

#### On the importance of multidisciplinary studies on insect vectors to better understand vector-borne plant diseases

Astrid Cruaud, Marguerite Chartois, Pauline Farigoule, Martin Godefroid, Xavier Mesmin, Ileana Quiquerez, Sabrina Borgomano, François Casabianca, Guénaëlle Genson, Anne-Alicia Gonzalez, et al.

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3rd European Conference on Xylella fastidiosa and XF-ACTORS final meeting







## On the importance of multidisciplinary studies on insect vectors to better understand vector-borne plant diseases

Astrid Cruaud<sup>1</sup>, Marguerite Chartois<sup>1</sup>, Pauline Farigoule<sup>1,2</sup>, Martin Godefroid<sup>1</sup>, Xavier Mesmin<sup>1</sup>, Ileana Quiquerez<sup>3</sup>, Sabrina Borgomano<sup>3</sup>, François Casabianca<sup>4</sup>, Guénaëlle Genson<sup>1</sup>, Anne-Alicia Gonzalez<sup>1</sup>, Laetitia Hugot<sup>3</sup>, Maxime Lambert<sup>1</sup>, Éric Pierre<sup>1</sup>, Sylvain Santoni<sup>5</sup>, Jean-Claude Streito<sup>1</sup>, Jean-Pierre Rossi<sup>1</sup> and Jean-Yves Rasplus<sup>1</sup>

<sup>1</sup> CBGP, INRAE, Montferrier-sur-Lez, France; <sup>2</sup> AgroParisTech, Paris, France; <sup>3</sup> CBNC, OEC, Corte, France. <sup>4</sup> LRDE, INRAE, Corte, France; <sup>5</sup> AGAP, INRAE, Montpellier, France astrid.cruaud@inrae.fr

















#### WHY VECTORS?

- Because nobody will ever study malaria or dengue without studying mosquitoes..
- Studies on the biology /ecology /distribution of vectors are crucial to understand and control the spread of Xf
- Virtually nothing was known for EU vectors before XF-ACTORS
- Significant progresses have been made
- I will report on studies conducted by my research group in the framework of XF-ACTORS and other national (french) projects.





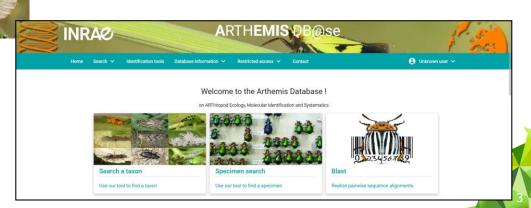
- Identification based on morphological characters can be difficult, sometimes impossible (eggs, nymphs) and always time consuming
- Massive molecular survey of communities of vectors (e.g. through metabarcoding)
   will be possible only with a reference database built by taxonomists





https://arthemisdb.supagro.inra.fr/

- Several specimens / species
- Different locations when possible
- Non destructive DNA extraction
- 2 step PCR + MiSeq sequencing (Cruaud et al. 2017)
- Bioinformatic workflow with quality controls (contaminants; NUMTs; heteroplasms; introgression)





#### A CURATED COI DATABASE FOR VECTORS

- All frequent EU species (#74) have been barcoded + 32 non-EU species
- Pictures of habitus and genitalia
- Identification of a query sequence by BLAST
- Reliable identification for 80% of the species



- P. spumarius and P. tesselatus same COI and morphologically close → synonyms?
- Possible species complexes within
   Neophilaenus and Aphrophora → clarification in progress

#### https://arthemisdb.supagro.inra.fr/





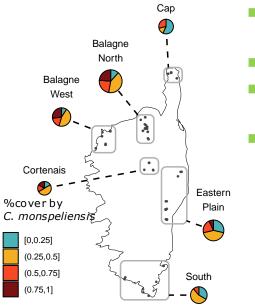
DOI:10.5281/zenodo.4680659



## No.

#### **PHILAENUS SPUMARIUS: KNOW YOUR ENEMY**

- Extensive research on Ps in different countries in the course of XF-ACTORS.
- Climate and vegetation drivers influencing its abundance still poorly known, esp. in semi-natural habitats.



- density of Ps monitored 3 times / year during 3 years in 64 plots throughout Corsica
- Visual counting of white foam blobs / sweep netting of adults
  - Description of the vegetation + daily temperature retrieved from weather databases

GLMM







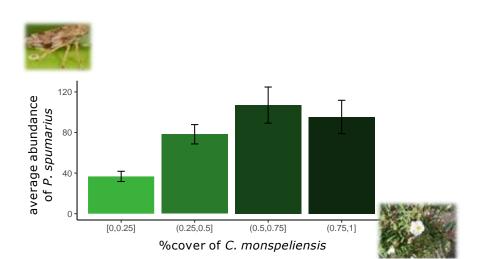


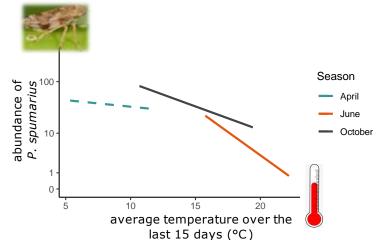




#### **PHILAENUS SPUMARIUS: THE CORSICAN EXCEPTION?**

- Abundance of Ps positively correlated with density of Cistus monspeliensis
- → contrast with mainland France and other countries in EU, more polyphagous
- → Xf expansion facilitated by disturbed habitats / firebreaks?
- Abundance of Ps negatively correlated with increasing temperatures
- → Role of Ps as vector is summer is reduced (other vectors?)



