

Xylella fastidiosa in France



Plant Health Laboratory



Characterisation of *Xylella fastidiosa* strains of coffee

Alert given by positive **ELISA tests** on symptomatic **coffee** plants (2012) in confinement facilities then eradicated



Source : Anses LSV

Chlorotic leaves

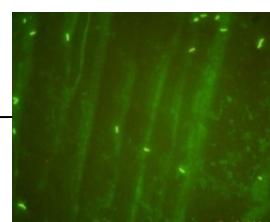
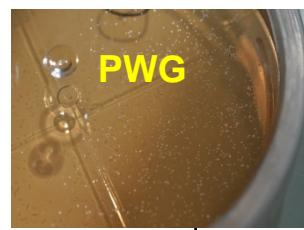


Source : Anses LSV

Marginal scorches

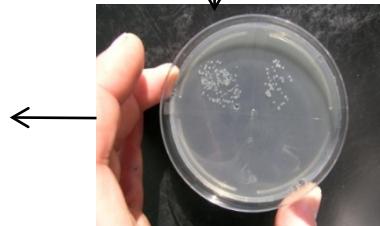
Processing symptomatic samples in
PBS

Isolation



Immunofluorescence

PCR (Minsavage et al. 1994)
Pooler & Hartung 1995 /
Firrao & Bazzi 1994)
Real Time PCR (Harper et al.
2010)

