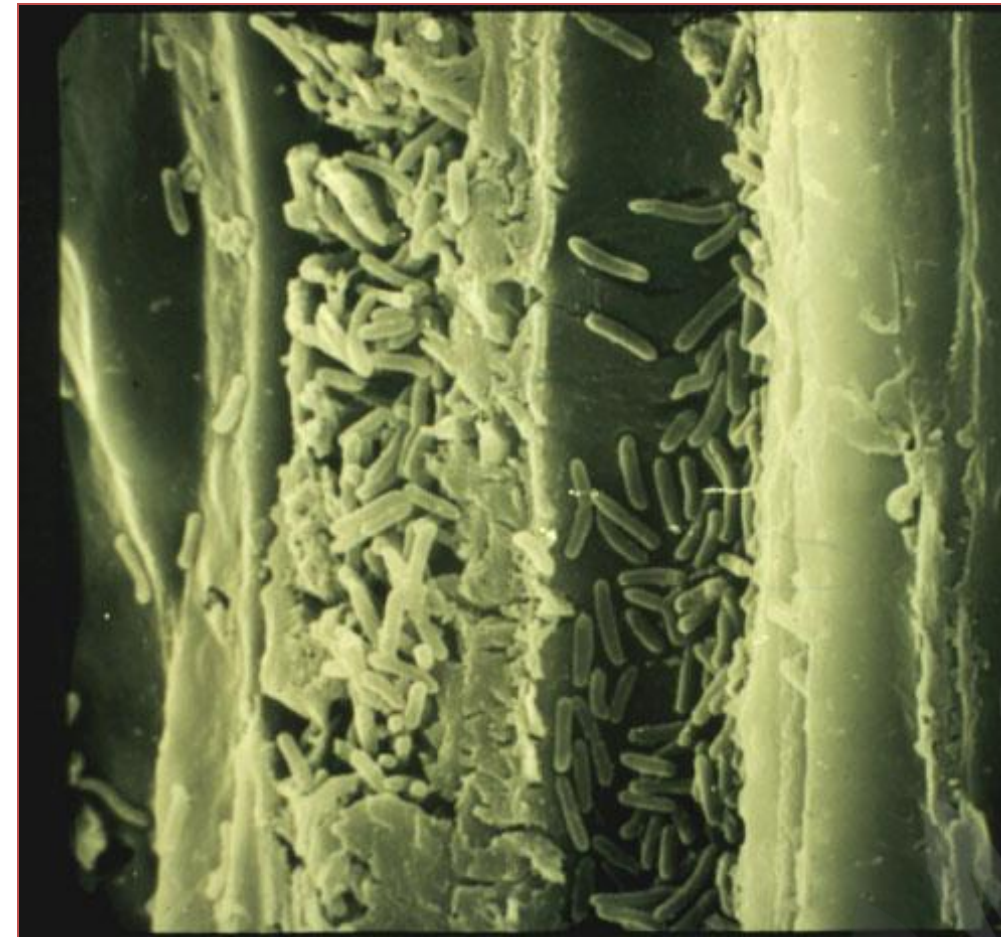
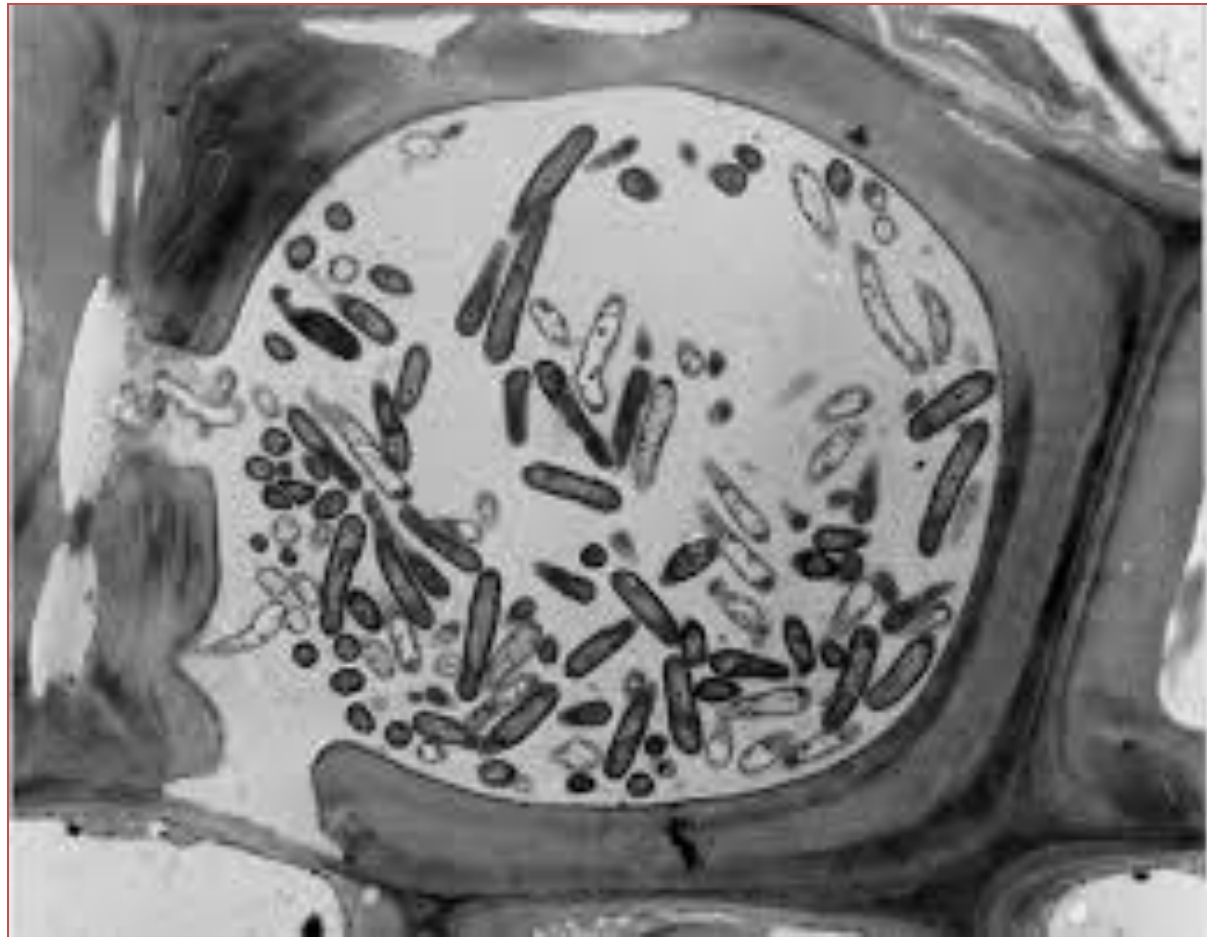
E-mail: donato.boscia@cnr.it



Xylella fastidiosa is a Gram-negative bacterium, non sporing,
slow-growing in axenic culture



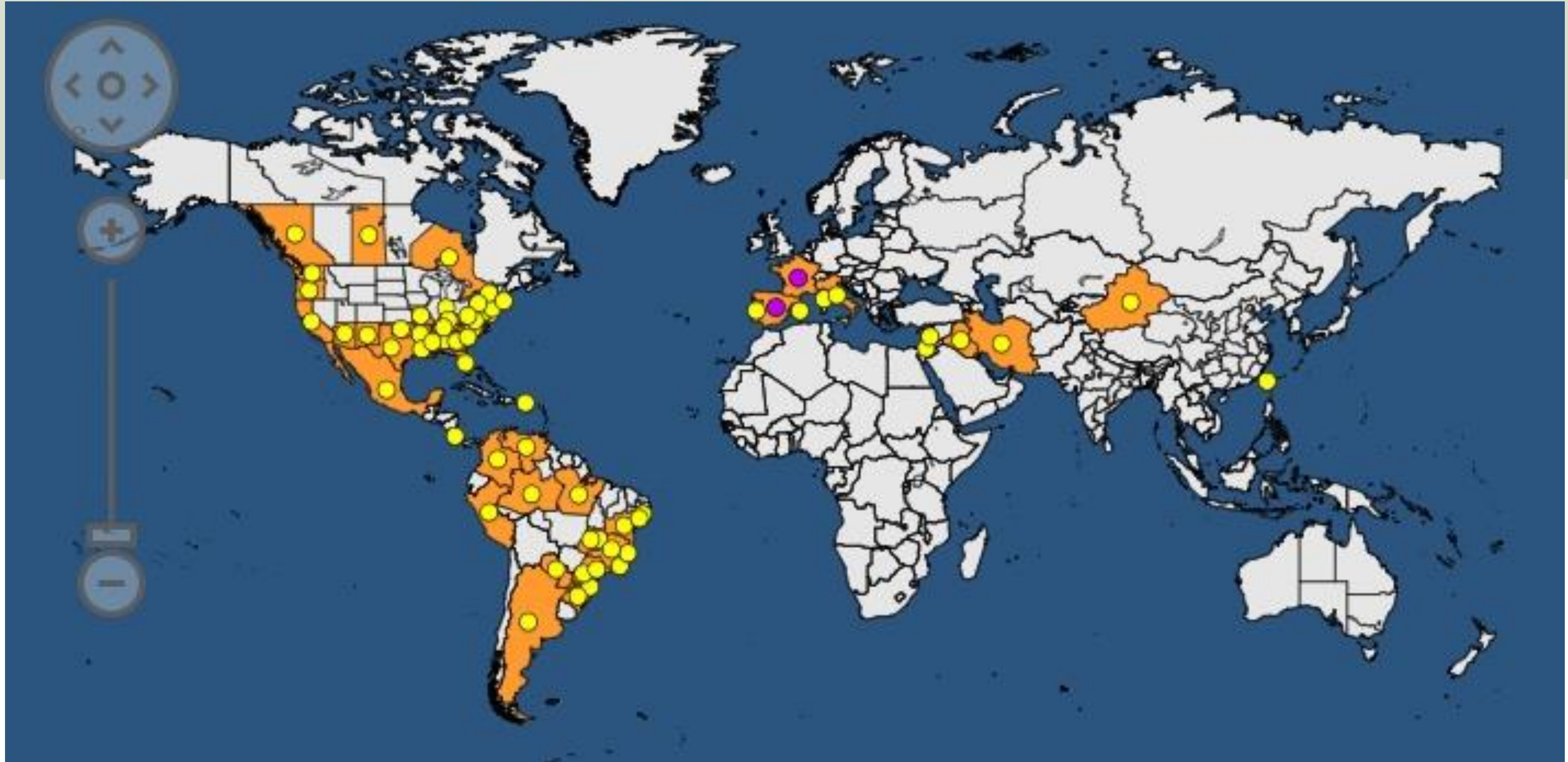
It localizes in the xylem vessels



In addition to the movement of infected plants, responsible for the spread over long distances, the natural spread of the bacteria from plant to plant occurs via xylem feeding insects belonging to the Order *Hemiptera*.



Wide geographical distribution



Its natural host range includes
71 families and 210 genera, for a total of 463 plant species.
It is the recognized agent of infectious diseases
affecting many economically important crops.



Pierce's disease



Kills grapevines, which cannot be grown in large areas
of the USA and the Gulf of Mexico



Citrus variegated chlorosis

(Central America, Argentina, Brazil)



Leaf scorch of stone fruits

Almond



Cherry



Plum

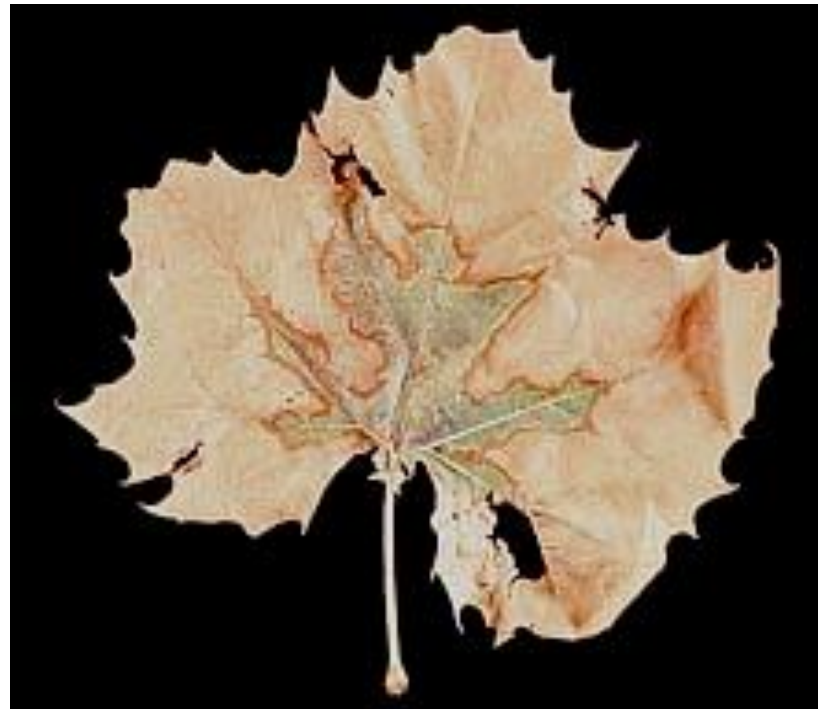


Olive Quick Decline Syndrome

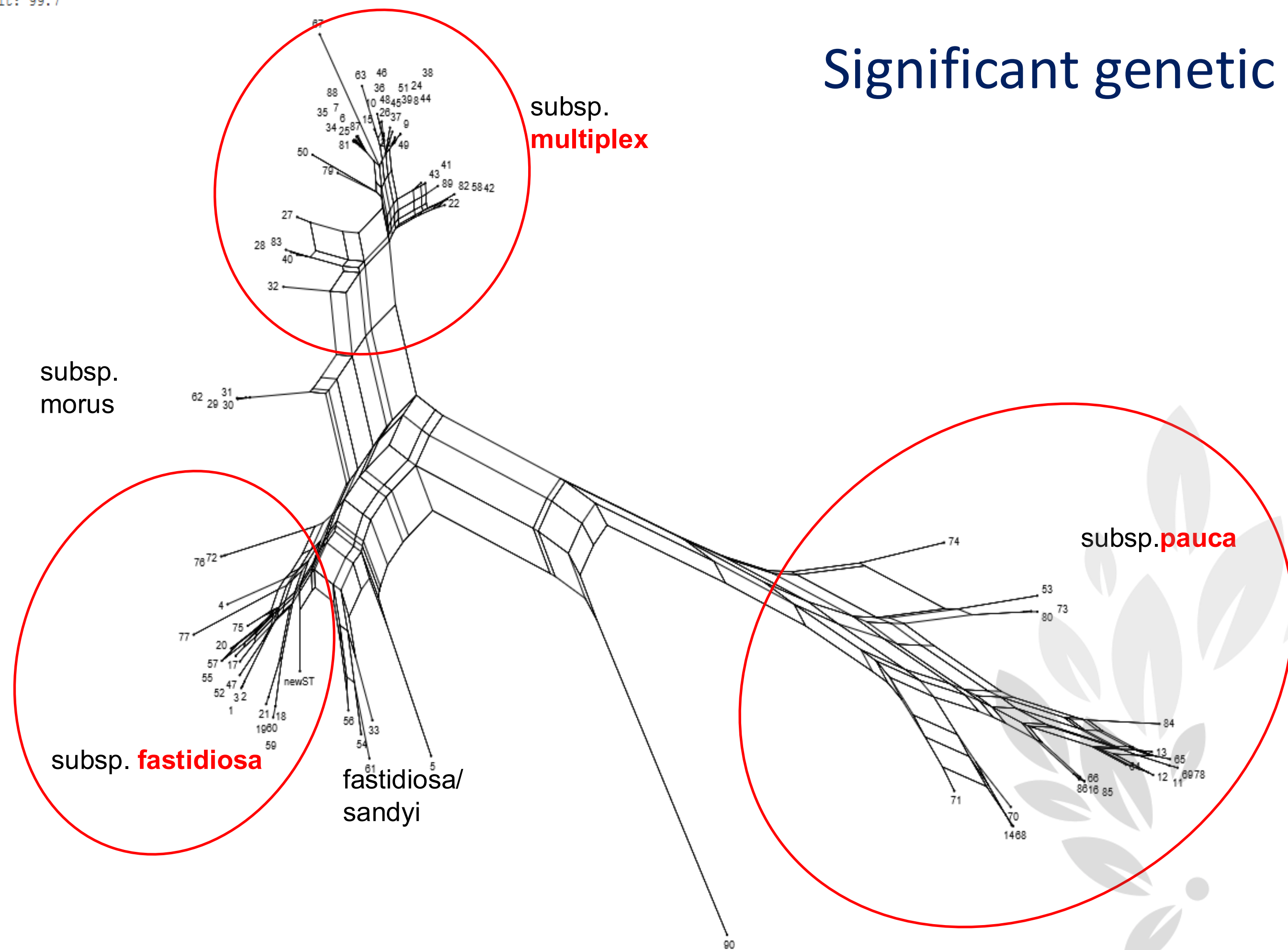
(Apulia, Argentina, Brasil, Balearic islands)



Leaf scorch of forest and shade trees



Significant genetic variability



It is a quarantine organism included in the EPPO A2 list



EPPO A2 List of pests recommended for regulation as quarantine pests

- version 2025-09 -

EPPO recommends its member countries to regulate the pests listed below as quarantine pests (A2 pests are locally present in the EPPO region). The EPPO A2 List is reviewed every year by the Working Party on Phytosanitary Regulations and approved by Council.



