



CONSIGLIO NAZIONALE DELLE RICERCHE

Istituto per la Protezione
Sostenibile
delle Piante, UOS Bari



UNIVERSITÀ
DEGLI STUDI DI BARI
ALDO MORO

Dipartimento di Scienze
del Suolo,
della Pianta e degli
Alimenti



Centro di Ricerca
Sperimentazione e Formazione
in Agricoltura "Basile Caramia"

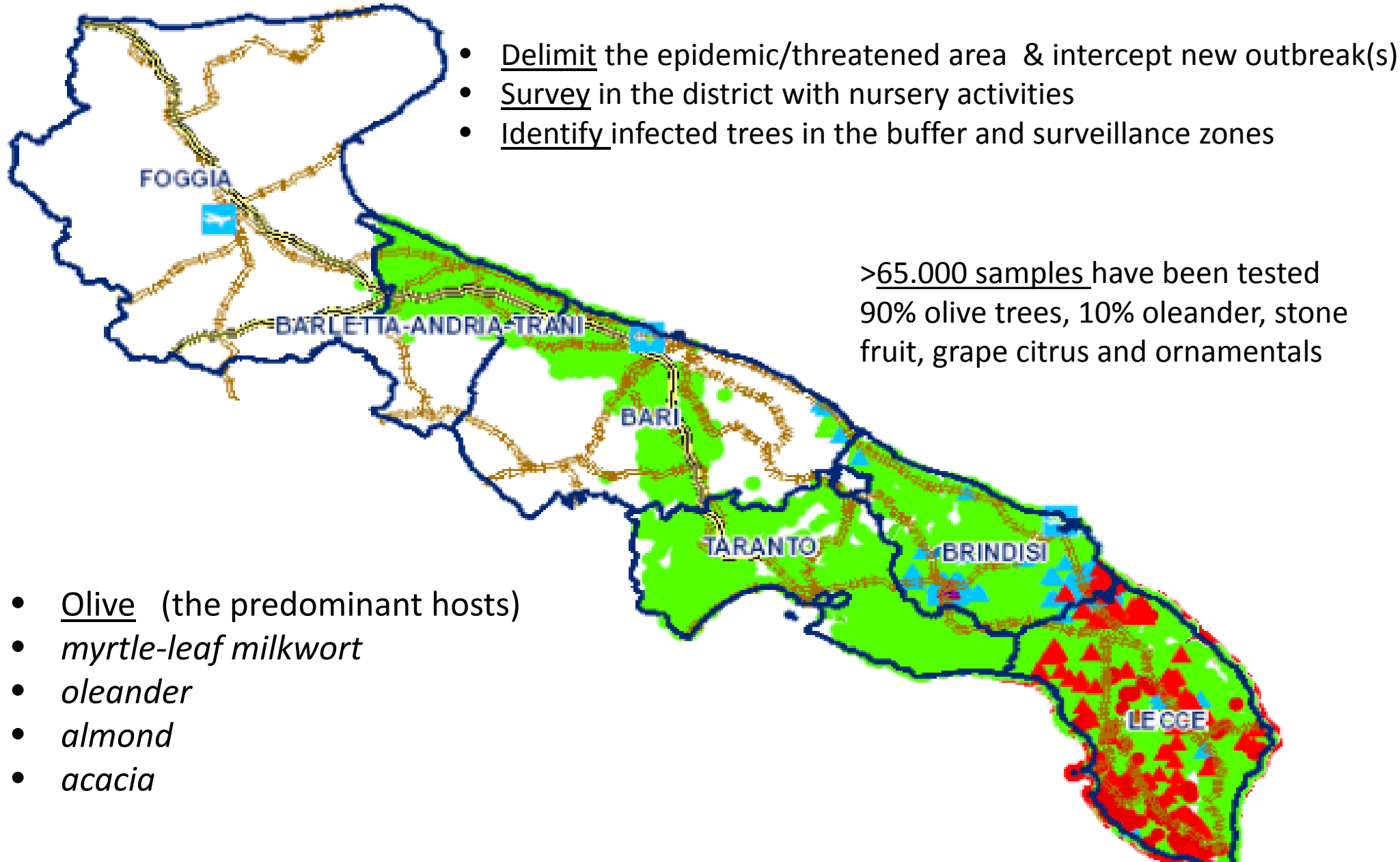


CIHEAM
IAM BARI

Diagnostic tools and protocols currently adopted in Apulia for large scale field monitoring of *Xylella fastidiosa*

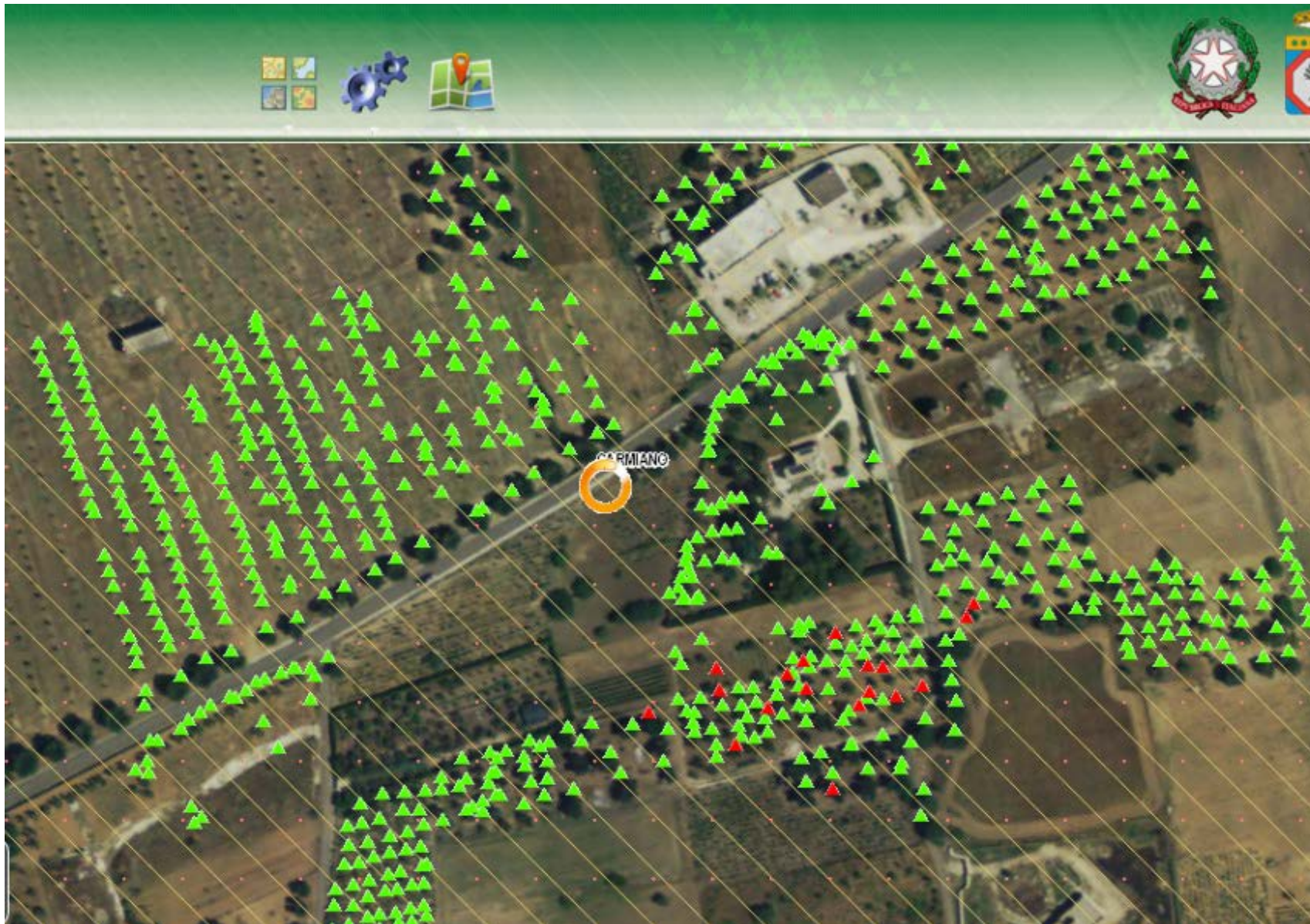
Maria Saponari, Giuliana Loconsole, Oriana Potere, Donato Boscia