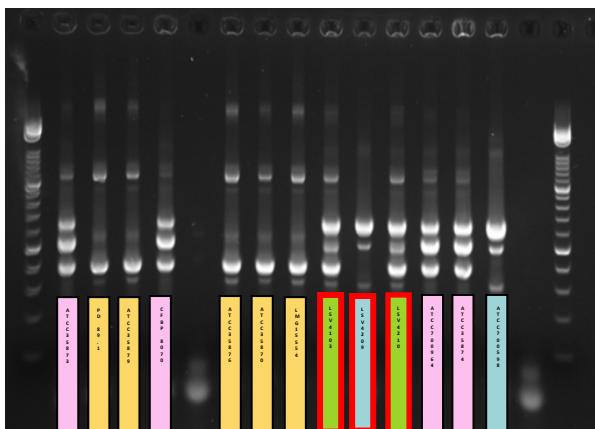


Typical
cells

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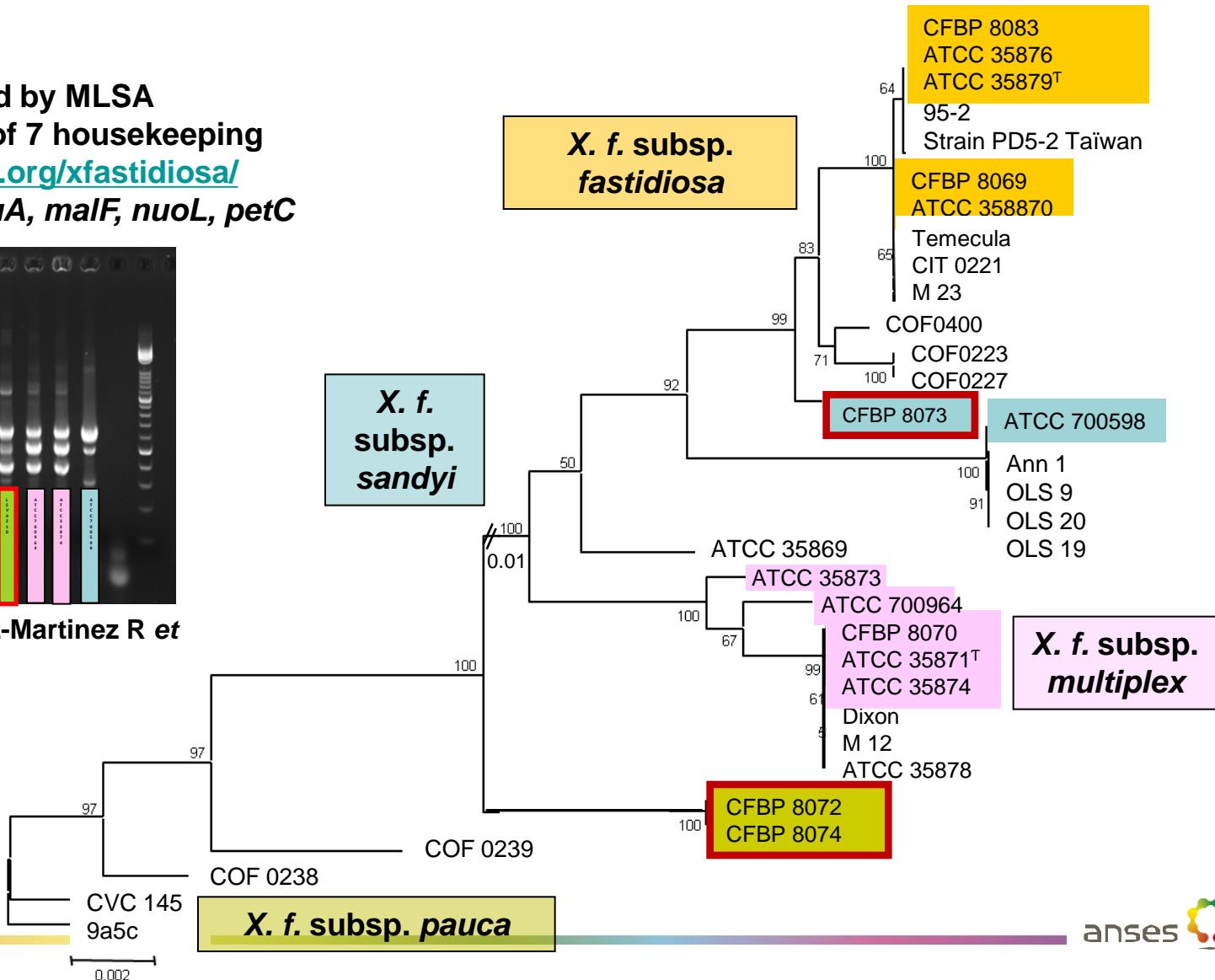
Phylogenetic concatenated tree (2012)

- Strains characterized by MLSA
- Partial sequences of 7 housekeeping genes: <http://pubmlst.org/xfastidiosa/>
- cysG, gltT, holC, leuA, malF, nuoL, petC*



PCR multiplex Hernandez-Martinez R et al., 2007

Strains isolated from coffee plants
2012



The first alert in Corsica: july 2015



The first alert in Corsica: july 2015



Before *Xylella*



After *Xylella fastidiosa*

Immediate emergency plan



In the infected zone : vector treatment, sampling of specified plants, destruction (priority to *Polygala* and any plant with symptom), no plantation of host plants

In all delimitated zone : no circulation of specified plants : in particular, no specified plant for planting goes out the delimited zone (including from nurseries), surveillance