In EU is regulated by

L 269/2

EN

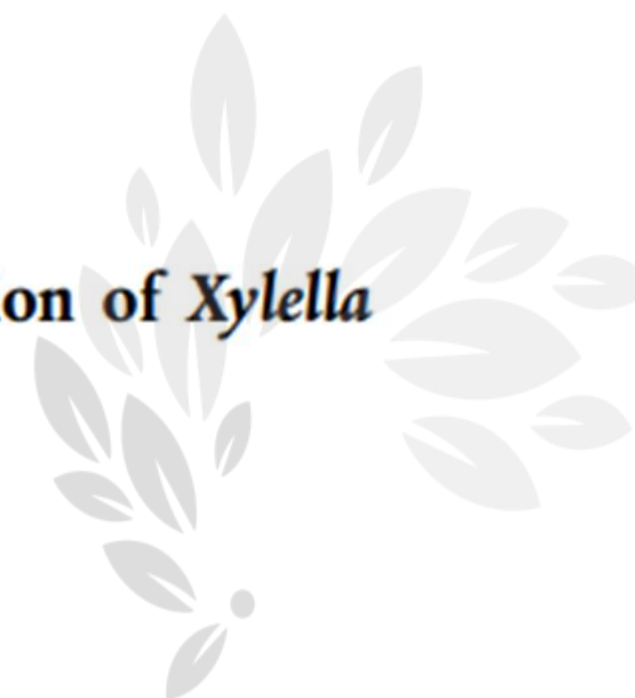
Official Journal of the European Union

17.8.2020

COMMISSION IMPLEMENTING REGULATION (EU) 2020/1201

of 14 August 2020

as regards measures to prevent the introduction into and the spread within the Union of *Xylella fastidiosa* (Wells *et al.*)



October 2013:
In Apulia first discovery of established infections of
X. fastidiosa in EU

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}



2015: annual monitoring obligation for all member states

L 125/36

EN

Official Journal of the European Union

21.5.2015

DECISIONS

COMMISSION IMPLEMENTING DECISION (EU) 2015/789

of 18 May 2015

as regards measures to prevent the introduction into and the spread within the Union of *Xylella fastidiosa* (Wells et al.)

Article 3

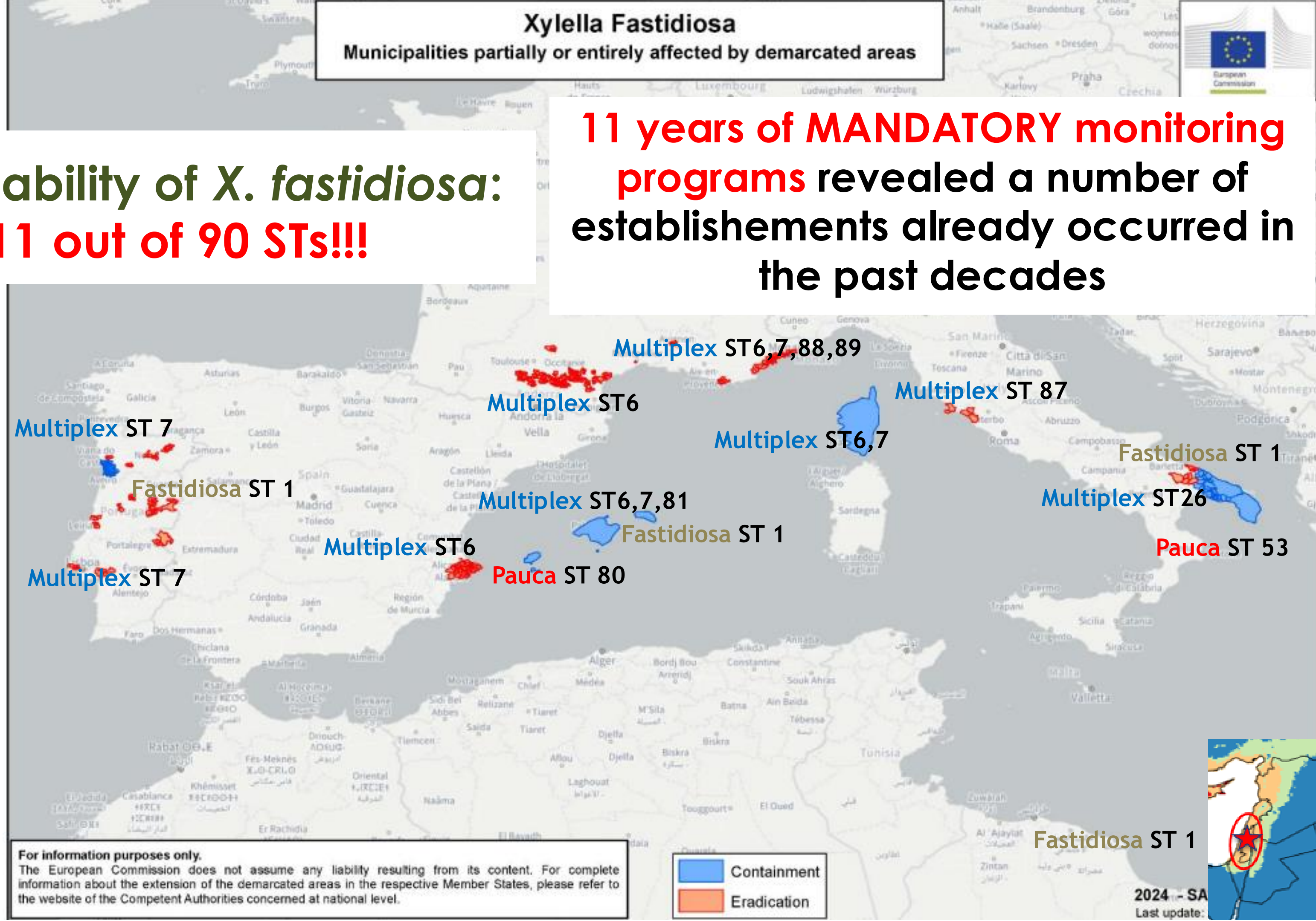
Surveys of the specified organism in the territories of the Member States

Member States shall conduct annual surveys for the presence of the specified organism in their territory on the specified plants.



Genetic variability of *X. fastidiosa*: in EU 11 out of 90 STs!!!

11 years of **MANDATORY** monitoring
programs revealed a number of
establishments already occurred in
the past decades

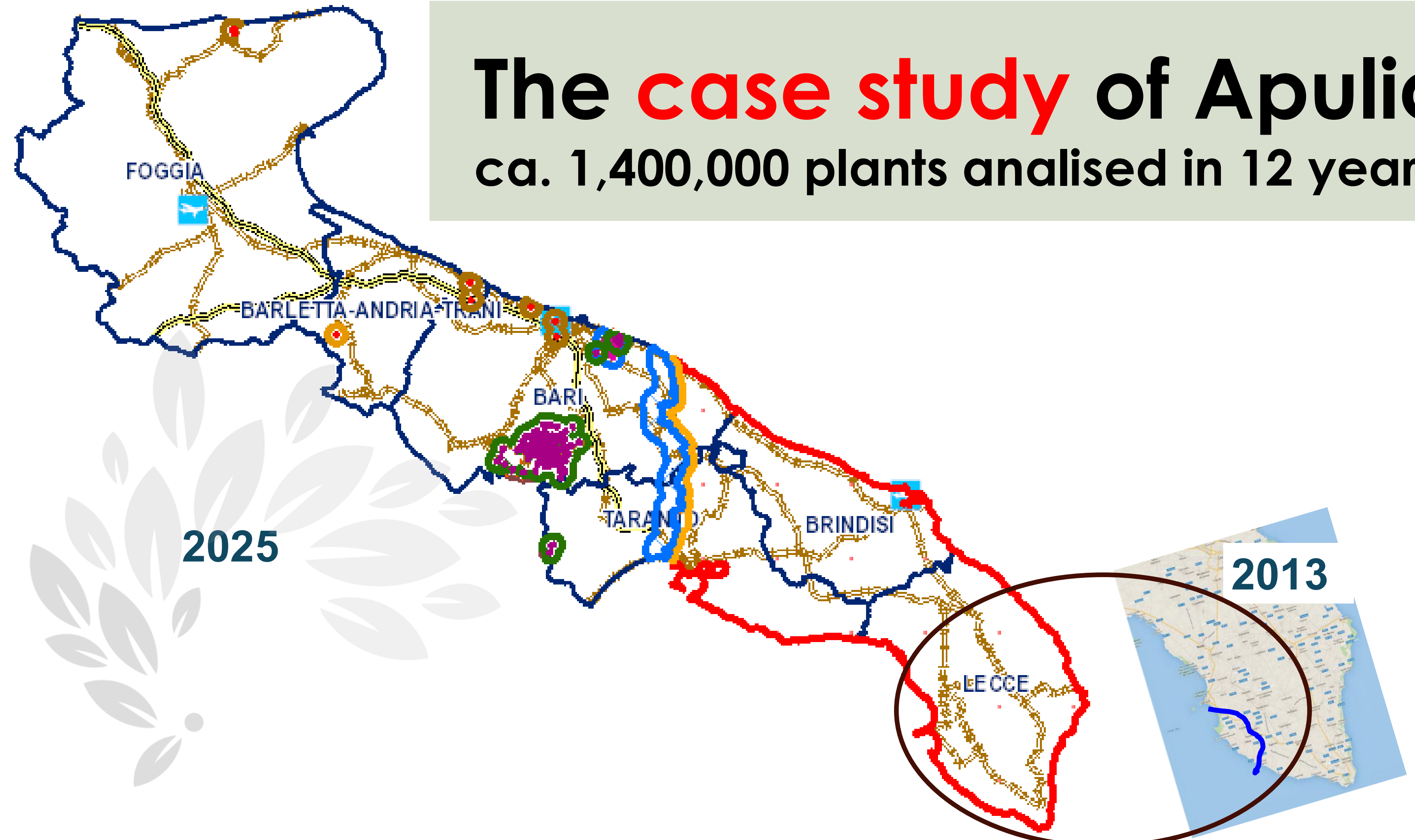


«pauca» was discovered following the phytosanitary investigations on the devastating OQDS, the **other subspecies** were found following mandatory monitoring campaigns



The **case study** of Apulia

ca. 1,400,000 plants analysed in 12 years!



2013-2025: annual monitoring campaigns

Green: negatives

Red: positives

(Yellow: visual inspection)