Regional Monitoring Program: scope and target host plants



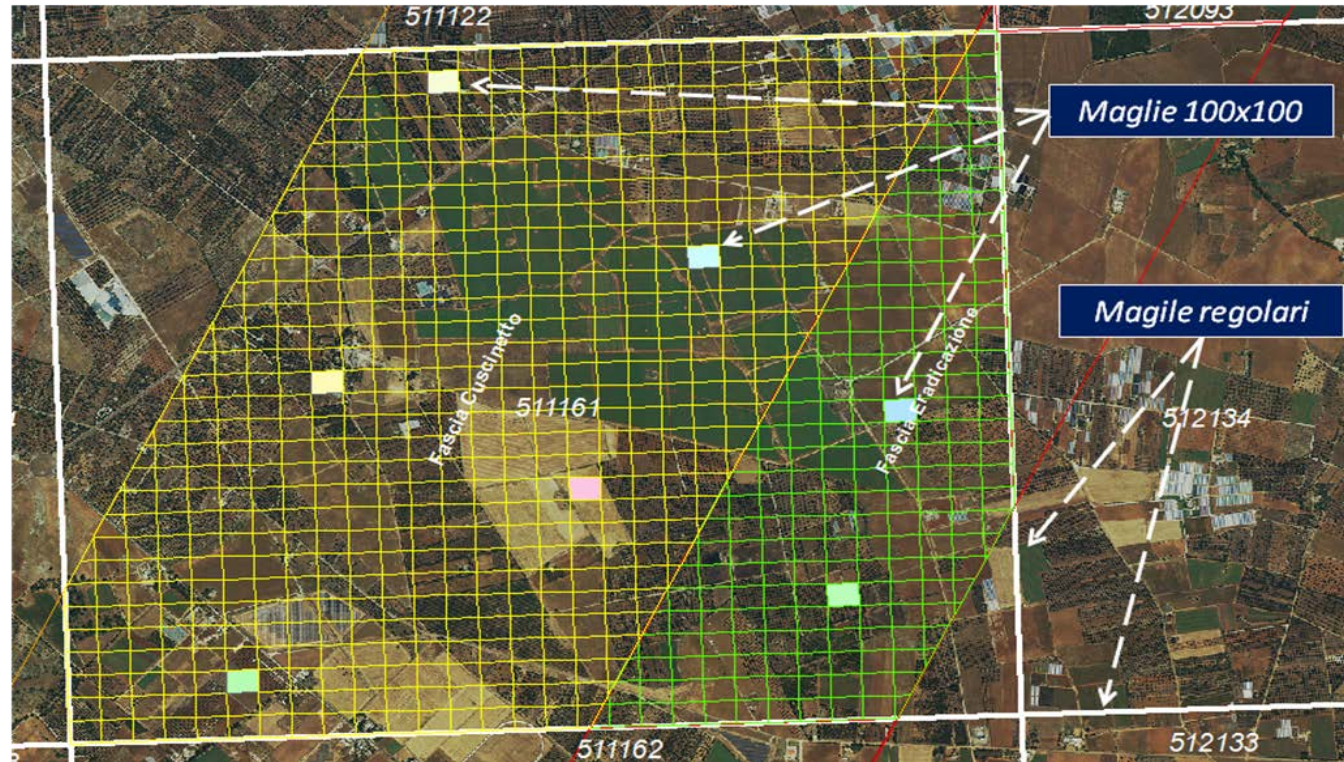
Close-up of the map

Survey in the surveillance area (single tree/plant of the specified host)



ELEMENTS characterizing an effective monitoring program

- Knowledge of the host range
- Knowledge of the symptoms associated to the infections
- Sites to be prioritized for monitoring
- Criteria for sampling
- Seasonal fluctuation of the bacterial population
- Laboratory tests for processing large scale number of samples



Sampling: the experience gathered upon the establishment of the bacterium under the Mediterranean conditions (climate, hosts, isolate introduced, etc.)