Survey

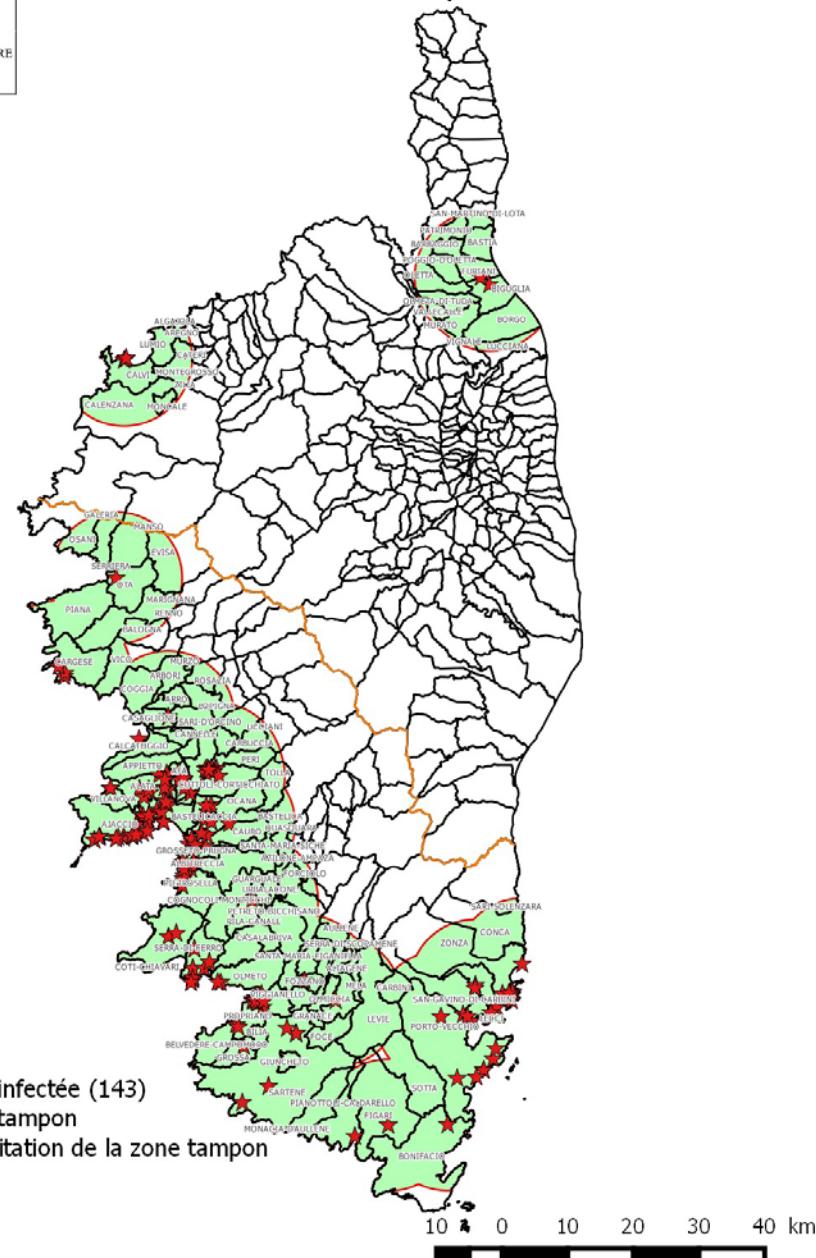
Samples analysed by rt-PCR
(Harper et al, 2010/2013)

January to
15th October 2015

	Samples	% positives
Total France	~ 3700	
Corsica samples	~2500	
Polygala myrtifolia	~1000	29%
Olea europea	~490	0
Oleander	~160	0
Coffee	~130	11%



Carte des zones tampons de 10 km autour des zones infectées par Xylella fastidiosa.