2013-2014

2014-2015

2016-2017

2017-2018

2018-2019

2019-2020

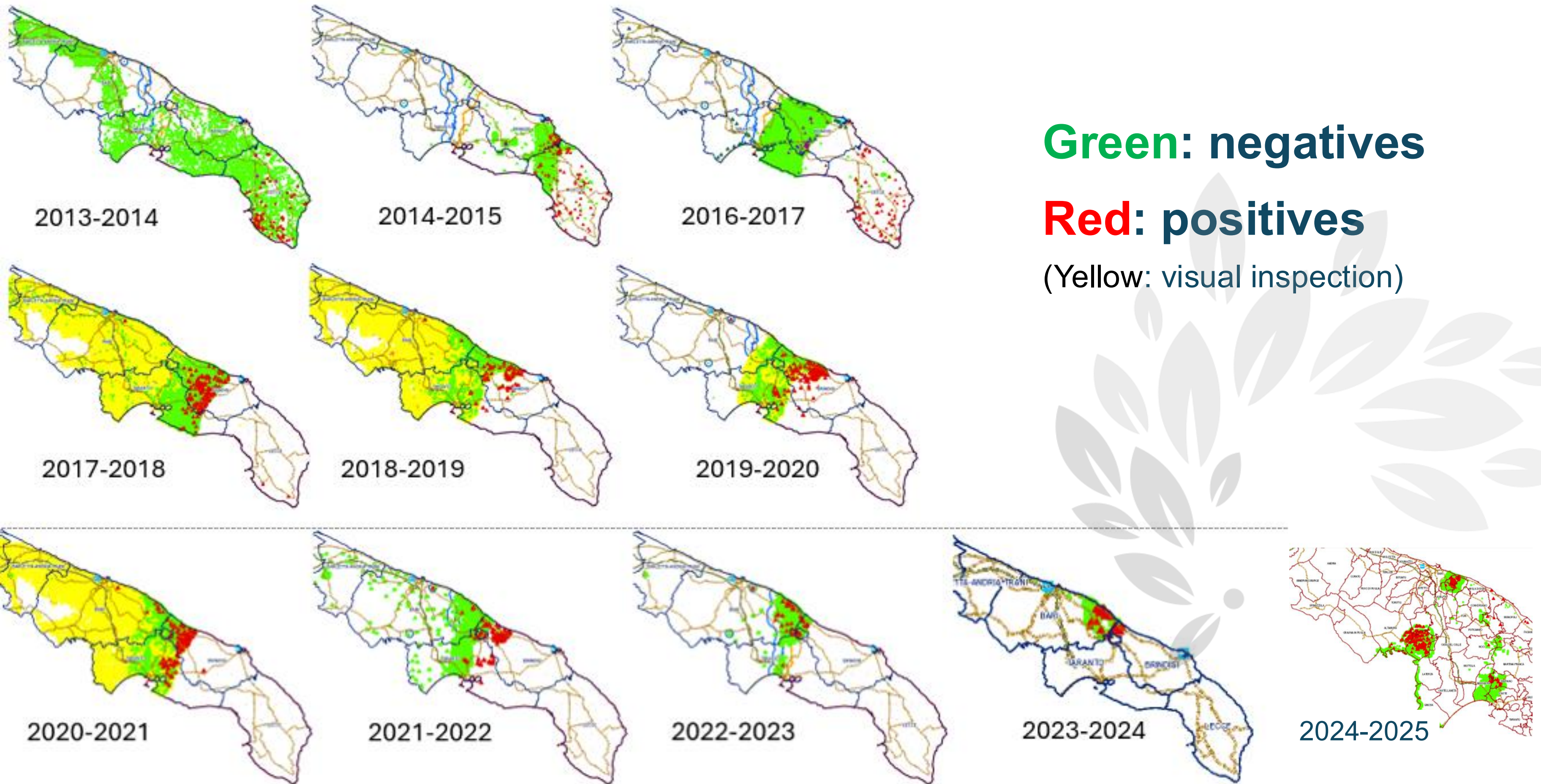
2020-2021

2021-2022

2022-2023

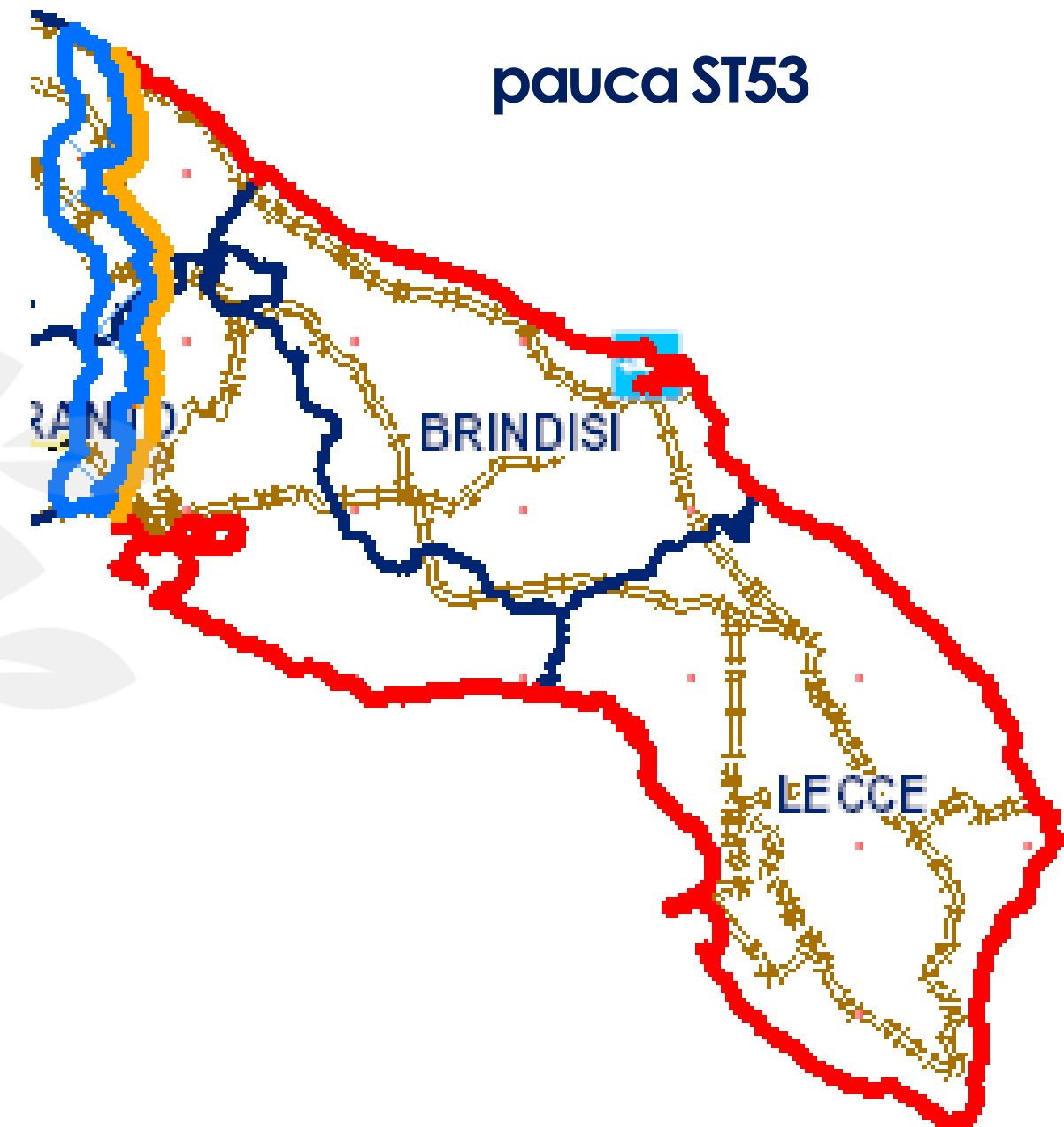
2023-2024

2024-2025



TWO SCENARIOS

UNDER CONTAINMENT



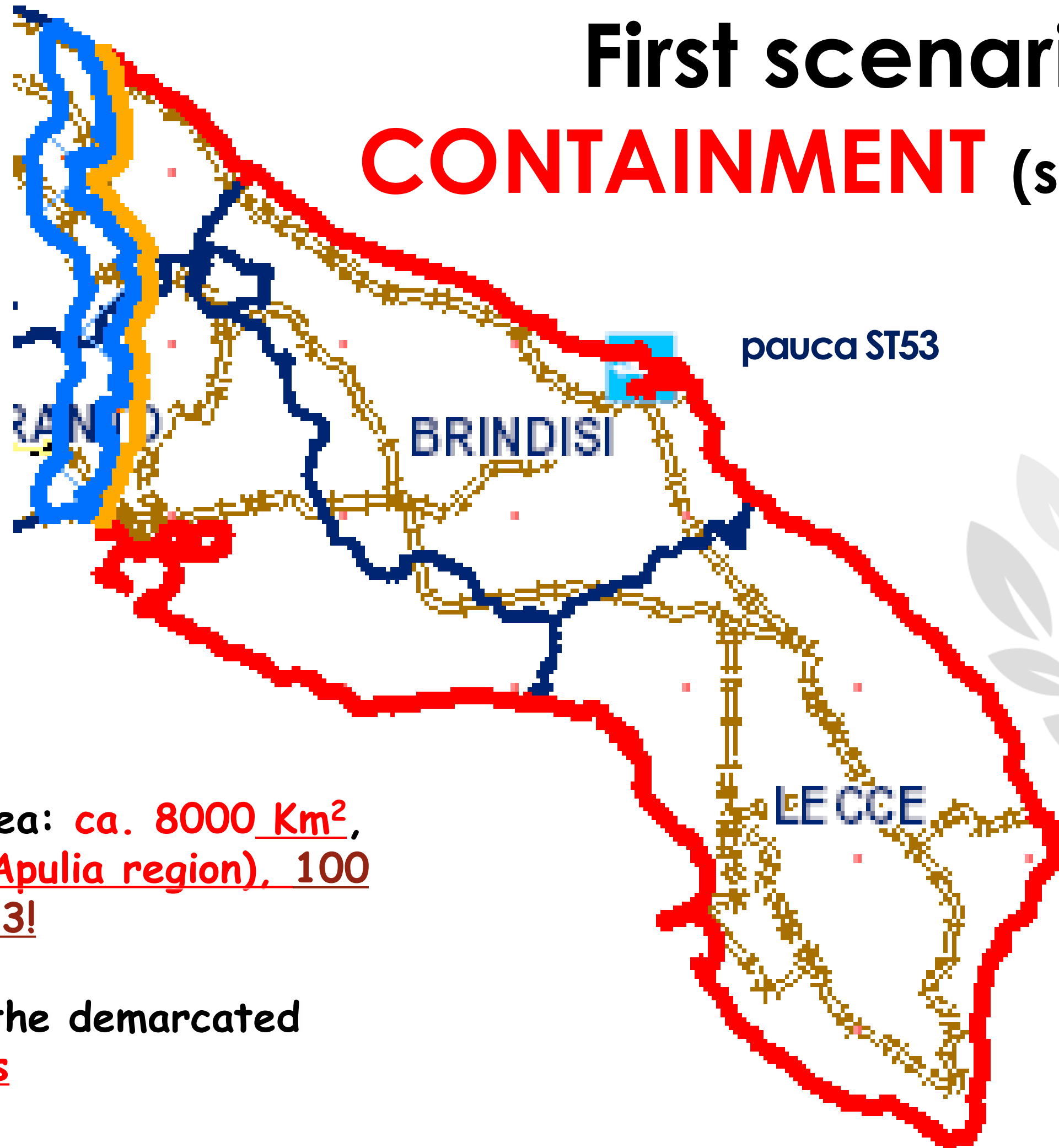
UNDER ERADICATION

- ★ multiplex ST26
- ★ fastidiosa ST1
- ★ pauca ST53



First scenario

CONTAINMENT (since 2015)



October 2013



Actual demarcated area: ca. 8000 Km², 160 Km (40% of the Apulia region), 100 times bigger than 2013!

Nr. of olive trees in the demarcated area: over 25 millions

ALLEGATO 1

Legenda

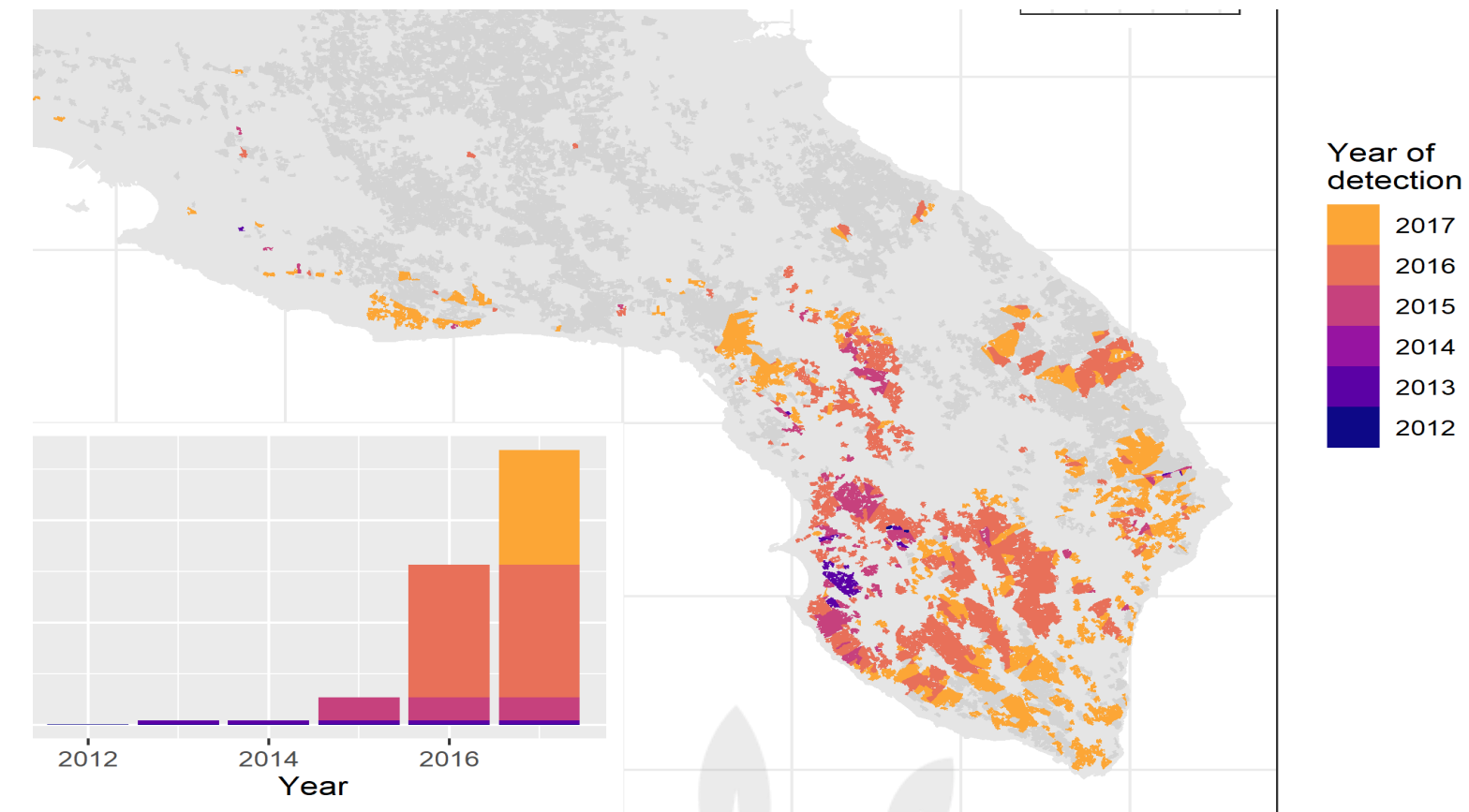
- Confini comunali
- Confini provinciali
- Focolai Canosa di Puglia, Monopoli, Polignano a Mare
- Zona Cuscinetto Canosa di Puglia, Monopoli e Polignano a Mare
- Zona Infetta
- Zona Cuscinetto
- Zona Containment

IL PRESENTE ALLEGATO E' COMPOSTO DA N. 1 FOGLIO
IL DIRIGENTE VICARIO DELLA SEZIONE
DOTT. LUISTROTTA

SUBSP. PAUCA: IMPACT PARTICULARLY SEVERE

SEVERAL MILLIONS OF TREES
DEAD OR SEVERELY COMPROMISED

A TOTAL **LOSS OF 132 MILLION**
EUROS/YEAR OF PROFITABILITY
AND A TOTAL LOSS OF 1,050,000 H
OF WORK/YEAR (*CALDERONI ET AL.,*
2025)








Xylella could cost Europe €20 billion, Wageningen University-led study finds

Researchers say the economic costs of Xylella could run to more than €20 billion.

RESEARCH ARTICLE

Impact of *Xylella fastidiosa* subspecies *pauca* in European olives

 Kevin Schneider,  Wopke van der Werf,  Martina Cendoya, Monique Mourits,  Juan A. Navas-Cortés,  Antonio Vicent, and Alfons Oude Lansink

PNAS April 28, 2020 117 (17) 9250-9259; first published April 13, 2020 <https://doi.org/10.1073/pnas.1912206117>

Edited by Charles Perrings, Arizona State University, Tempe, AZ, and accepted by Editorial Board Member Simon A. Levin March 3, 2020 (received for review July 16, 2019)



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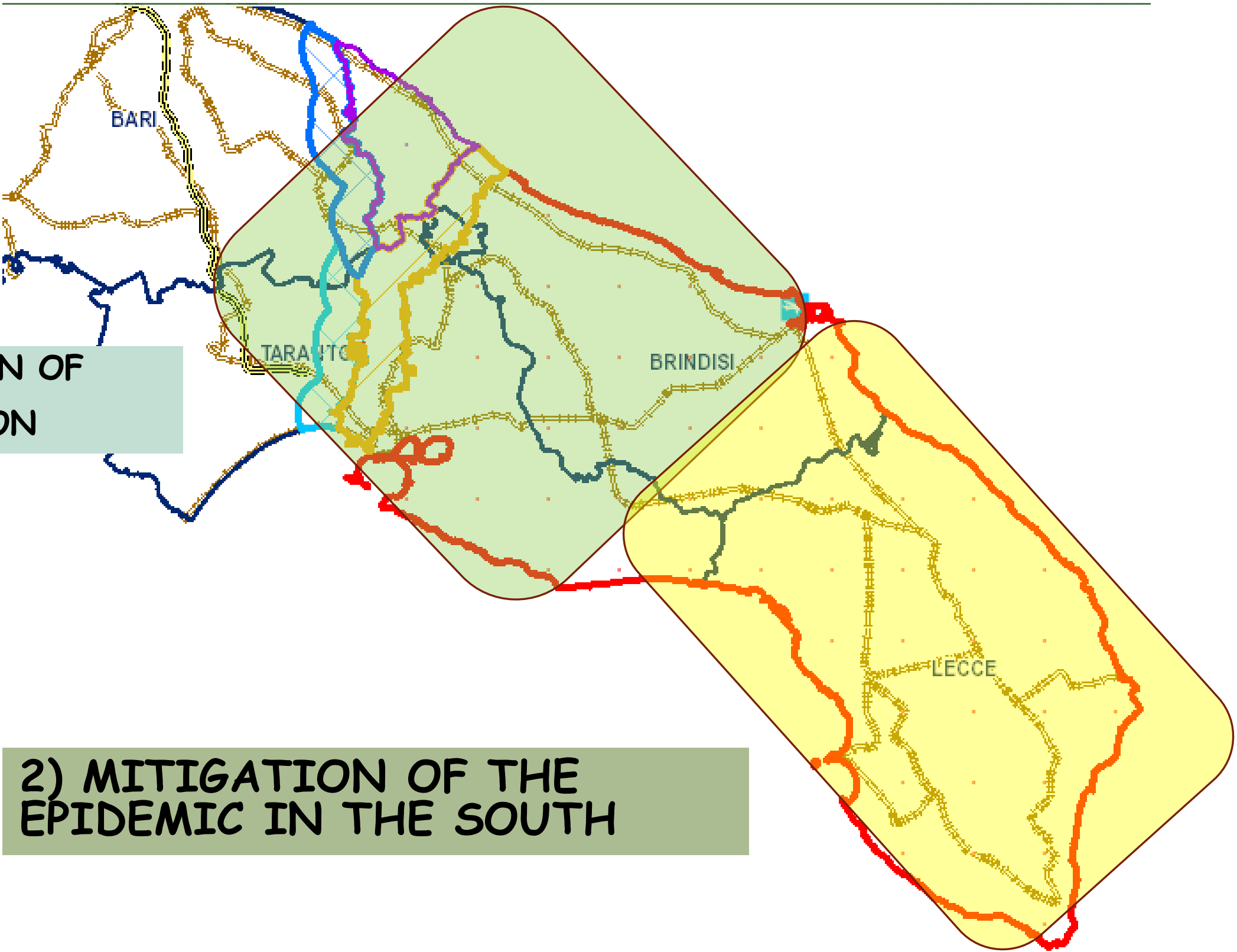
In addition:

- **Loss of land value**
- **impact on the attractiveness of tourist facilities**
- **impact on the climate**
- **growing phenomenon of abandonment**
- **increase in summer fires**
- **Etc.**



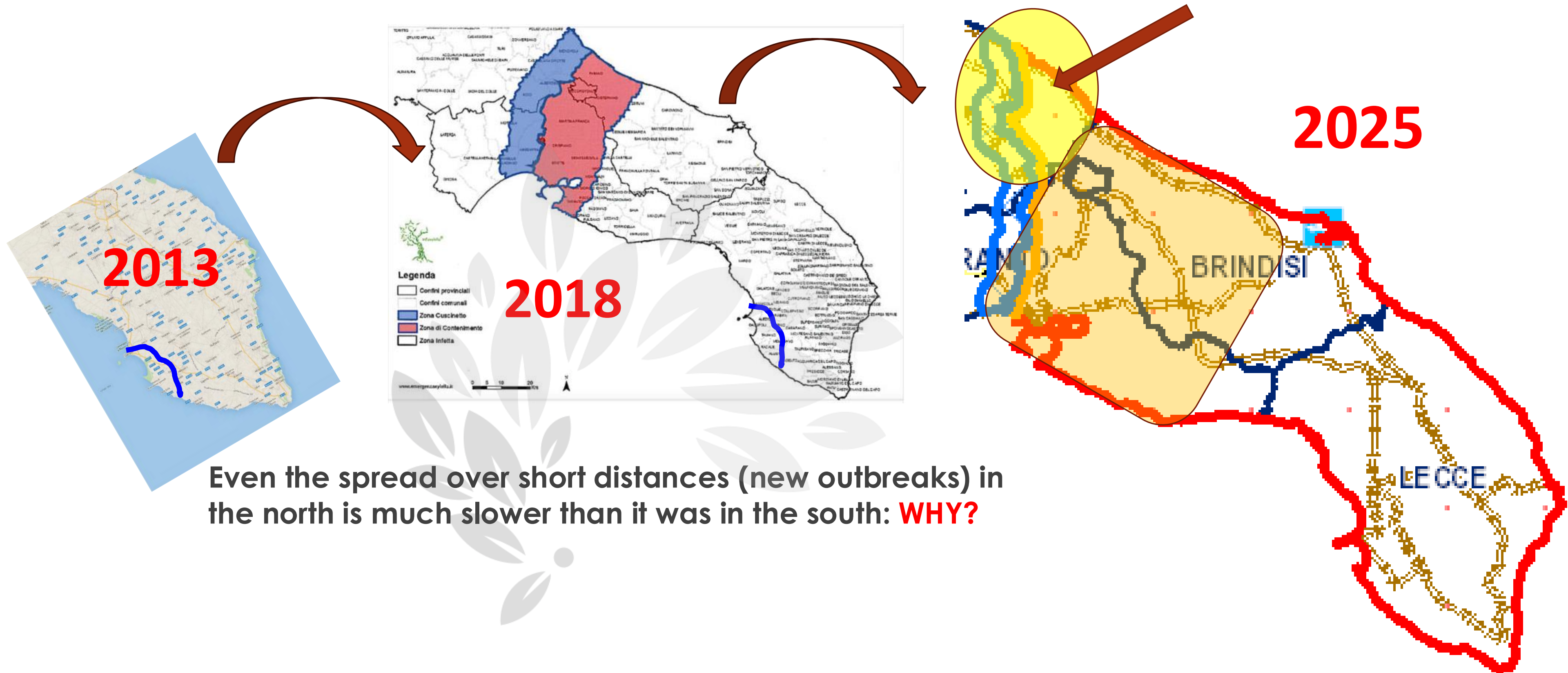
THE CURRENT STATUS: TWO DIFFERENT SCENARIOS

1) SIGNIFICANT SLOW DOWN OF
THE NORTHWARD DIFFUSION



2) MITIGATION OF THE
EPIDEMIC IN THE SOUTH

After 2018 significant slowdown in spread



SLOW DOWN OF THE NORTHWARD DIFFUSION: THE REASONS

- **Less suitable CLIMATE**
- Different land management (differences in agronomic and phytosanitary practices)
- APPLICATION OF CONTAINMENT MEASURES (monitoring and removal of infected plants, vector control) WHICH HAVE THE OBJECTIVE OF SLOWING DOWN THE SPREAD
- Less abundant VECTOR POPULATION



Estimated climatic suitability map for *X. fastidiosa* subsp. *pauca* according to a SDM ensemble model (EFSA 2019)