- Major susceptible crops and host species affected show symptoms, while few susceptible hostm plants appear to be symptomless

VISUAL INSPECTIONS IS A CRITICAL ASPECT

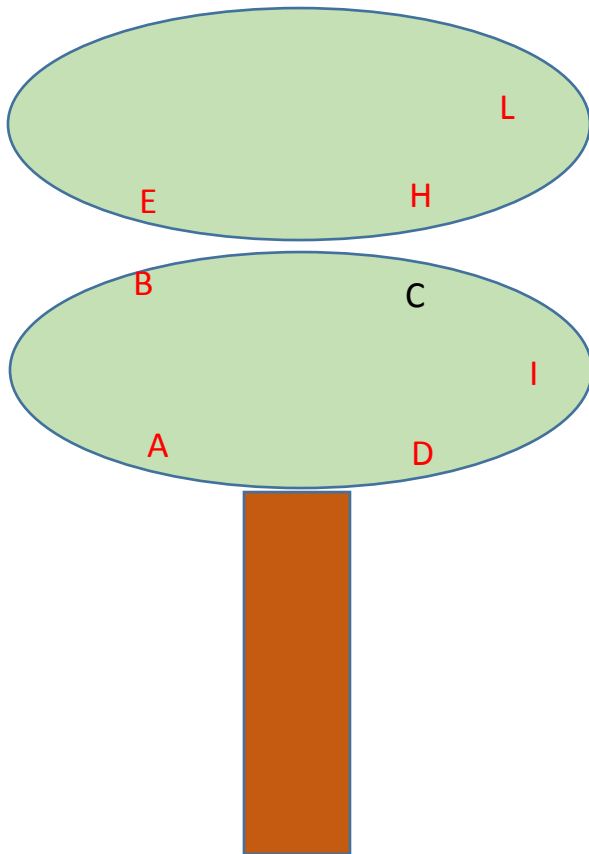
- Host species consisting in deciduos trees: the bacterium is undetectable in the leaves (qPCR/ELISA) in spring and early summer; in the wood it is detectable thought the entire year.
- Host species consisting in evergreen trees/shrubs: the bacterium is detectable in the leaves (qPCR/ELISA) through the entire year (referred to the specific conditions in the contaminated area); isolations from olive failed only in late summer; isolation from other susceptible hosts failed in late summer and in the winter season