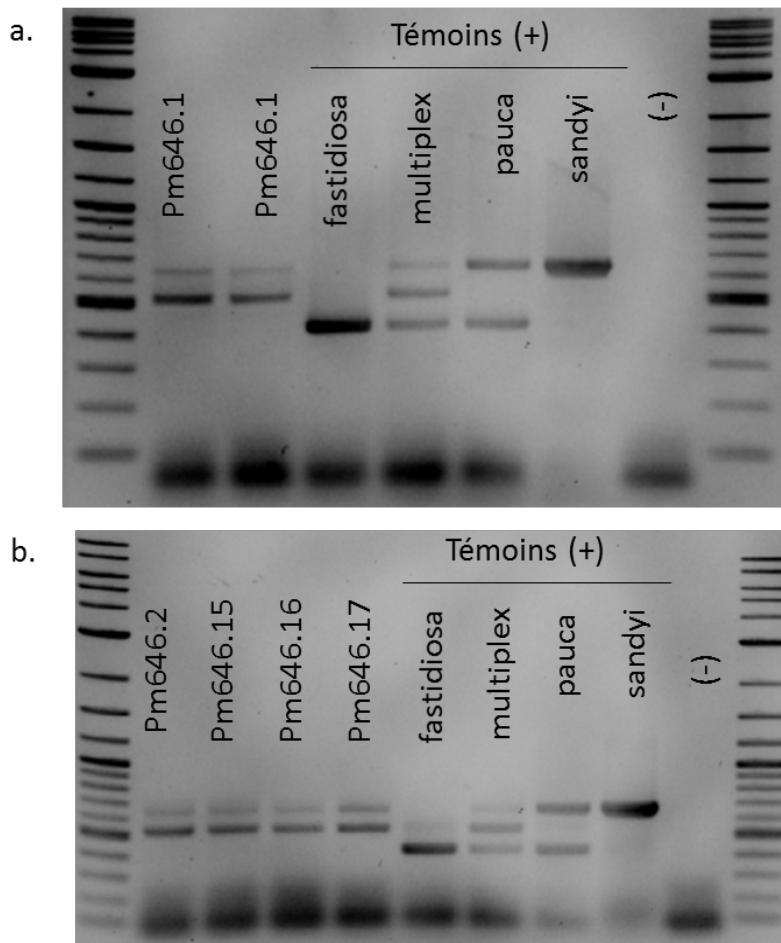


Légende

- ★ Zone infectée (143)
- Zone tampon
- Délimitation de la zone tampon



Identification of the X.f subspecies on corsica samples of *Polygala myrtifolia*



PCR multiplex Hernandez-Martinez et al. (2006):
subsp. *multiplex* (2 types founded)

- genotype **ALSII (Griffin-1)** (Chen et al., 1995 - 2013).
- Oak (*Quercus rubra* (OLS), USA (non known on *Vitis*)
- genotype **ALSI (Dixon (ST6), M12 (ST7))**
- Almond (*Prunus dulcis*), *Olea europea*, USA

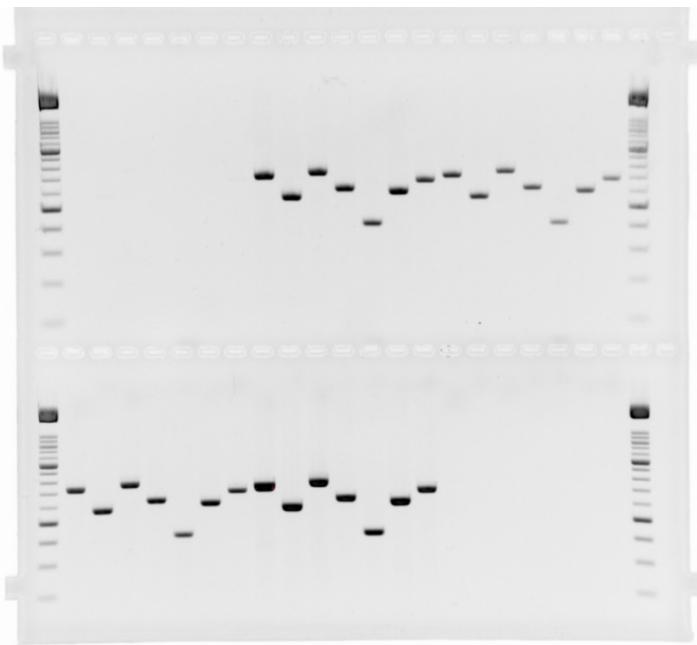
Figure 1. Affiliation par PCR à une sous-espèce de *X. fastidiosa*