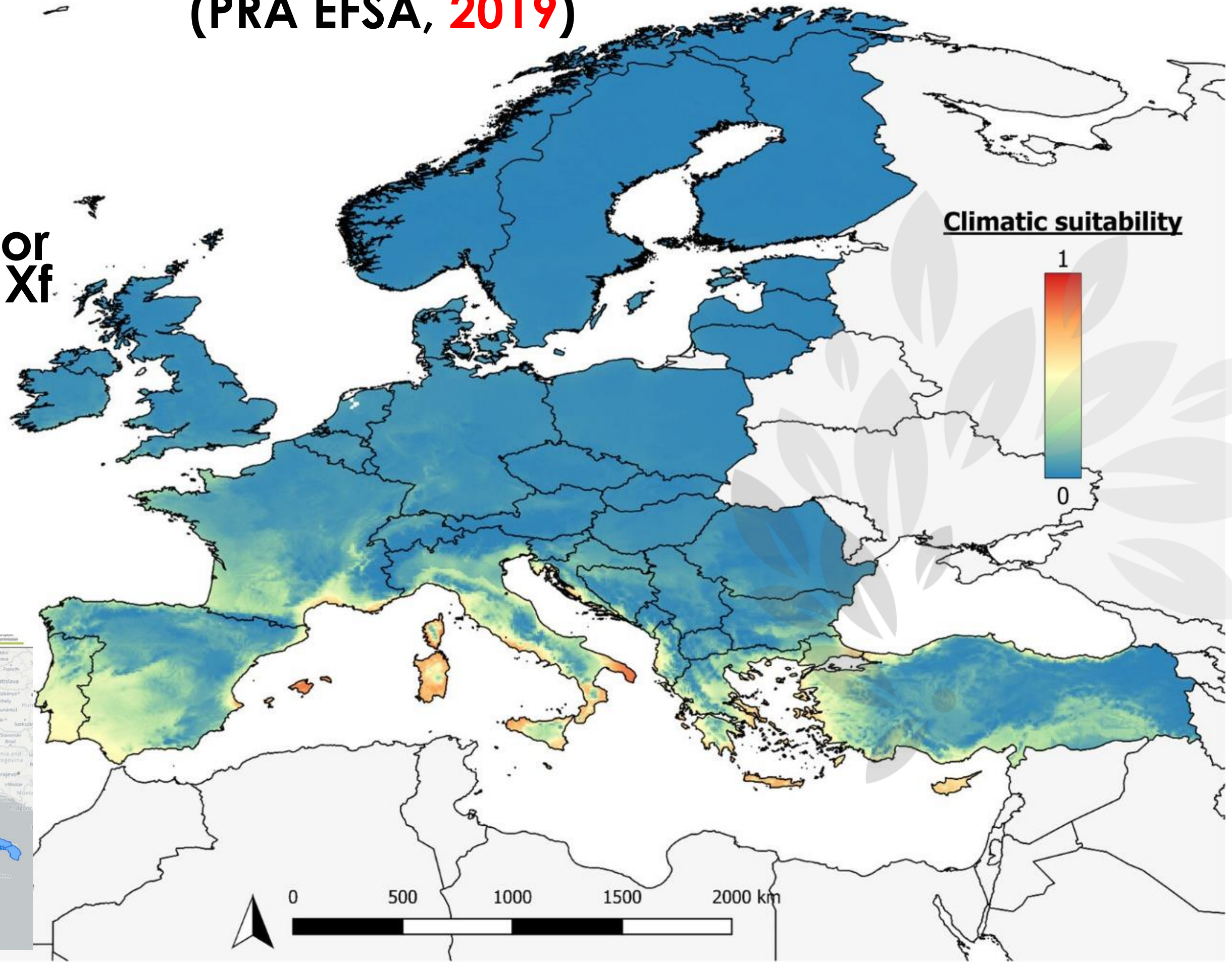
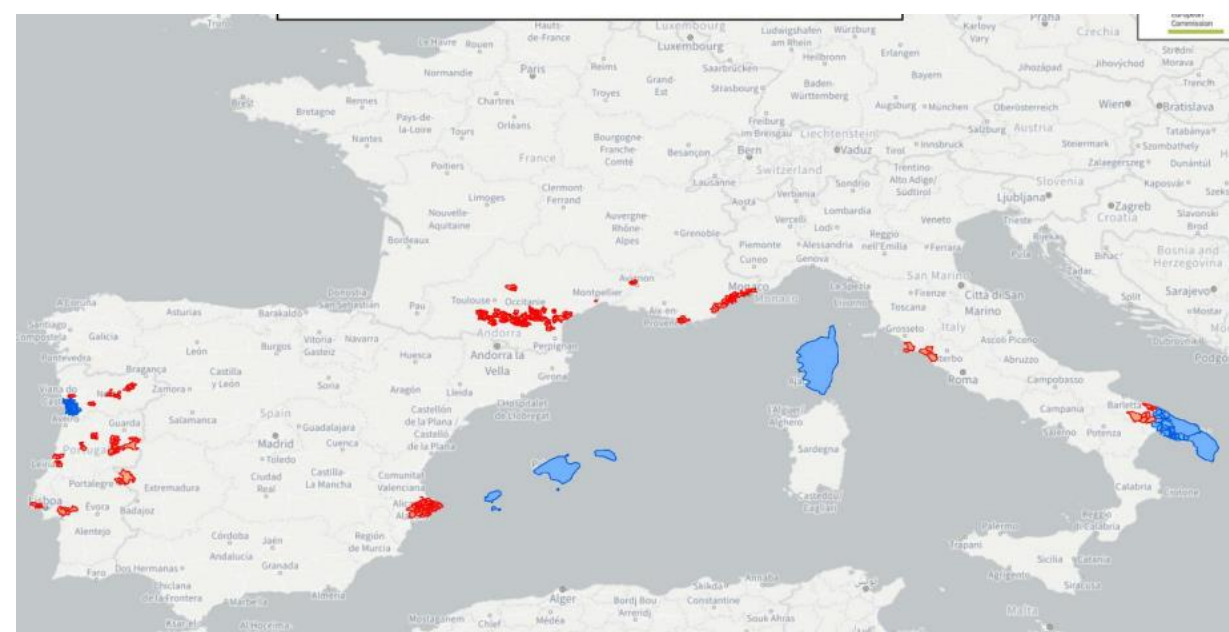


Estimated climatic suitability map for *X. fastidiosa* according to the SDM ensemble model

(PRA EFSA, 2019)

The climate: a key factor
for the epidemiology of Xf

**All the foci discovered
so far fall in areas with
high/moderate climatic
suitability**

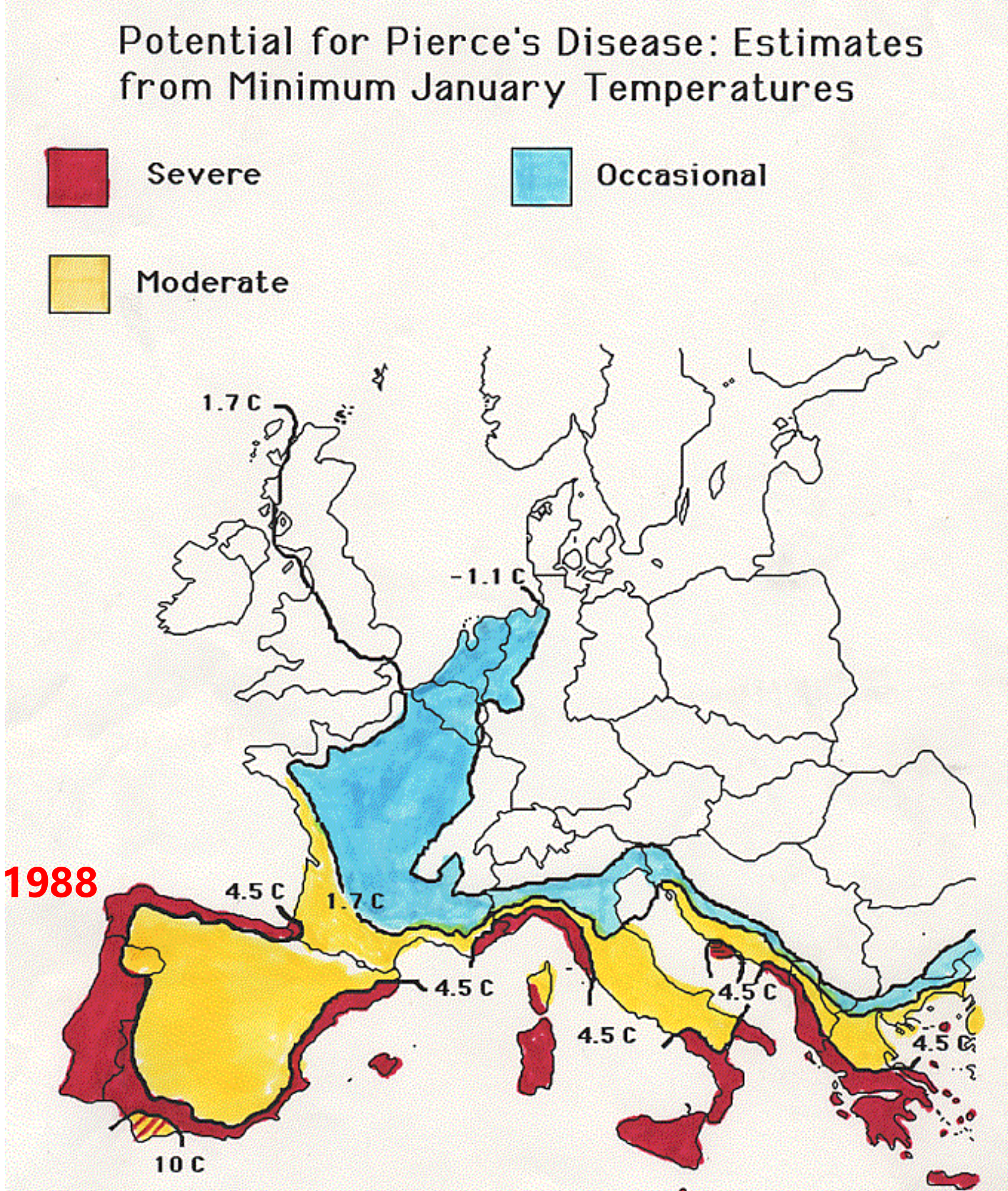
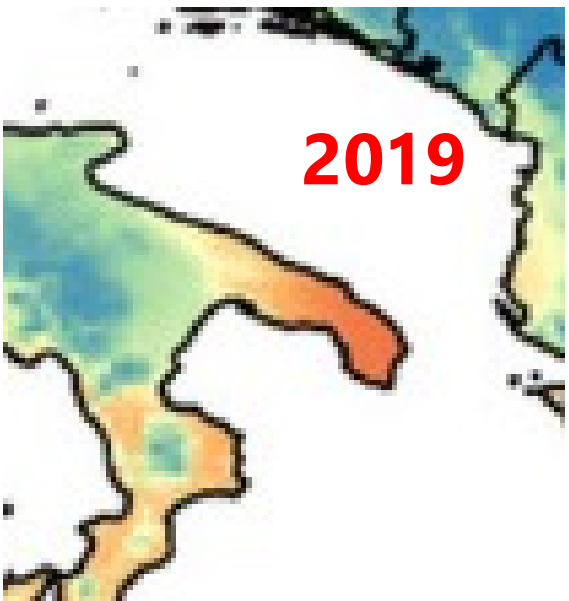


Prof. H.
Alexander Purcell



"I created this map for a talk that I gave in a symposium in Budapest (in 1988?)

The article in *Phytoma* (France) was its first official publication, but it was on the Internet before that."

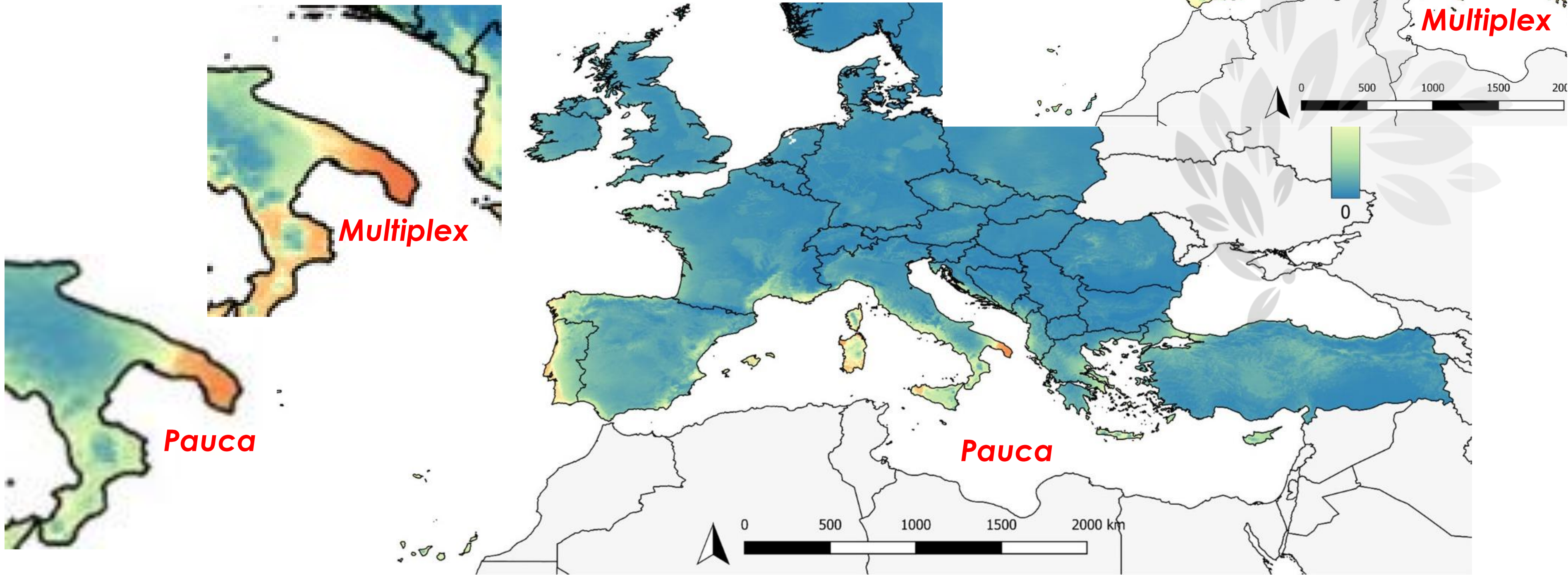


Sforza and Purcell, 2002, Phytoma La Défense des Végétaux 550:10-14

The climatic suitability varies among subspecies

***DIFFERENCES AMONG SUBSPECIES (AND STRAINS)
AFFECTS NORTHER BOUNDARIES***

***I.E.: PAUCA IS MORE WARM-LOVING THAN MULTIPLEX
(EFSA, 2019)***

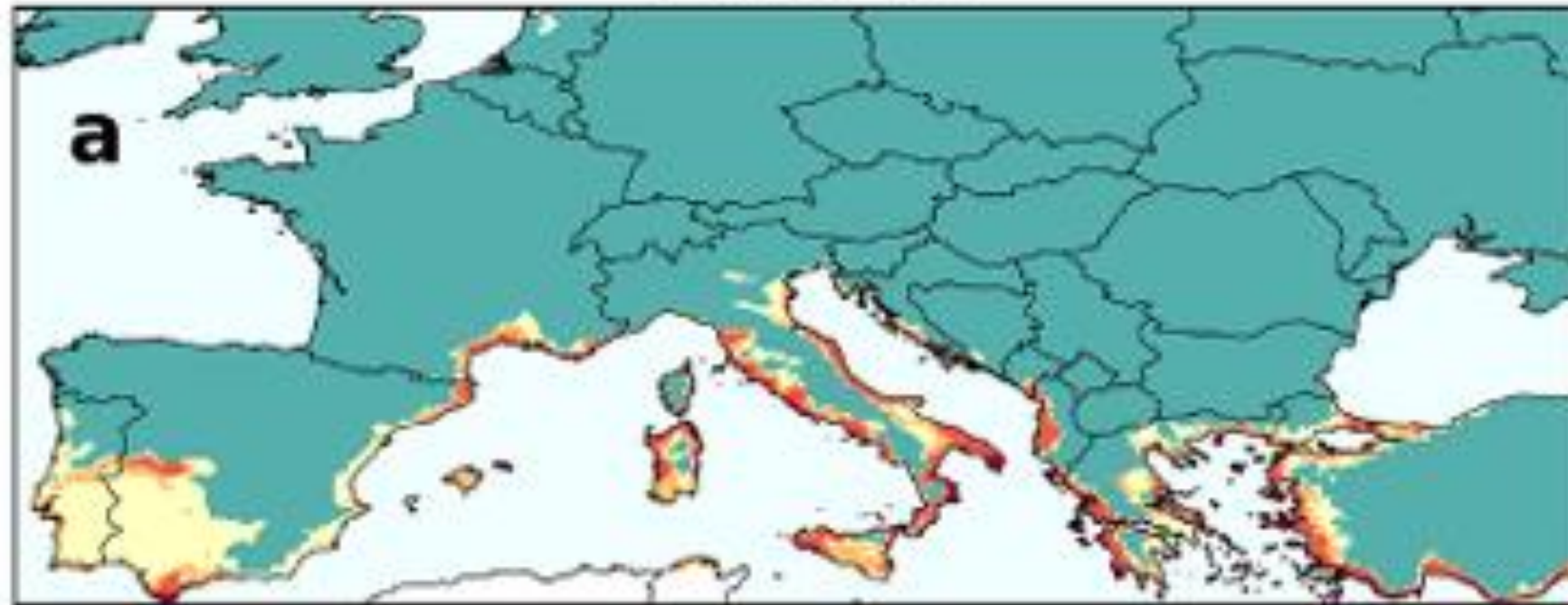


The effect of the climate change

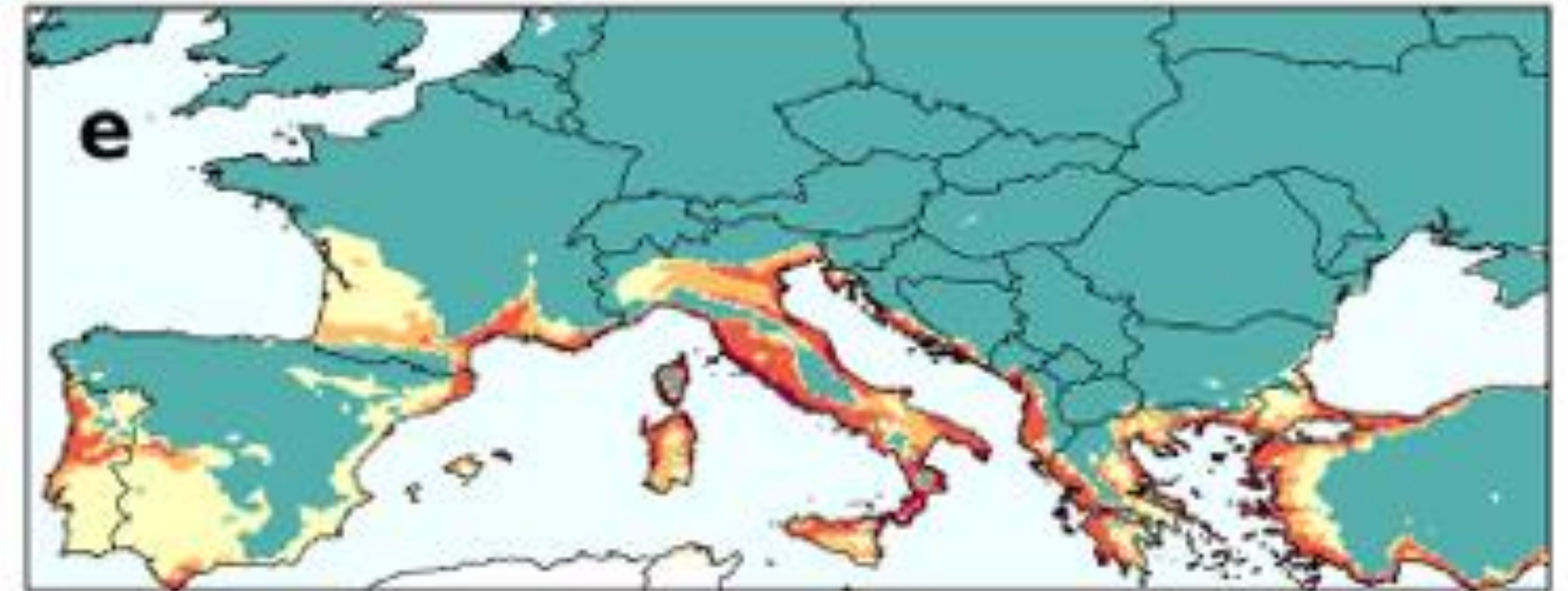
PD RISK MAPS UNDER DIFFERENT CLIMATE PROJECTIONS (GIMENEZ-ROMERO ET AL., 2024)