PERIOD FOR SAMPLING – RELATED TO THE DIAGNOSTIC TESTS

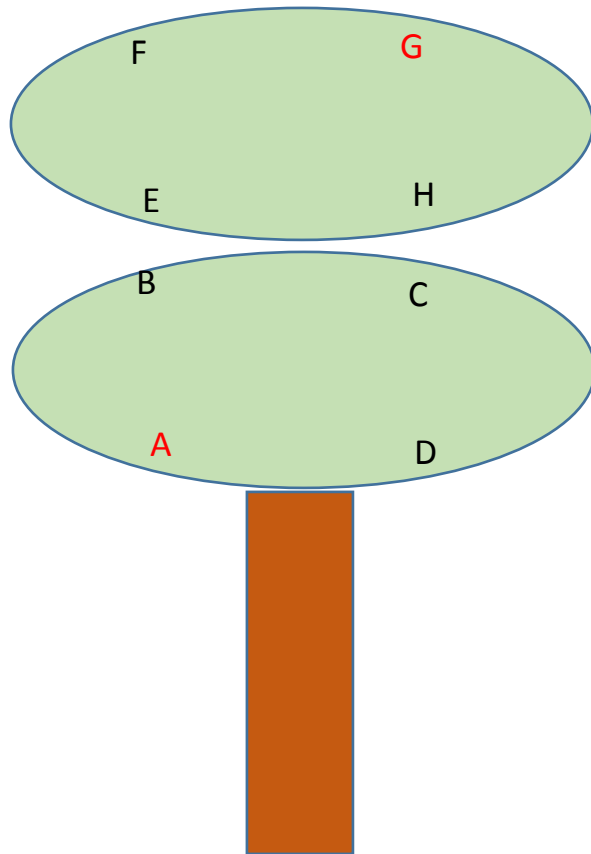
Sampling from symptomatic trees

7/8 subsamples tested positive qPCR/ELISA



Sampling from trees showing mild symptoms

2/8 sub-samples tested positive qPCR/ELISA



Correlation symptoms – Xf detection







Young portions



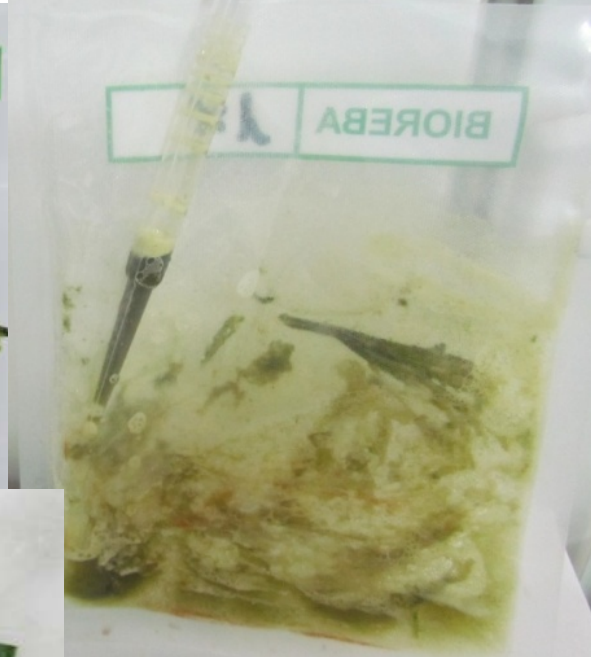
Mature cuttings



Diagnostic approaches tested for the CoDiRO strain

- *ELISA*
- *DTBIA*
- *PCR*
- *qPCR*
- *LAMP*

SAMPLE PREPARATION for DNA purification and ELISA/LAMP sap preparation



Advantage: process
0,5-1gr of tissue

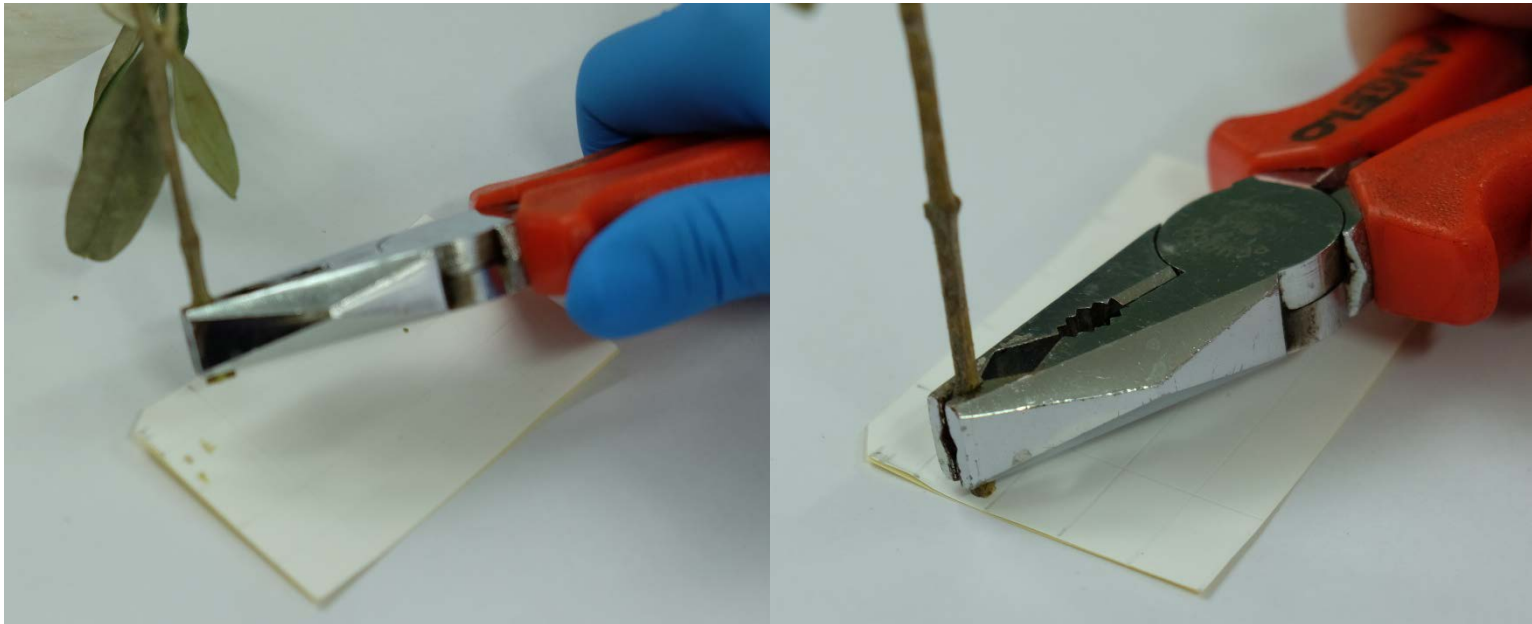
*Representativeness
of the sample*

Update on the implementations developed by the Apulian research groups:

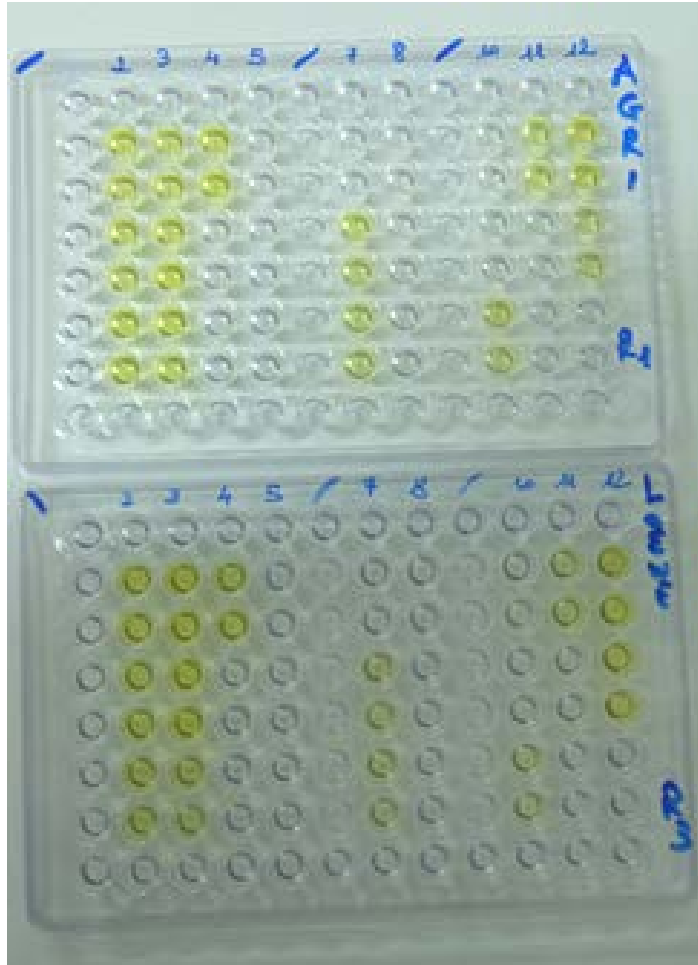
Serological approaches

- Antisera specific to the CoDiRO

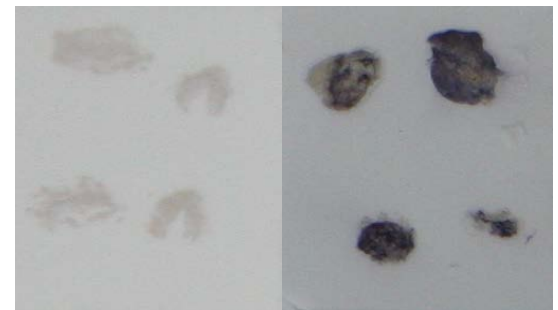
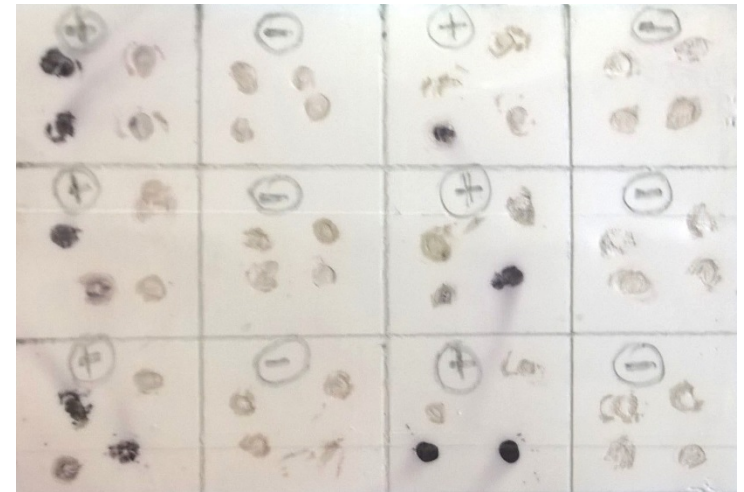
- DTBIA



ELISA (LOEWE & AGRITEST)