Similarity sup 99,5%

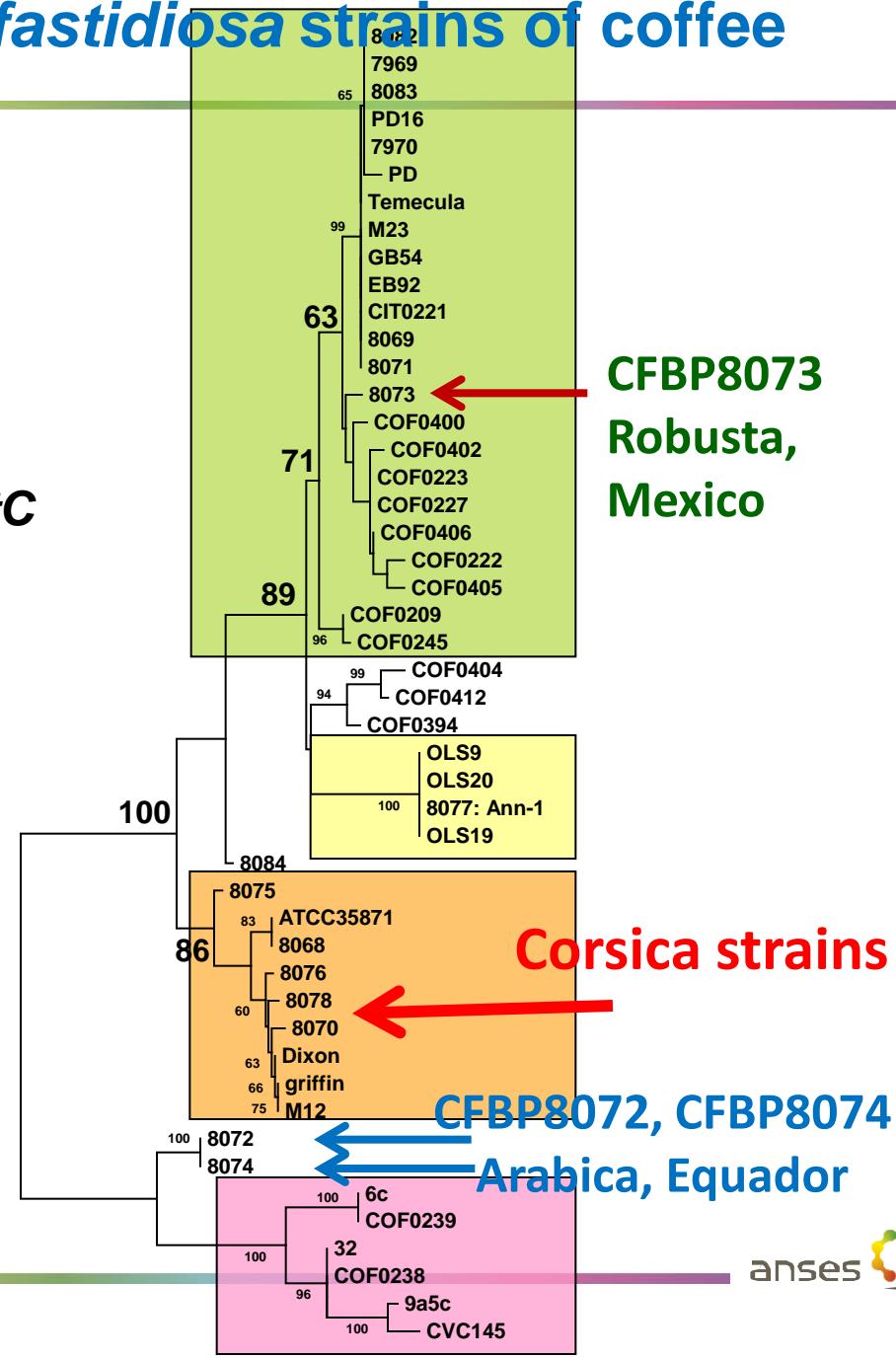
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- Characterization by MLSA (MultiLocus Sequence Analysis)
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fastidiosa sandyi multiplex pauca



Progressing host range in Corsica

Polygala myrtifolia

Pelargonium graveolens

Cytisus racemosus

Spartium junceum

Hebe sp.

Lavandula dentata hybride

Cistus creticus

Cistus monspeliensis

Genista ephedroides

Quercus suber

Myrtus communis

Rosmarinus officinalis

Acer pseudoplatanus



Research projects

- Evaluation and validation of detection method (NGS) of Xf on insects in collaboration with INRA
- Evaluation of ddPCR for complex matrices
- Validation of PCR for identification of subsp
- Pathogenicity test on panel of hosts in collaboration with INRA
- EU project H2020 Ponte (2015-2018)



Philaenus spumarius



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