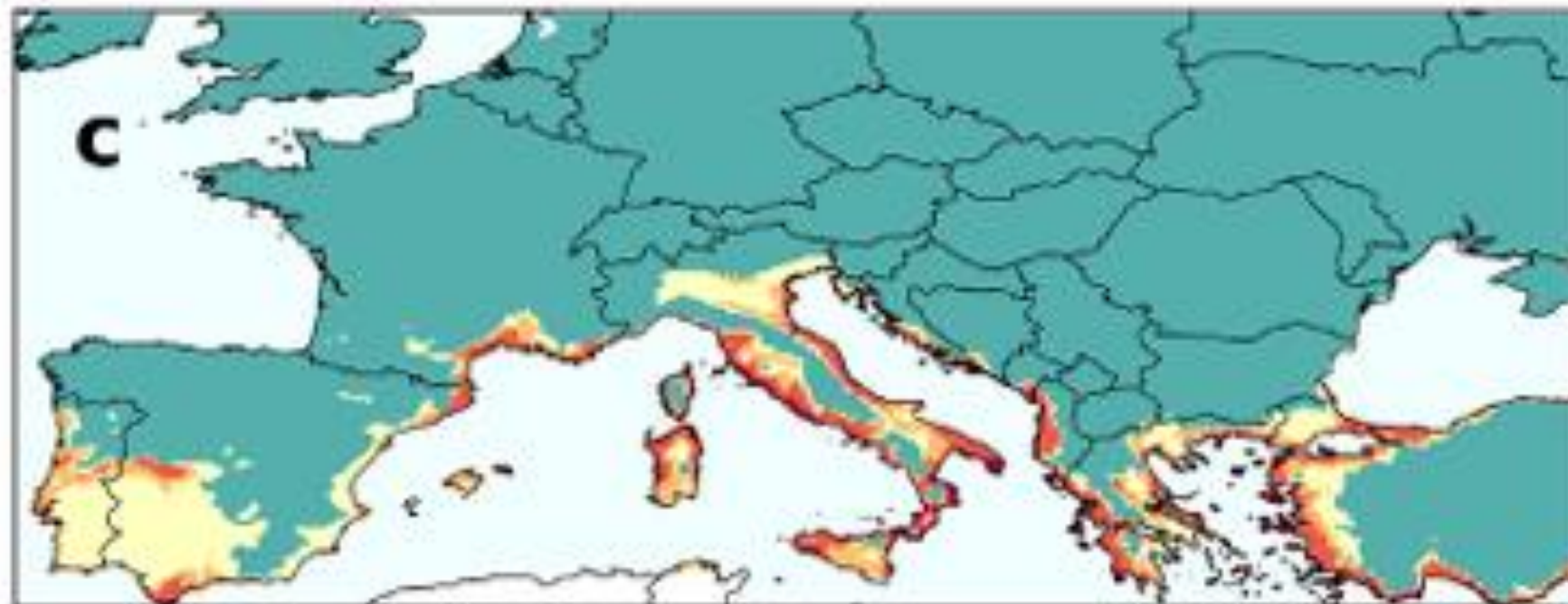
+1.5 °C



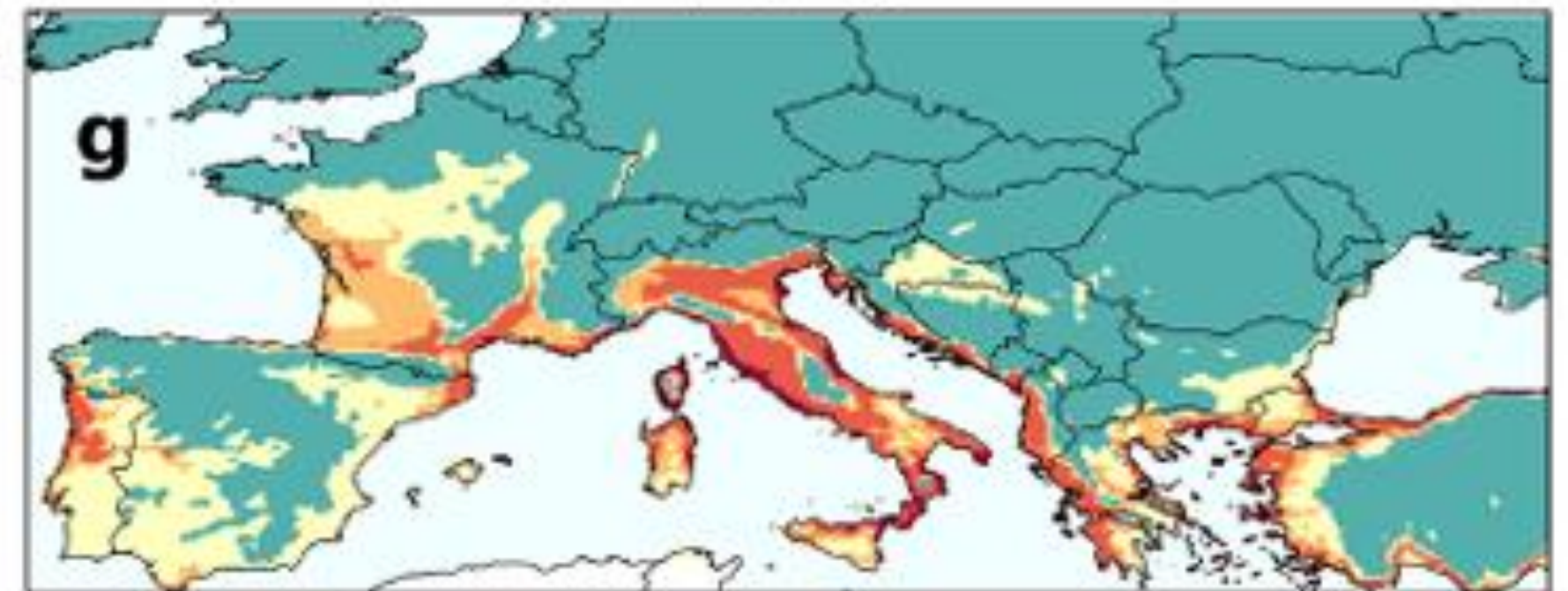
+3 °C



+2 °C



+4 °C



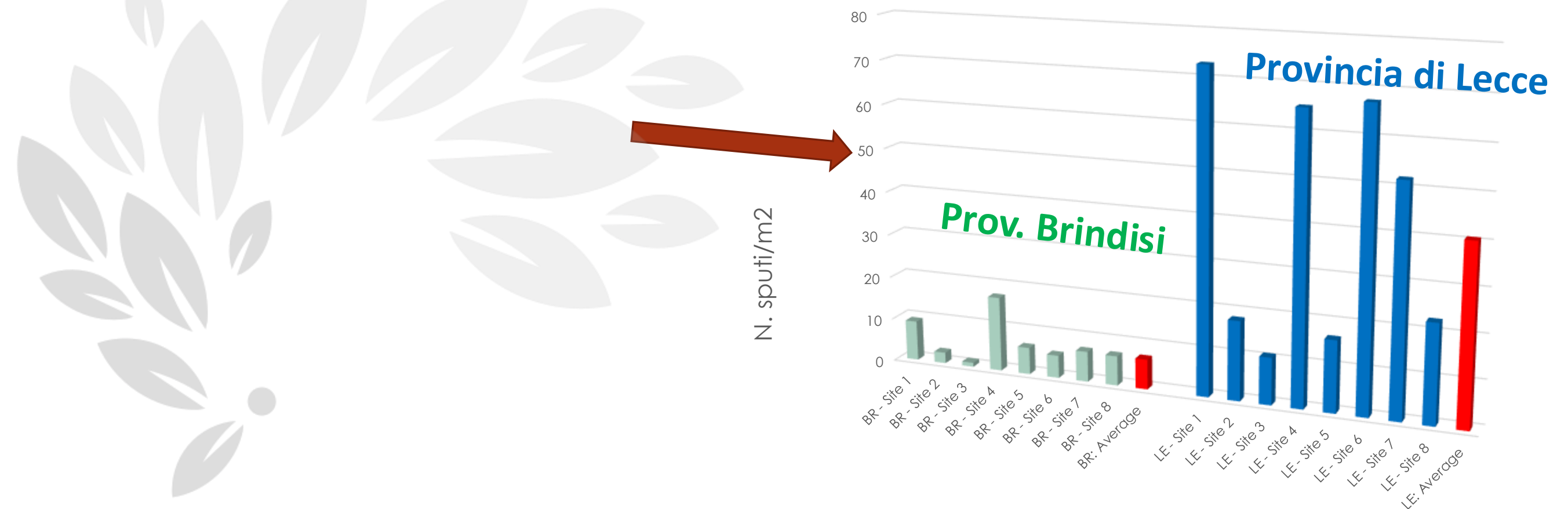
SLOW DOWN OF THE NORTHWARD DIFFUSION: THE REASONS

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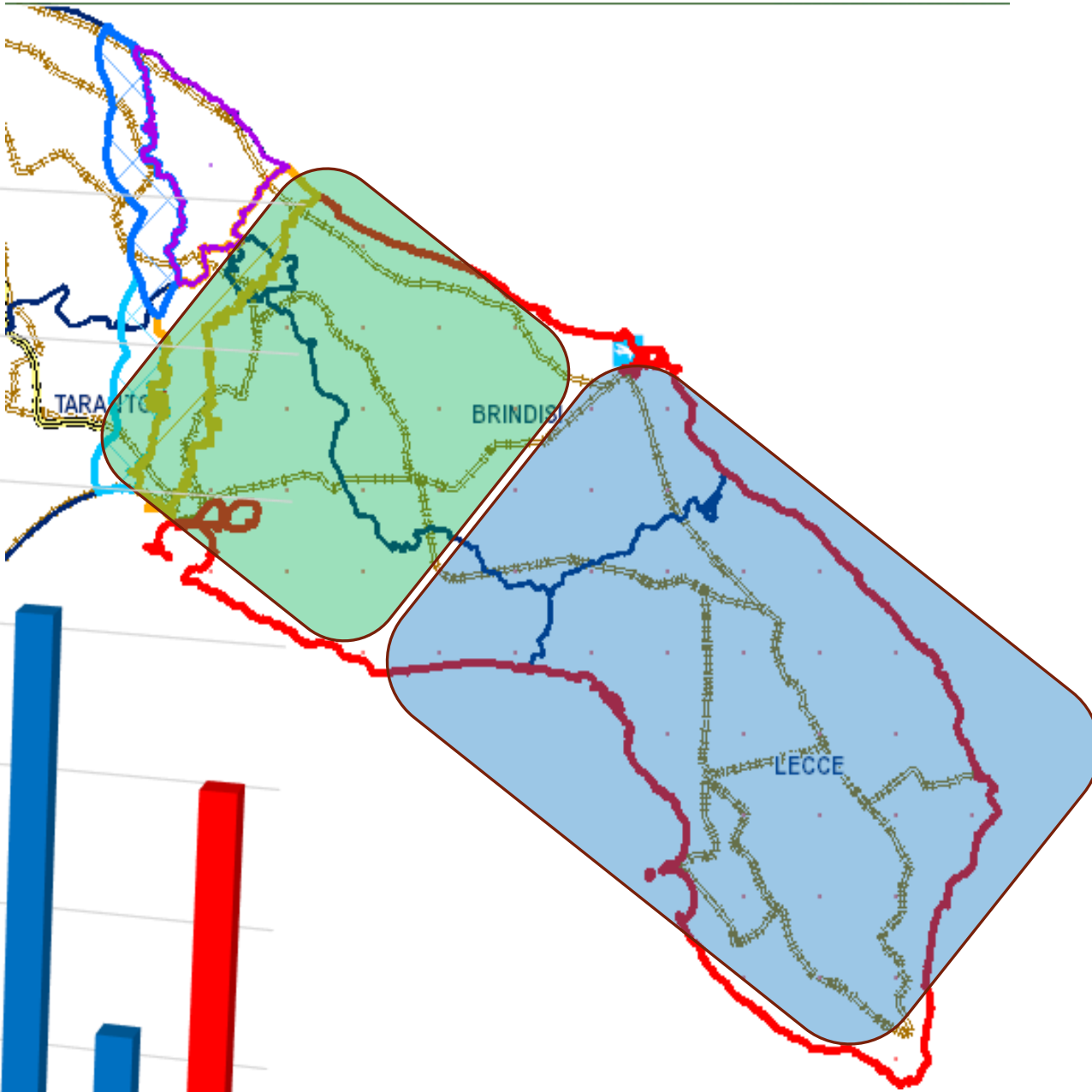
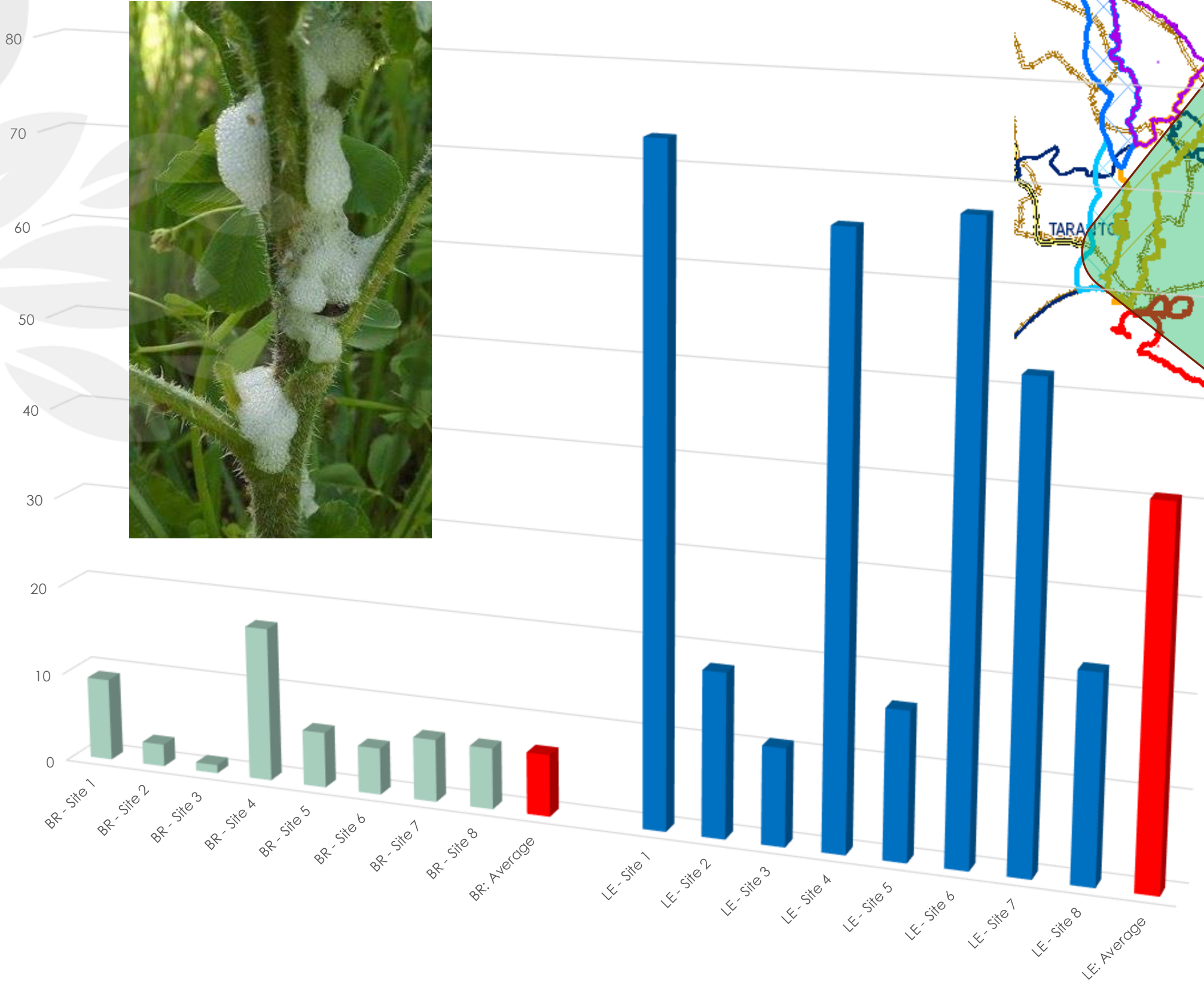


Population density of juvenile forms of Ps in 8 southern (blue) and 8 northern (green) sites. Data (n. individuals/m2) collected in spring 2023 when the peak of stage 4-5 of juvenile forms was reached.