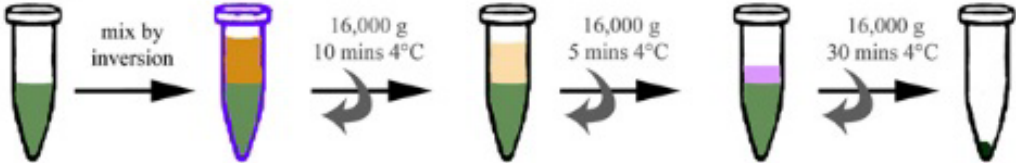
DTBIA



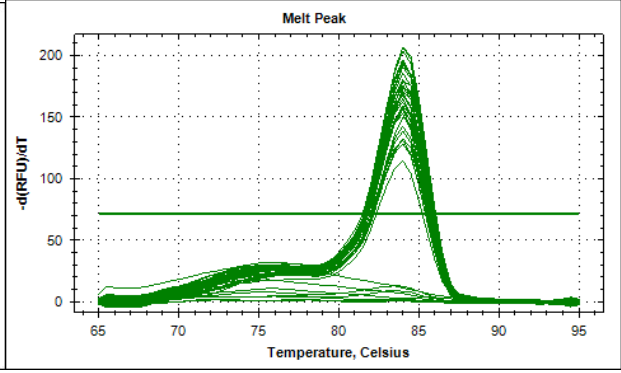
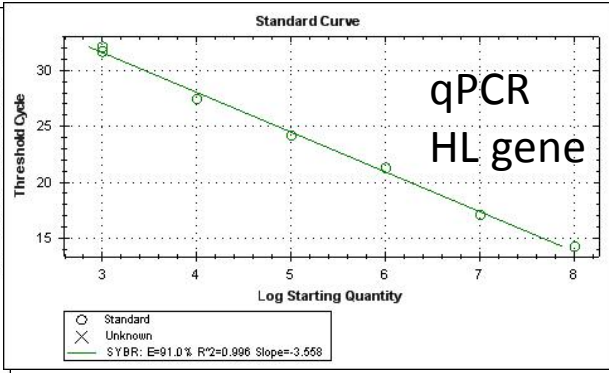
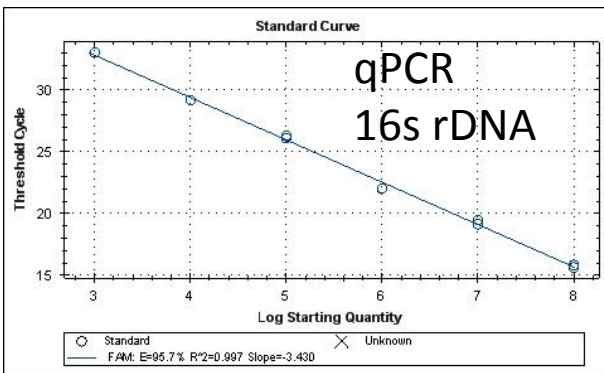
LAMP ASSAYS: no extraction



qPCR assays



CTAB extraction → Isopropanol precipitation → Purified DNA



Analytical sensitivity and interlaboratory validations

- ILV – every six months the laboratories involved

Panel of blind samples: - field infected samples of olive and oleander
 - artificially spiked samples (quercus, citrus, almond and grape)

TESTS	OLIVE	OLEANDER	GRAPES	ALMOND	CITRUS	OAK
ELISA	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%
PCR (RST31/33)	PA= 80% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%
qPCR	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%	PA= 100% NA=100%
DTBIA	PA= 100% NA=100%					
LAMP	PA= 100% NA=100%	PA= 100% NA=100%				

ELISA & qPCR on different matrices

Matrices (3 spiked samples)	Harper et al., 2010 Cq values	ELISA OD ₄₀₅ values
OLIVE	14.57 – 15.26 (Δ Cq 0.69)	2.829 – 3.021
ALMOND	15.27 – 15.56 (Δ Cq 0.29)	2.559 - 2.963
OAK	14.89 – 15.28 (Δ Cq 0.49)	2.734 – 3.062
GRAPE	14.55 – 14.94 (Δ Cq 0.39)	2.578 – 2.923
CITRUS	14.87 – 14.91 (Δ Cq 0.04)	2.938 – 3.060

Field samples (reaction background can occur)

Notes

qPCR: In 2 out 5 laboratories, negative samples produced Cq 35-38

DTBIA: Produced reliable results but their interpretation could be a limiting factor

LAMP: False negative if using only small piece/sample

Alternatively crude sap of purified total nucleic acid can be used to overcome such erratic reaction

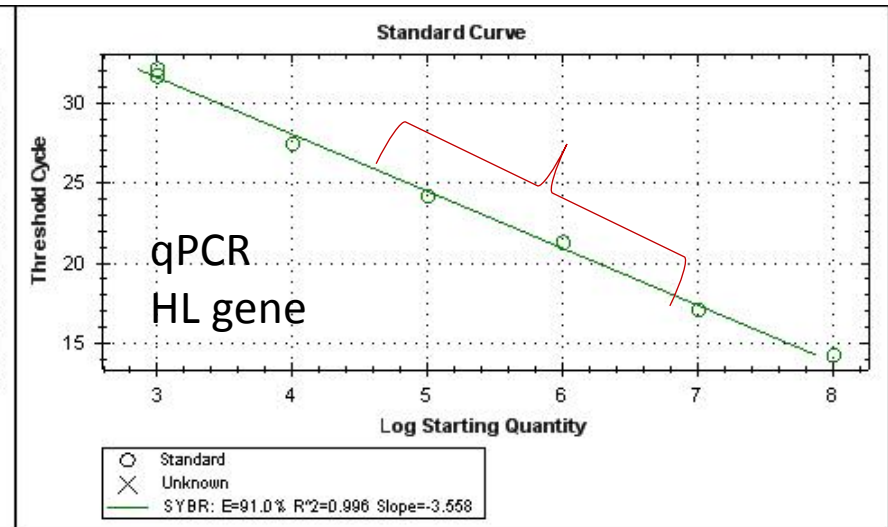
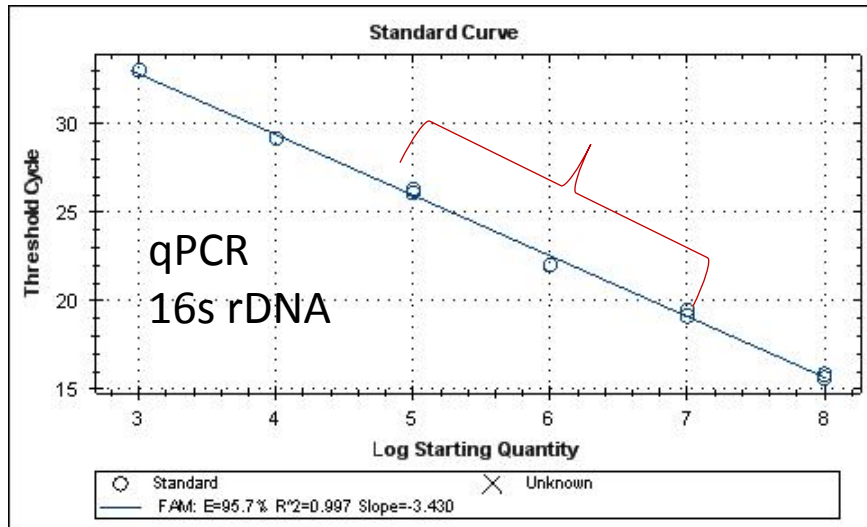
[Conc] & range for field samples