In red the mean



N. spittles/m2

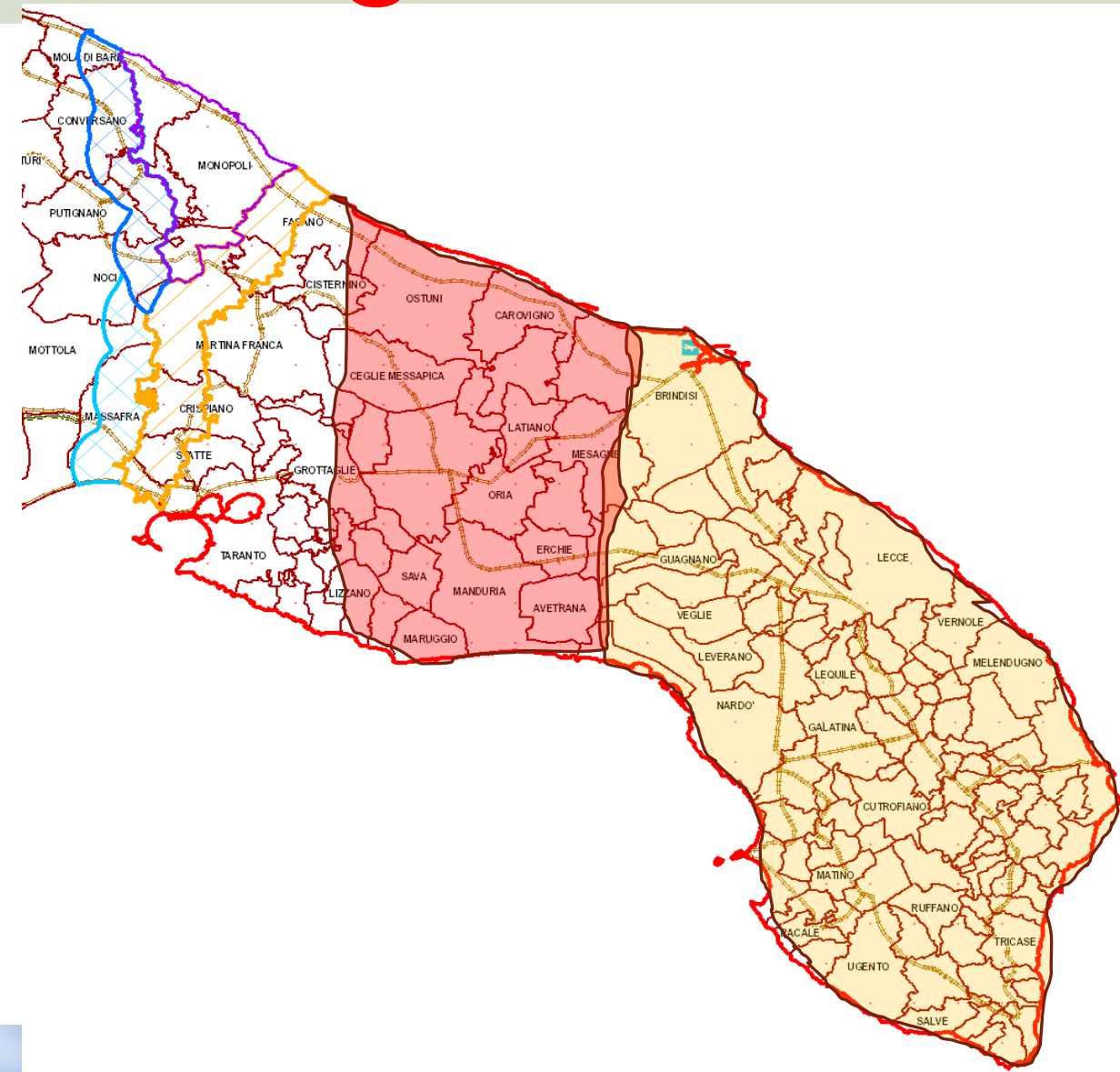


CONCLUSIONS

- A notable slow down of the northward diffusion has been observed in the past years
- Several factors seem to contribute to this matter
- A relevant role seems to play the lower vector population abundance and the different climatic suitability in comparison with the southern part of the Salento



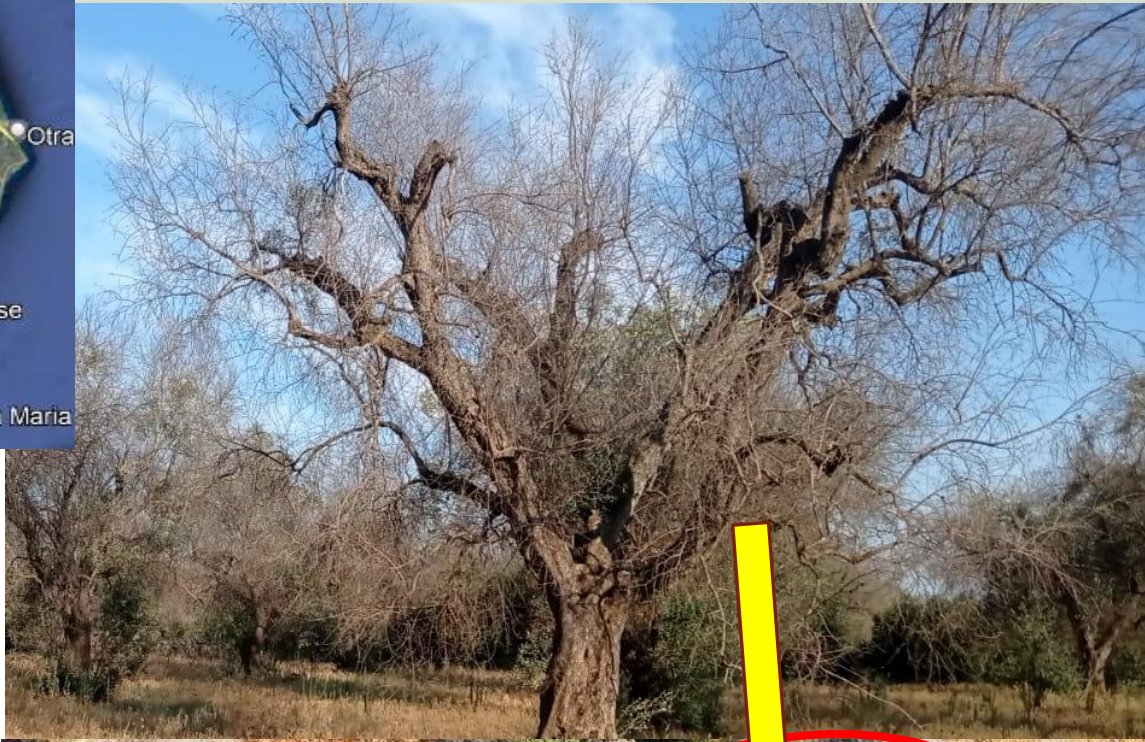
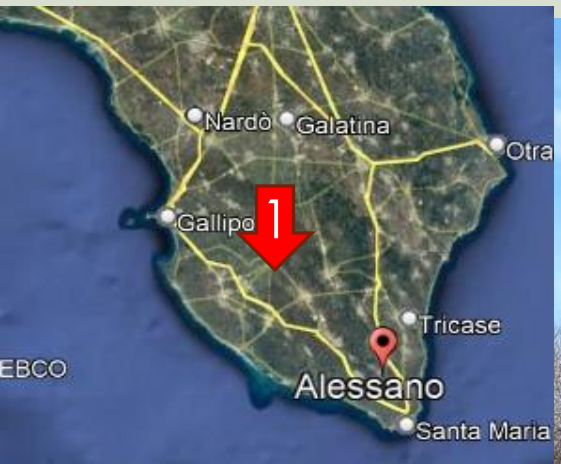
Mitigation of the epidemic in the south (South of Brindisi)



- New symptoms of dessication are milder and less frequent (see yellow area)
- in the last 3-4 years an increasing number of reports of symptoms remittance in survived trees has been recorded in adult trees (age up to 60-70 years old, not century old), more evident in cv. “Cellina” than in “Ogliarola”
- On the contrary, in the northern territory (red area) the disease is active and turning to the worst

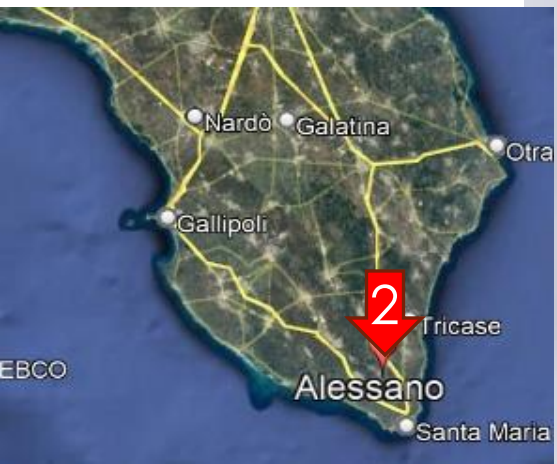


Few examples

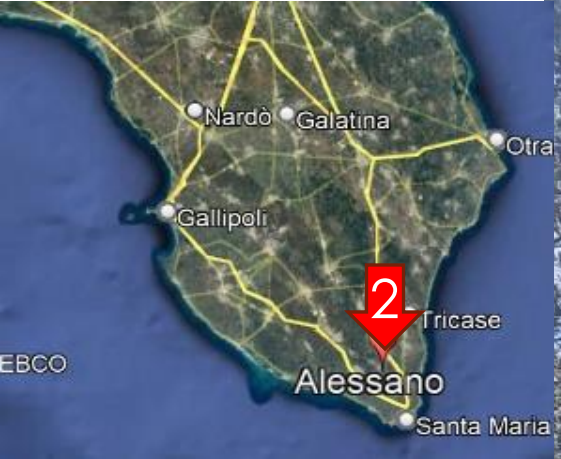


Cv. Cellina di Nardò

OCTOBER 2021



Cv. Cellina di Nardò

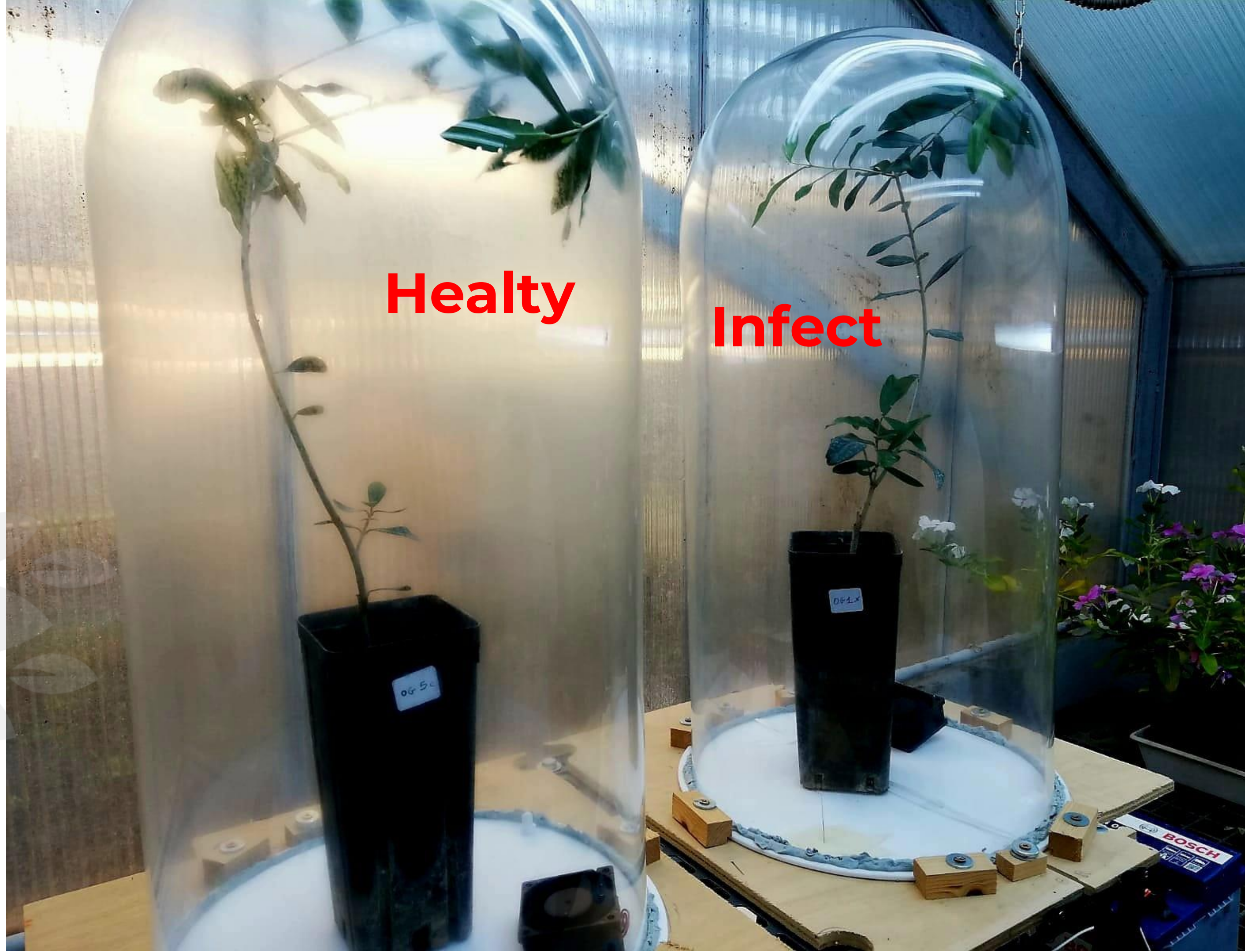


JUNE 2023



**"Partial improvement"
is not synonymous
with "recovery"**

Reduced transpiration



IN RESISTANT GERMPLASM THE BENEFICIAL EFFECTS SEEM TO BE HIGHER

Leccino (resistant), Ugento (le), august 16, 2023



What has changed?

Occurrence of relevant mutations in the *Xf* population?

As 12 years ago, all new isolated strains harbor ST53:

NO EMERGENCE OF NEW/DIFFERENT "ST"
REGISTERED SO FAR.

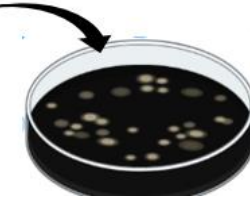
NO EVIDENCES OF RELEVANT GENETIC
MUTATIONS

NO EVIDENCES OF RELEVANT GENETIC MUTATIONS

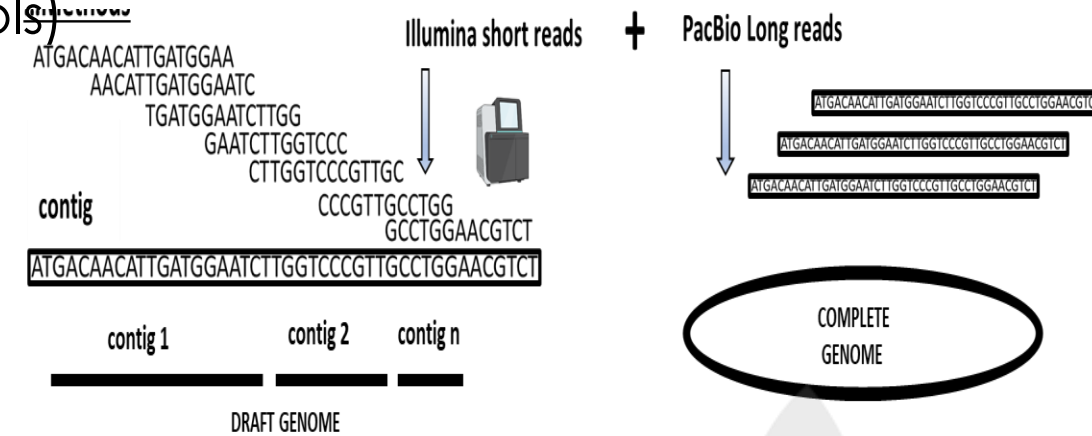
Changes in the bacterial genome?



X. fastidiosa



NGS: on a very limited n. of isolates (1 whole & 3 draft genomes reconstructed using different tools)



MLST typing: all isolates - **ST53**

- ✓ Mapping the newly recovered whole genome - reads (13,725,672) versus De Donno reference: 99,54% mapped with **homogeneous coverage**

So far, the only difference found with the reference strain «De Donno» is a deletion of 38nt repeat sequence

**However it corresponds to an intergenic region >>>
no effect on gene function!**

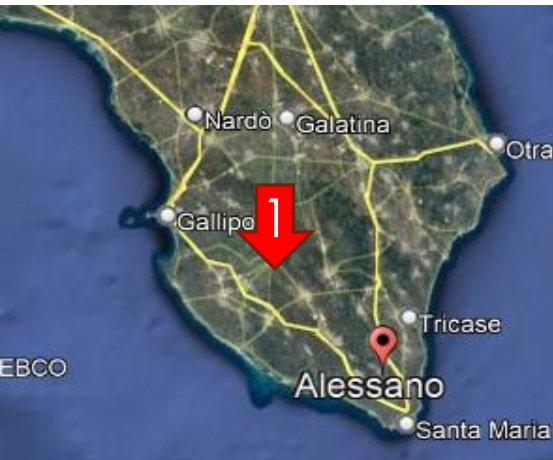
Wath has changed?

Reduction of bacterial titer in infected plants?