Analytical sensitivity

Technique	CONCENTRATION – CFU/ml OF PLANT EXTRACT						
	10 ⁷	10 ⁶	10 ⁵	10 ⁴	10 ³	10 ²	10
ELISA Average (AgriTest/Loewe)	Positive 3,345	Positive 3,360	Positive 2,801	Positive 0,403	Negative 0,186	Negative 0,025	Negative 0,023
PCR	Positive	Positive	Positive	Positive	Negative	Negative	Negative
qPCR	Positive	Positive	Positive	Positive	Positive	Positive	Negative
LAMP Enbitech	Positive	Positive	Positive	Positive	Positive	Positive	Negative

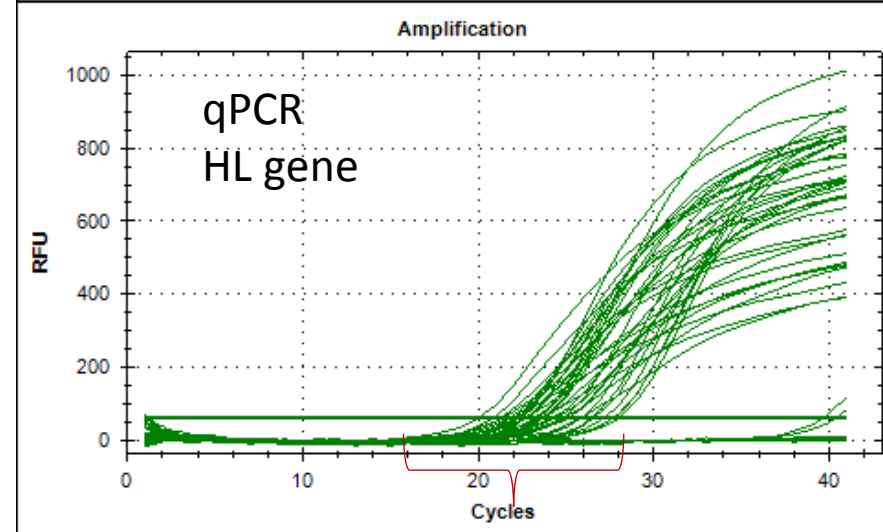
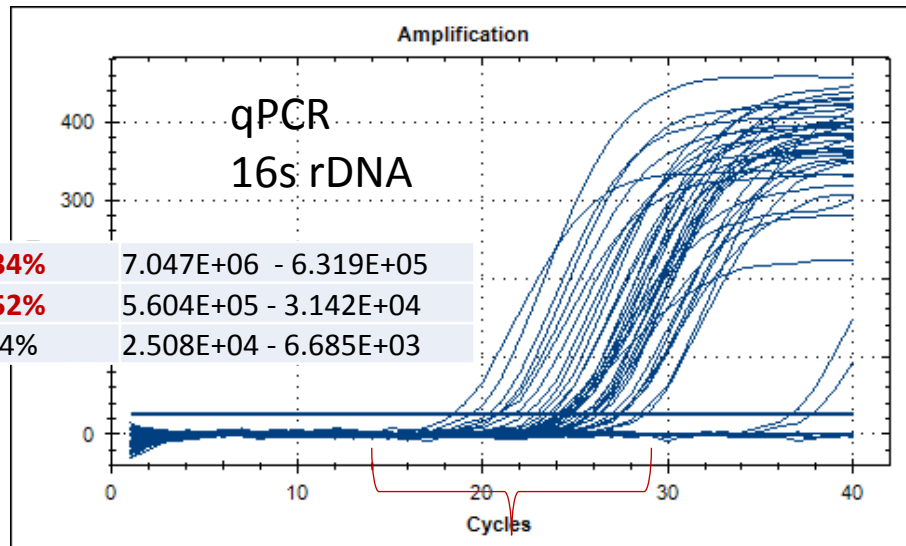
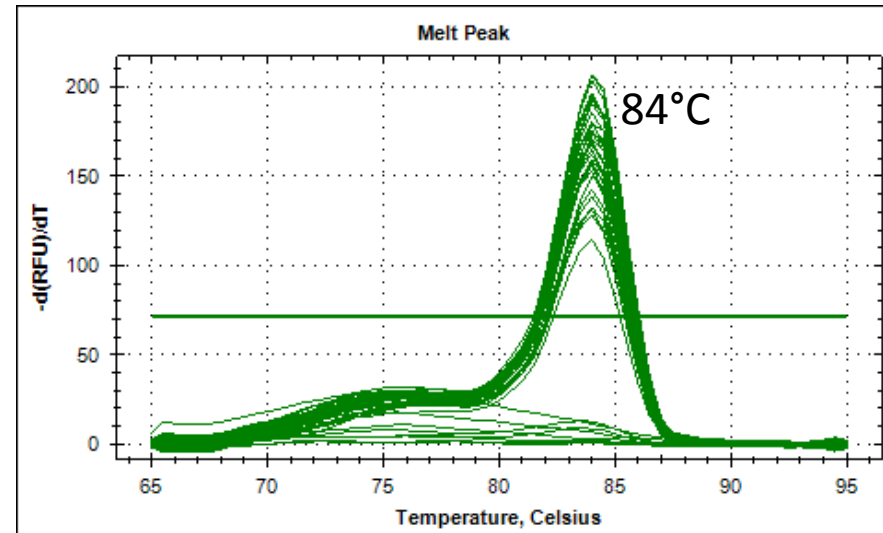
Suspected tree

Positive



Validation of serological and molecular tests in the field in new the outbreaks (Brindisi province)

ELISA	Average of the OD ₄₀₅
POSITIVE and SUSPECTED SAMPLES (ca. 75%)	57% (>1,000) 43% (0,400 - <1,000) 20% (0,100- < 0,400)
NEGATIVE SAMPLES (ca. 25%)	---