NO EVIDENCES OF SIGNIFICANT
REDUCTION

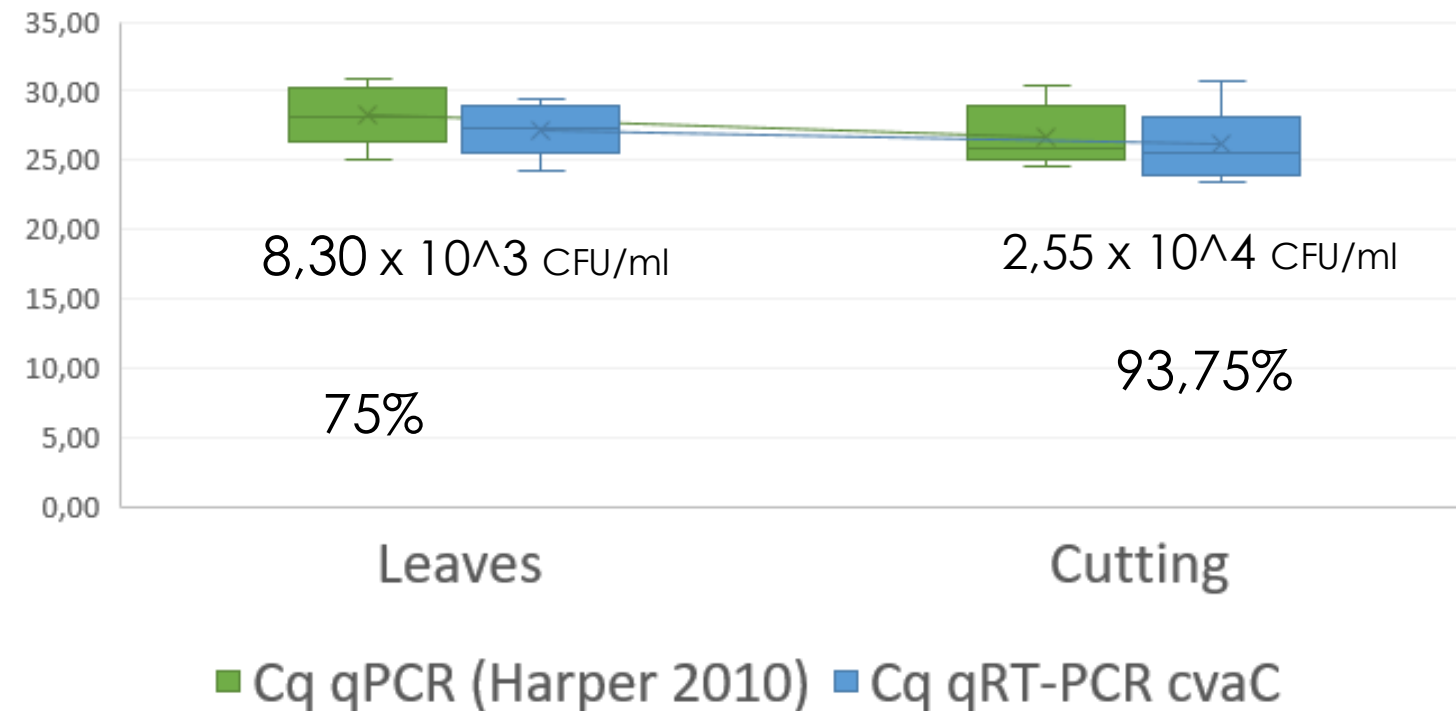


MONITORING THE XF-STATUS OF THE PLANTS

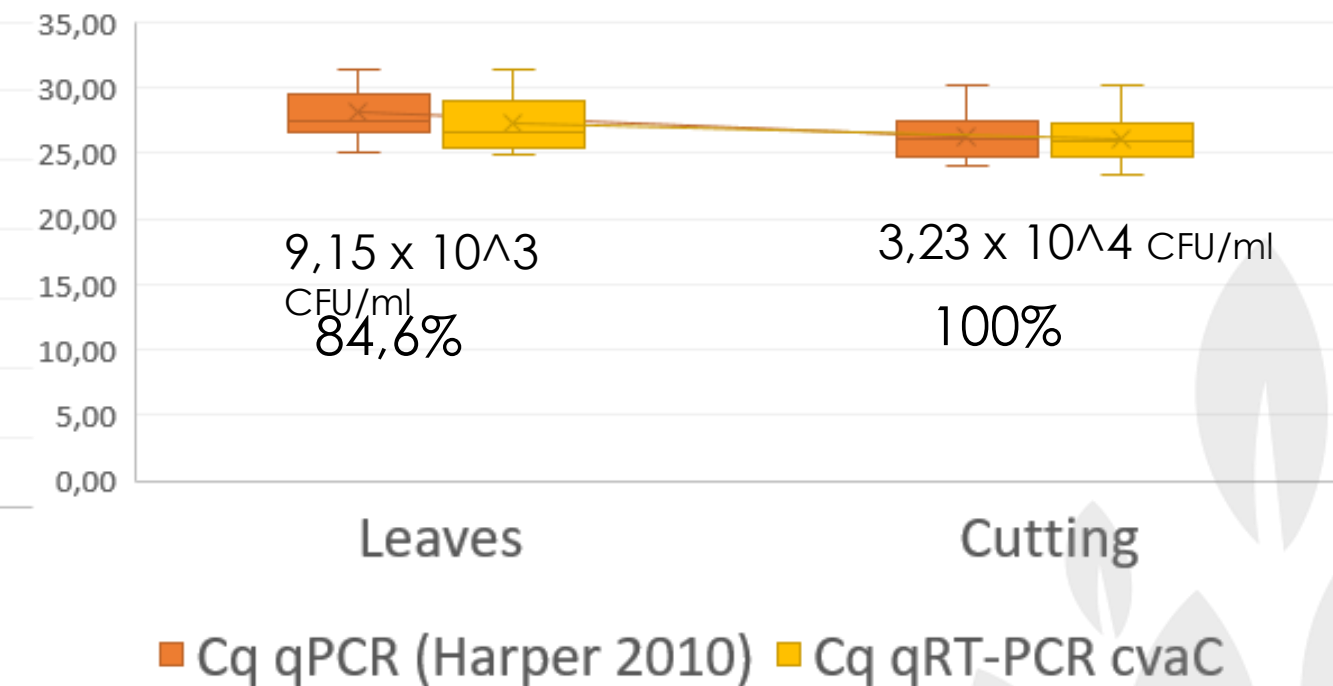


Sampling and testing a representative n. of trees: standard detection & estimation of the bacterial multiplication rate by quantification of the bacterial RNA (encoding a microcin-like protein highly expressed in infected trees)

Detection and quantification of Xf in trees showing symptom remission



Detection and quantification of Xf in symptomatic control trees



Drastic reduction of reservoir of inoculum: the reasons

Massal reduction of vegetation with high concentration of bacterium due to the infection



Drastic reduction of reservoir of inoculum: the reasons

Massal reduction of vegetation with high concentration of bacterium due to the infection



Drastic reduction of reservoir of inoculum: the reasons

Significant removal of severely damaged olive groves, often replaced (over 3 milions of plants) with resistant cultivars (low bacterium titre when infected)



Drastic reduction of reservoir of inoculum: the reasons

High frequency of fires in olive groves



Drastic reduction of reservoir of inoculum: the reasons

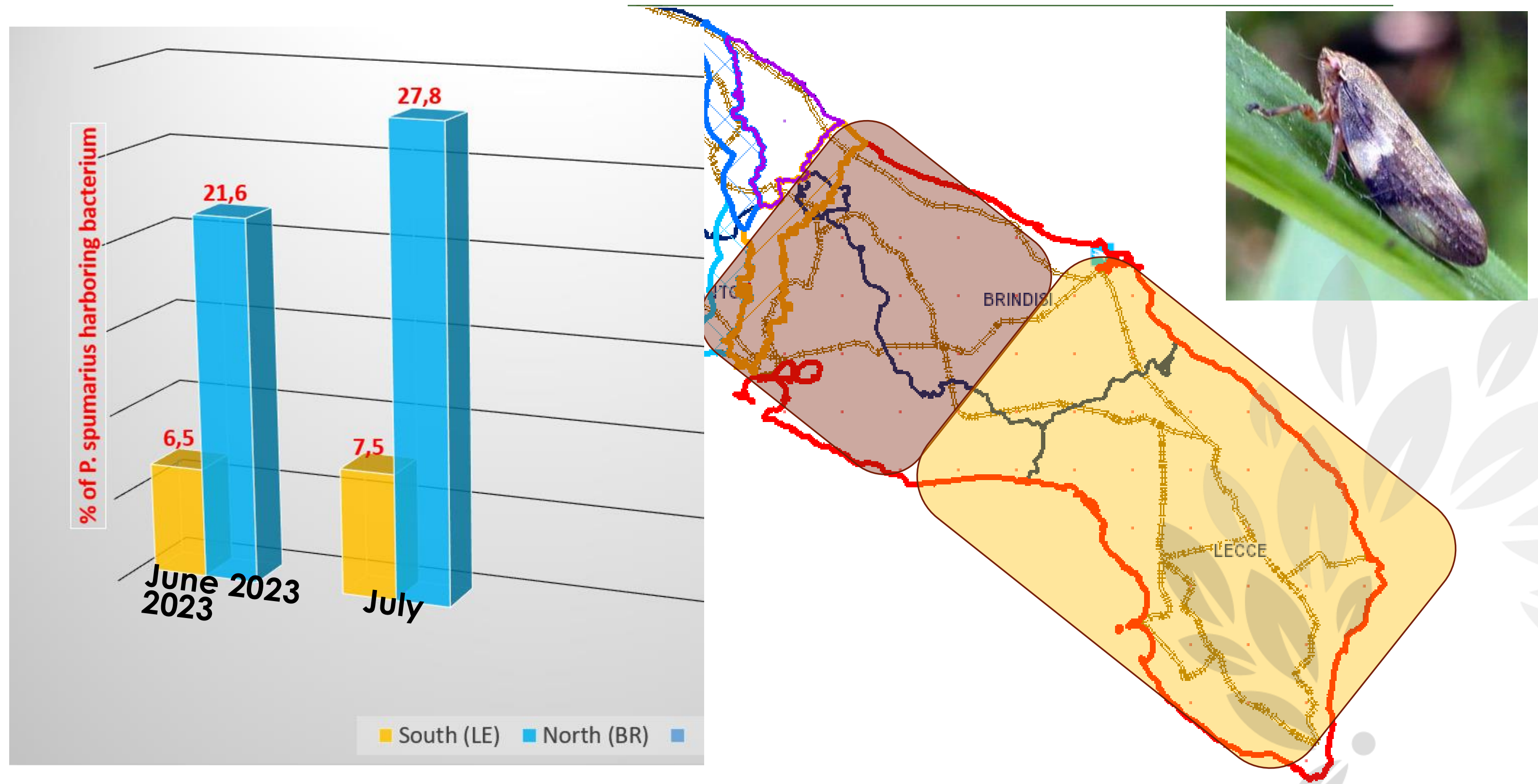
- Massal reduction of vegetation with high concentration of bacterium due to the infection
- Significant removal of severely damaged olive groves, often replaced (over 3 millions of plants) with resistant cultivars (low bacterium titre when infected)
- High frequency of fires in olive groves
- **Proibition of planting higly susceptible cultivars**

Consequences: Drastic reduction of infected vectors

A decorative graphic of an olive branch with several leaves, rendered in a light gray color, positioned in the bottom right corner of the slide.

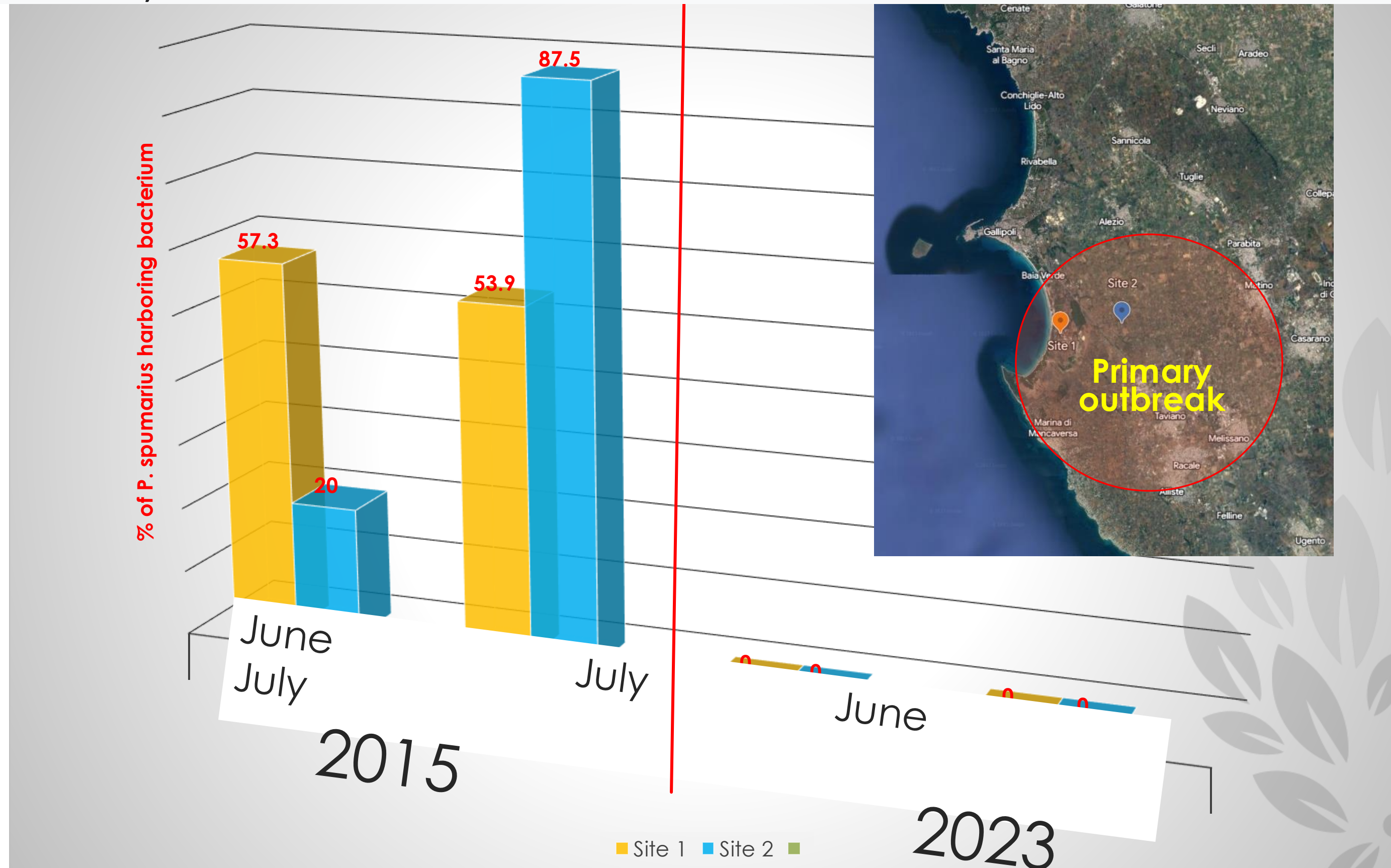
Drastic reduction of infected vectors

Comparison of % of infected vectors (qT-PCR) between southern (**6 sites, 173 specimens**) and northern (**5 sites, 141 specimens**) areas of the infected zone



Drastic reduction of infected vectors

Comparison of the percentage of positive vectors in different years in two sites close to the primary outbreak (80 specimens in 2023)



Mitigation of the epidemic in the south

CONCLUSIONS

Reservoir of inoculum drastically reduced!



Vectors efficiency dropped



Strong reduction of superinfections

In the meanwhile:

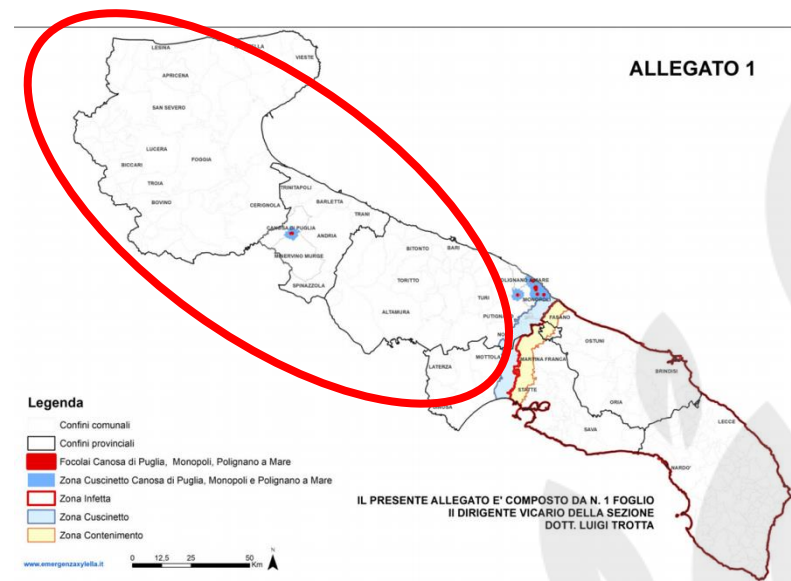
- 1) no significant mutations are recorded in the bacterial population (still only ST53)
- 2) the titre of bacterium in infected plants does not show significant reduction in sensitive cultivars

MAJOR ROLE OF THE SUPERINFECTIONS?



Second scenario: **ERADICATION**

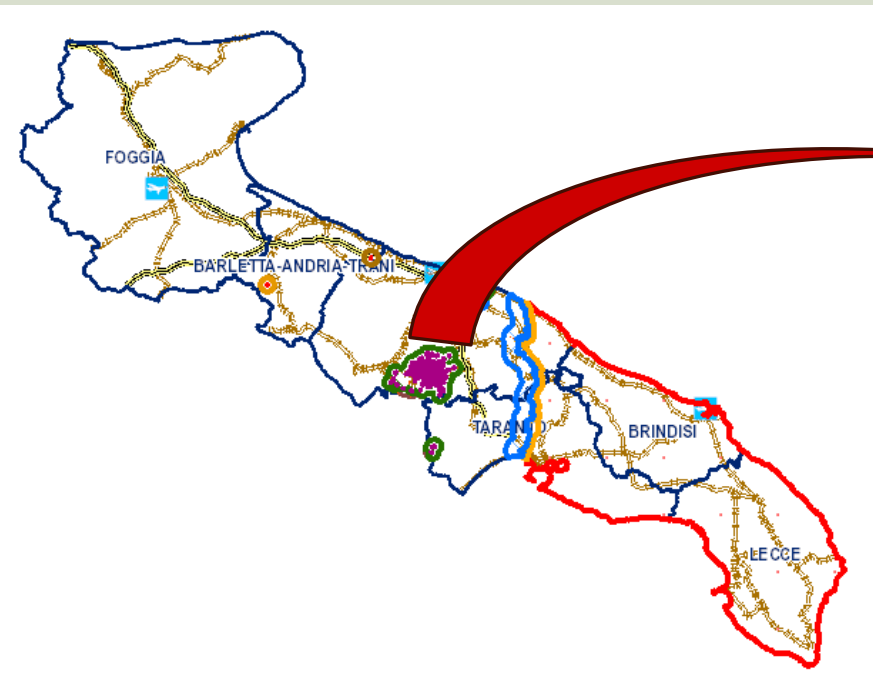
(2024-2025)



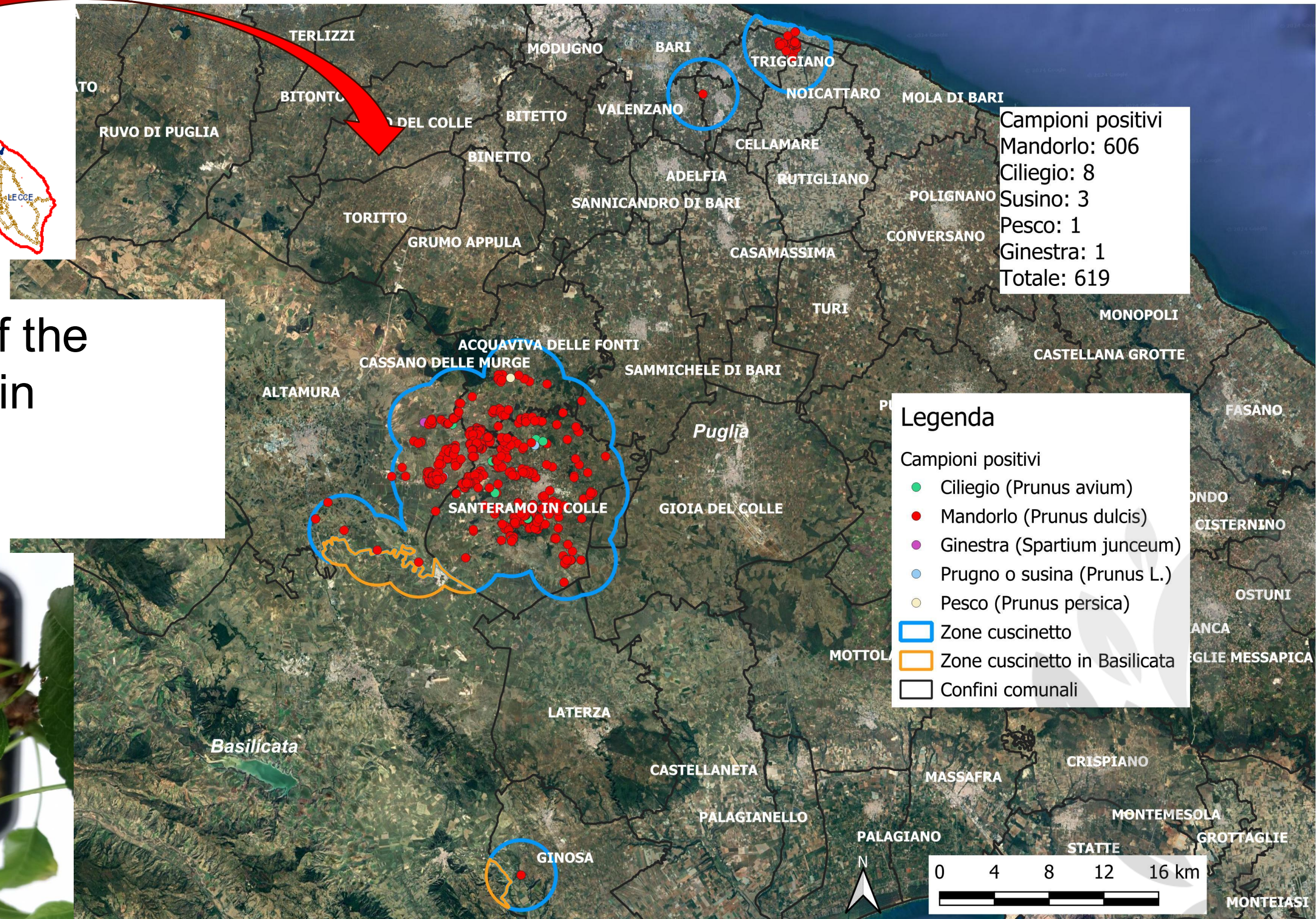
The **requirements** of the current EU Regulation for the control of *X. fastidiosa* are equally strict **regardless of the subspecies and the impact** of the strain present in a given area.

multiplex

First case of ST26 in EU



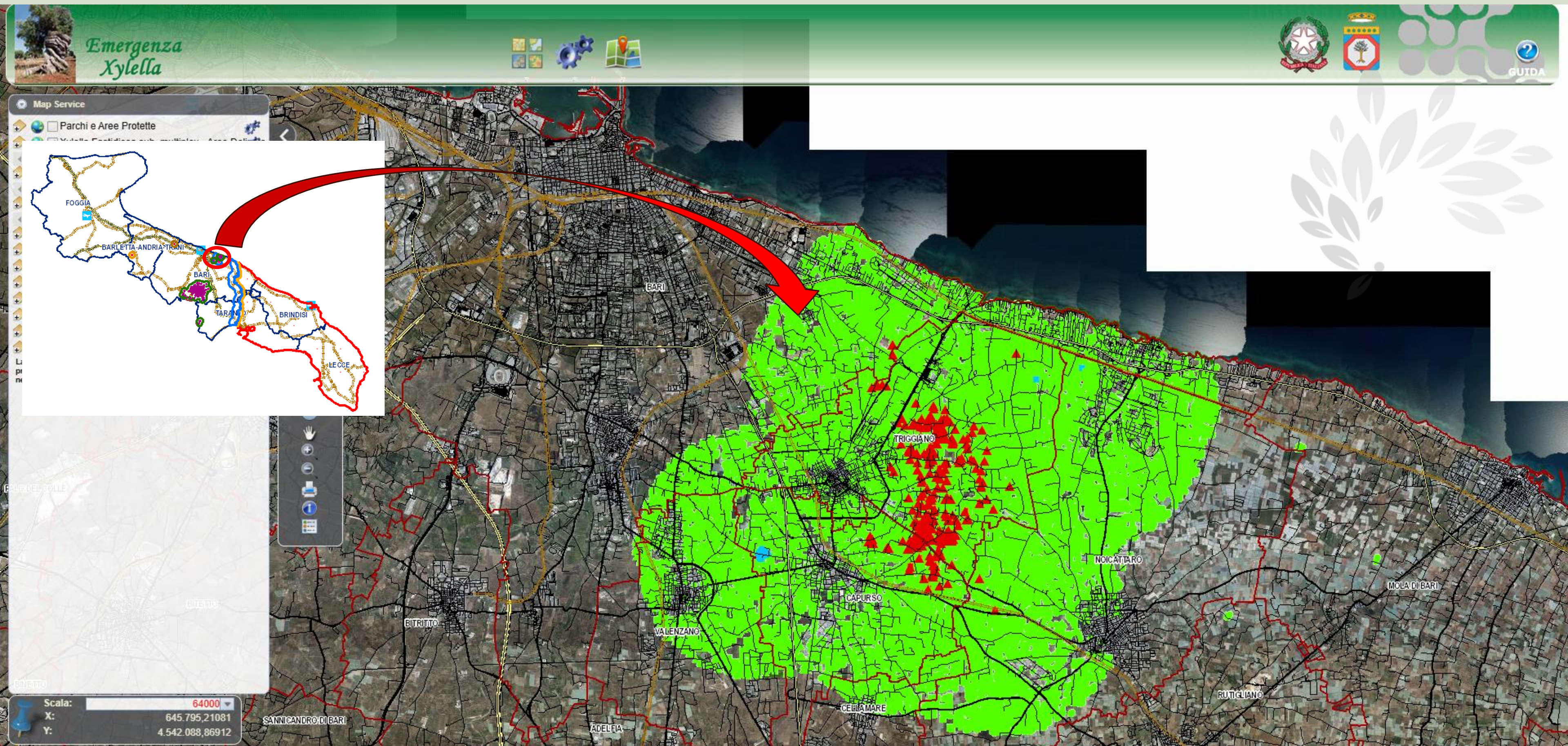
The delimitation of the boundaries is still in progress



18 MONTHS OF MONITORING OF MULTIPLEX

- the **impact** of bacterial infections appears to be **negligible**; symptoms, if any, are not distinguishable from those of drought, considered physiological in the NON irrigated almonds of this region
- the outbreak was discovered only as a consequence of the surveillance program, but so far in Puglia **no one has reported associated diseases**
- Beside the **almond**, the risk of infections of other «specified» plants seems to be very low
- The **boundaries** of the affected area are **not yet identified** (suspicion of old introduction and widespread presence)
- **Severe impact of the eradication program** (regulated as «mandatory»)

Xf fastidiosa ST1 (ca. 8 Km²)



20 months of monitoring «fastidiosa»

- Almond and Grapevine the major hosts
- Impact negligible on almond, to be defined in grape (major differences among cvs.)

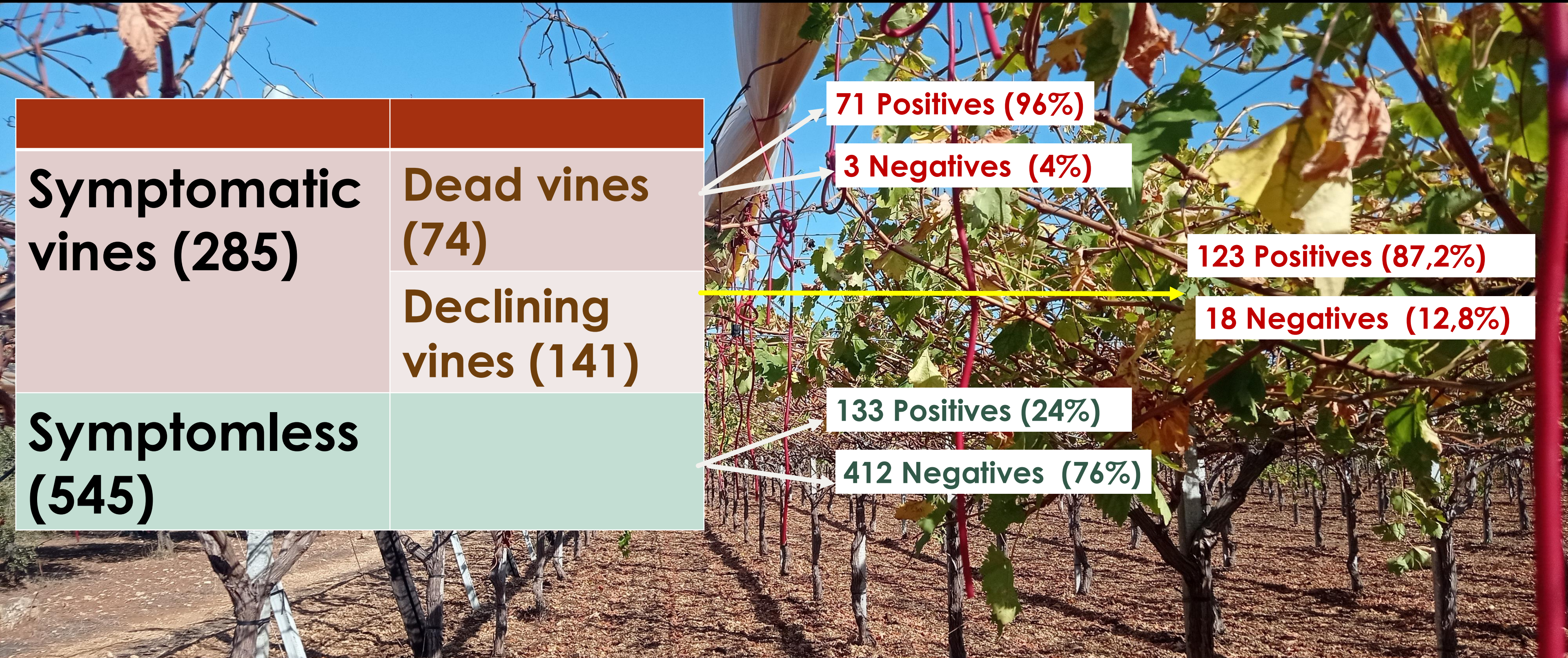
One example of susceptible Cv.: Superior 4 years old

Xf Positive ↓

Negative ↓



In dept analyses in one vineyard of cv. Superior (4 y old):



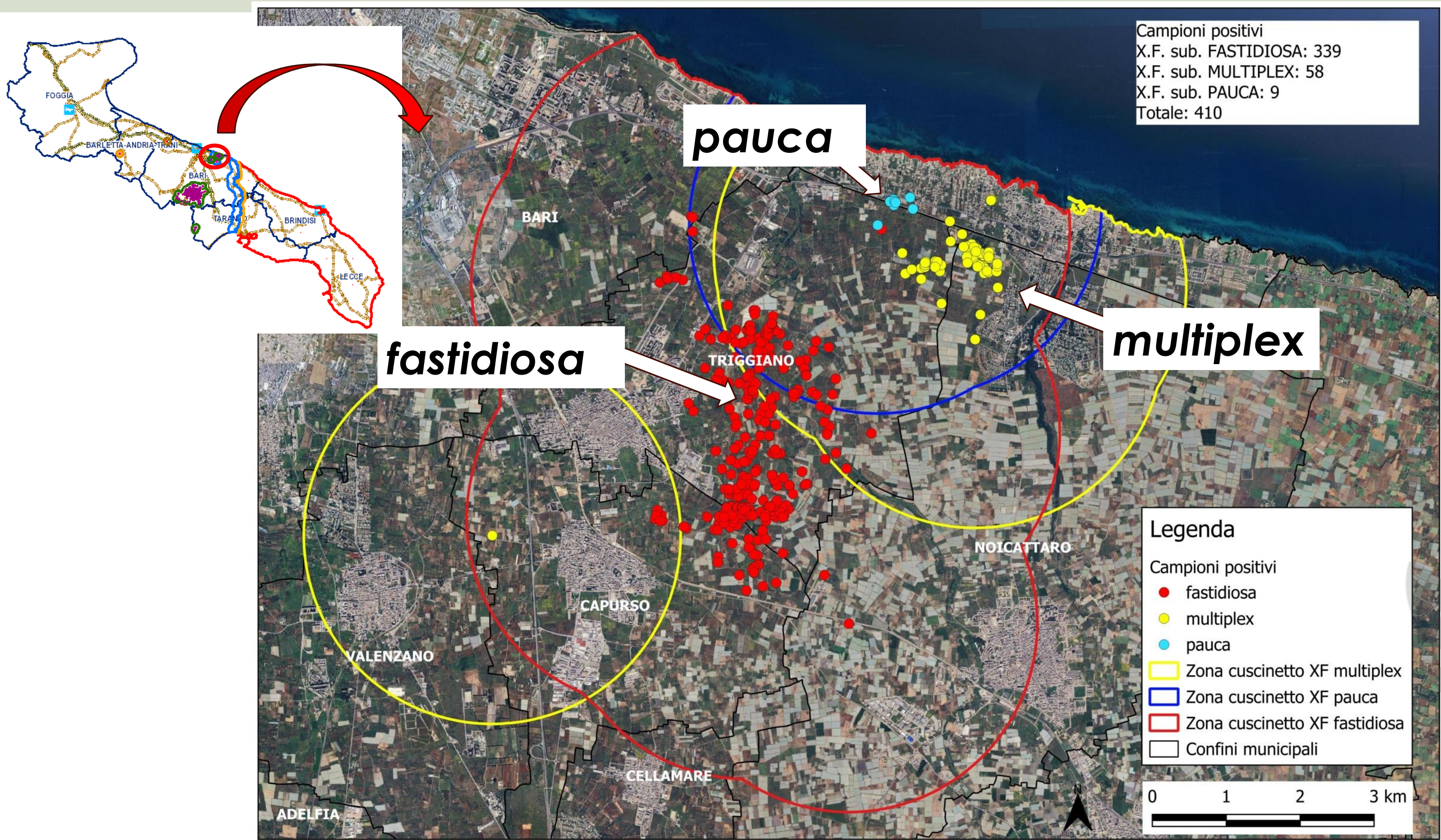
X. fastidiosa: The same species includes different pathogens

different diseases, host plants and impact

- *Pauca*
- *Multiplex*
- *Fastidiosa*




A complex picture (few Km from Bari)



NOT ALL BAD NEWS ARE NECESSARILY BAD (?)

Evidence suggests that **competition between strains may modulate disease prevalence** by influencing strain dominance in both hosts and vectors

i.e.: In the southern part of the island of Mallorca, where **the presence of multiplex (ST81) is higher**, the **incidence of Pierce's disease is lower** than in the rest of the island.

A decorative graphic of stylized leaves in a light gray color, located in the bottom right corner of the slide.

CONCLUSIONS: major changes since 2013

The complexity of the situation suggests the adoption of targeted and case-by-case adapted management measures, rather than a one-size-fits-all approach:

Is it time to consider revising the EU regulation?



THANK YOU!

donato.boscia@cnr.it



RIGENESI