Test performed on October 15th

Validation of serological and molecular tests in the field in new the outbreaks

GIVEN A PANEL OF ca. 900 FIELD SAMPLES CATEGORIZED BY ELISA TEST AS

Positive/suspected (700 SAMPLES)		Negative (200 samples)	
qPCR positive	97%	qPCR negative	98%
ELISA false positive	3%	ELISA false negative	2% (qPCR positive)
PCR RST31/33	94%	nd	
DTBIA	71%	nd	
LAMP*	56%	nd	

* Protocol Enbitech

Xf-free area

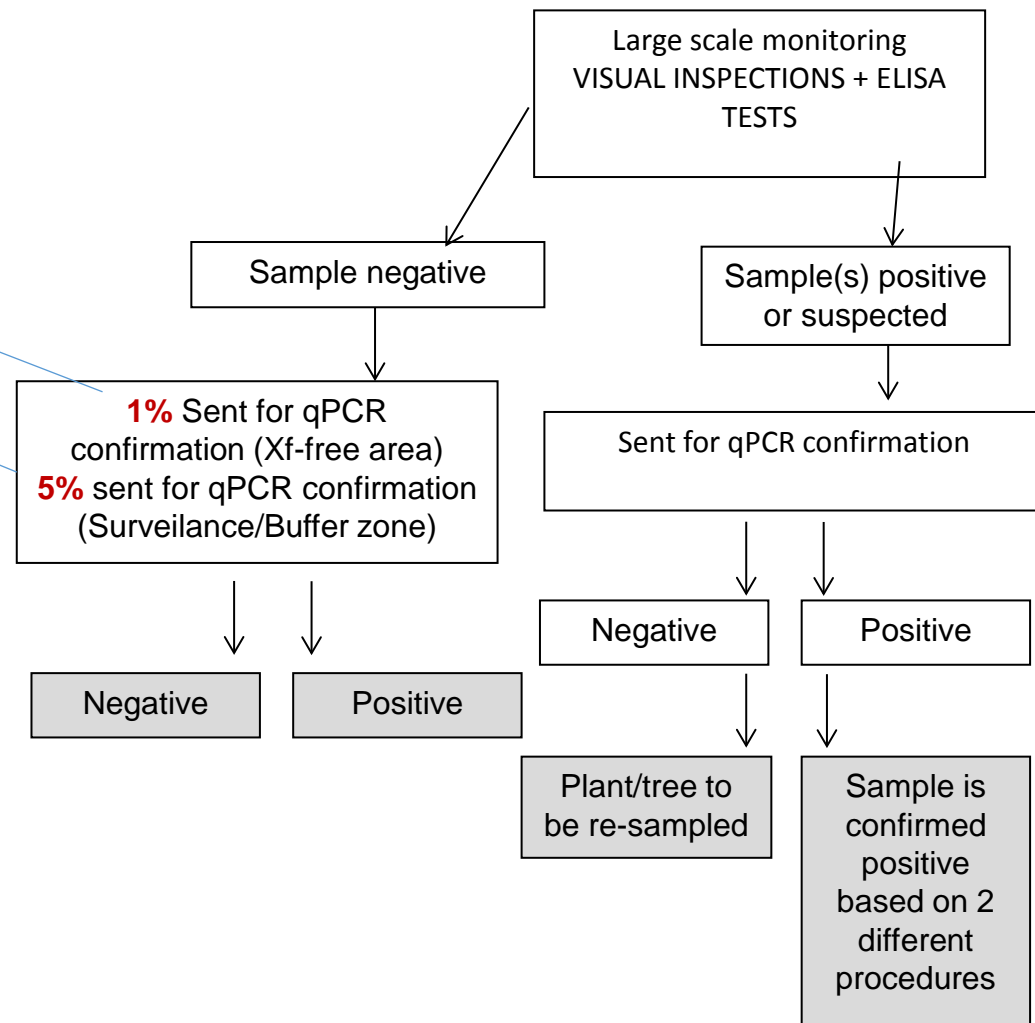
Surveillance
/buffer zone

Contaminated area

Legenda

- Zona infetta
- Zona cuscinetto
- Zona di sorveglianza
- Confini provinciali
- Confini comunali

www.emergenzaxifa.it



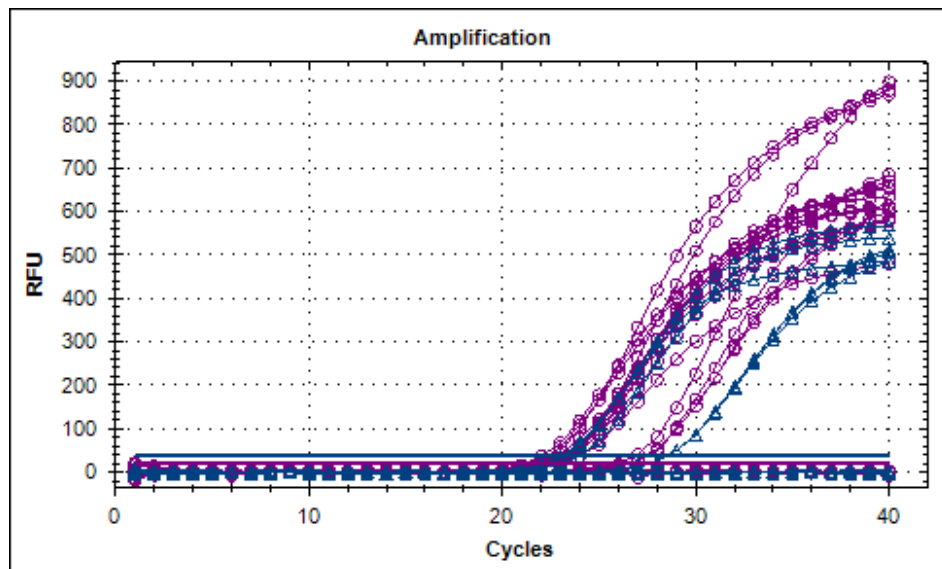
For samples that consistently give Cq values >32, it is suggested to proceed with an additional sampling

Ongoing researches:

1) Multiplex qPCR (target Xf gene [16s rDNA gene] and DNA plant internal control – COX)

Li et al., 2006

10-fold serial dilutions (Xf)	Singleplex		Multiplex	
	Cox- CY5	Xf-FAM	Cox-CY5	Xf-FAM
1.000E+08	18.16	18.13	18.18	17.44
1.000E+07	18.95	21.83	18.63	21.50
1.000E+06	18.74	25.54	18.21	24.64
1.000E+05	20.01	28.00	19.03	28.85
1.000E+04	19.11	32.54	19.60	32.34
1.000E+03	18.02	35.36	18.03	35.52
1.000E+02	18.19	37.40	17.99	37.15



- Almond, Cherry
- Citrus
- Grape
- Oleander
- Acacia saligna
- Myrtus and Rosmarinus
- Euphorbia terracina
- Rhamnus alaternus
- Eremofila maculata
- Grevillea juperina
- Lavandula stoechas
- Westringia glabra
- Cistus creticus
- Asparagus acutifolius
- Carissa macrocarpa
- Laurus nobilis
- Myoporim spp.
- Dodonea viscosa purpurea

Cq 16.77 – 19.45

2) Automatization of the extraction procedure

DNeasy *mericon* Food Kit (Modified CTAB procedure)

- 0,5-1gr of tissue
- 1 spin column



+



Qiacube

DNeasy *mericon* Food Kit (Modified CTAB procedure)

- 0,5-1gr of tissue
- 1 spin column

Parameters	ng/μl	ng/μl
	<u>CTAB</u>	<u>Mericon</u>
[DNA]	100-800	18-30
A260/208	>1,8	>2
Cq* Xf	21.68 - 28.62	16.37 - 25.42
Cq COX	15.53 - 25.66	18.06 - 20.28

*working on the detection limit

Evaluation of the qPCR master mix

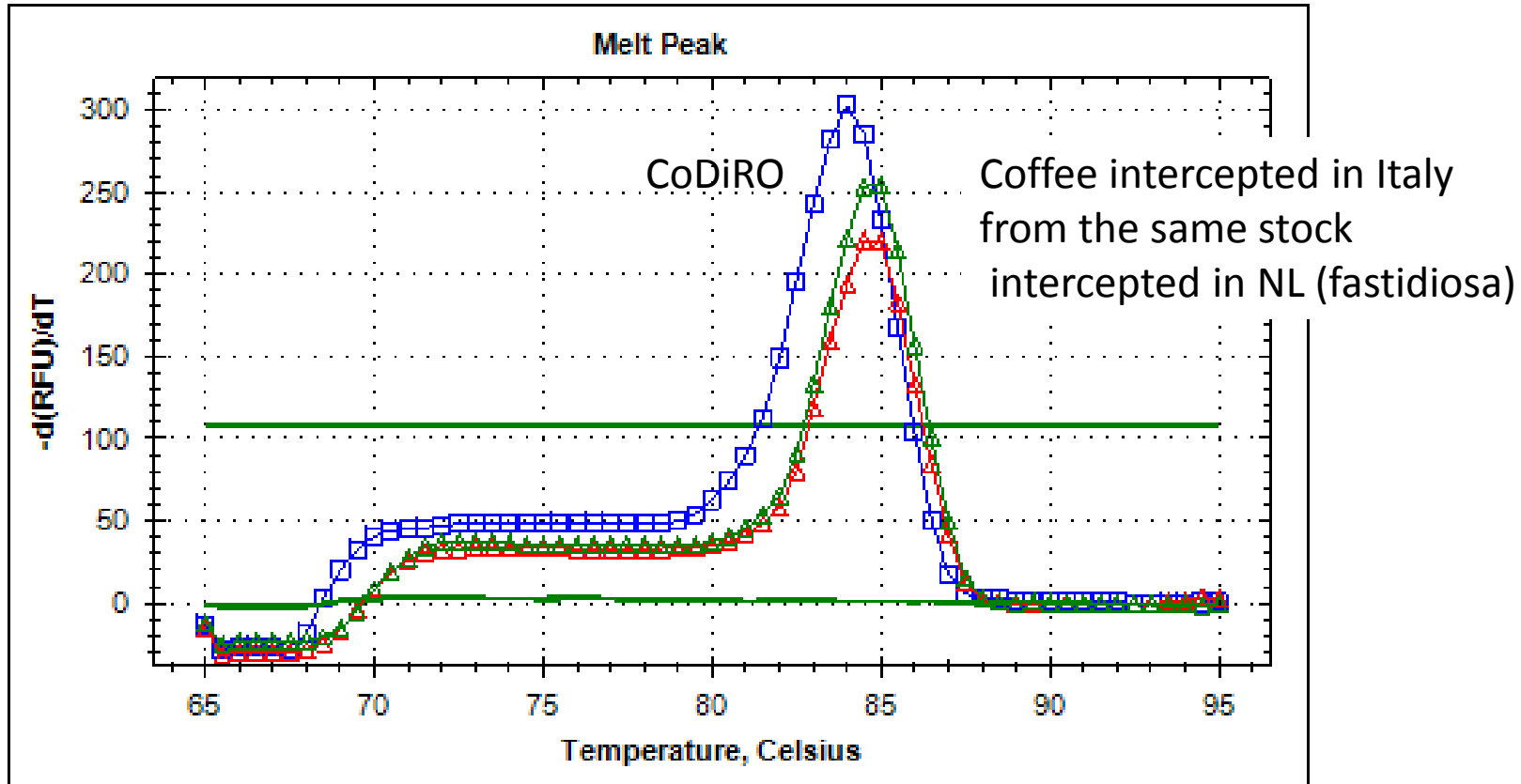
Master mix «A» and «B»

Master mix	Cherry		Oleander		Coffee		Almond		Olive	
	Undiluted	Diluted	Undiluted	Diluted	Undiluted	Diluted	Undiluted	Diluted	Undiluted	Diluted
«A»	(+)	(+)	(+)	(+)	(+)	(+)	(+)	(+)	(+)	(+)
«B»*	(+)	(+)	(+)	(+)	(-)	(+)	(-)	(+)	(+)	(+)

* Negative control Cq ≥ 36

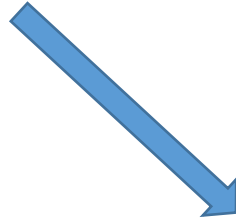
qPCR HL gene – Sybr green + melt curve analysis

Proved to be a useful tool for the preliminary isolate/strain discrimination



ISOLATION for the CODIRO: BCYE MEDIUM

Surface sterilization: 2% NaOCl – 70% ETOH



- 1) Isolation
- 2) Triple cloning
(no more than 3 passages)

Cuttings: olive, oleander, cherry,
almond, polygala, westringia



**Bacterial suspension in
PBS/Glycerol for storage**

Questions from the Plant Health Service:

- Sampling and tests to be used in the nurseries on propagating material
- Sampling scheme in the field (in the xf-free areas)

THANKS FOR YOUR ATTENTION

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