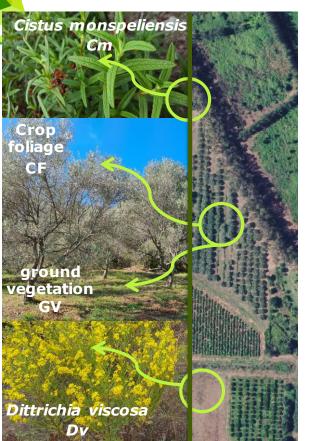




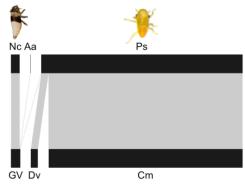


#### SPITTLEBUGS IN CROPS: NETWORK OF INTERACTIONS

Survey of spittlebugs in the vicinity of 16 organic clementine and olive groves (2 years)

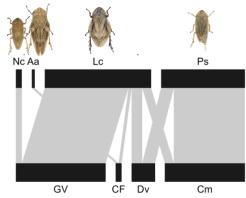


#### Spring 6647 spittles



- Strong host preferences
- No nymph found on crop foliage

#### Summer 119 adults

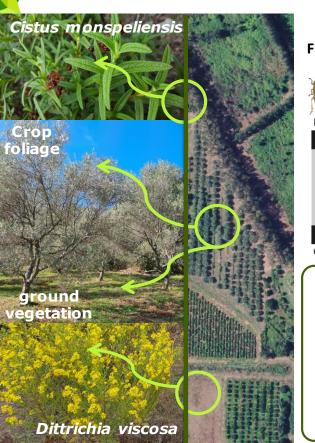


- Few spittlebugs
- Only A. alni and L. coleoptrata on crop foliage.
- No summer migration of *Ps* to crop foliage as observed in Italy

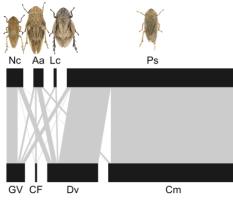


#### **PHILAENUS SPUMARIUS: THE CORSICAN EXCEPTION?**

Survey of spittlebugs in the vicinity of clementine and olive groves (2 years)



#### Fall 1595 adults

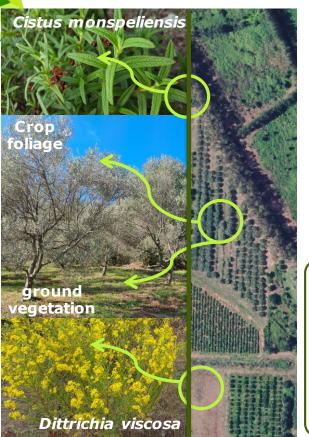


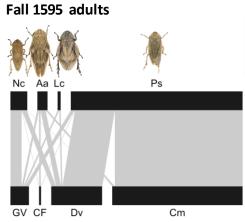
- complex network
- *Ps Cm* major component
- All species but Lc found on crop foliage.
- Possible role of other vectors need to be investigated



#### **PHILAENUS SPUMARIUS: THE CORSICAN EXCEPTION?**

Survey of spittlebugs in the vicinity of clementine and olive groves (2 years)



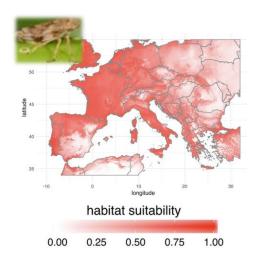


- complex network
- *Ps Cm* major component
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- Soil tillage against Ps probably less efficient than in Italy.
- Management of *C. monspeliensis* borders may decrease the threat posed by Ps to crops.

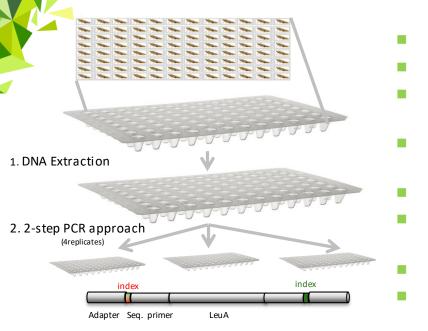


- Monitoring of Xf usually carried out by analyzing symptomatic plants.
- Plants are frequently asymptomatic
- Generalize the use of insect vectors as sentinels to complement plant survey.
- Ps is the perfect sentinel (largely distributed in EU acc. observations / SDMs)
- First test with nested PCR and Sanger sequencing; encouraging results
- High throughput approach needed



# OPEN SCIENTIFIC REPORTS Using insects to detect, monitor and predict the distribution of Xylella fastidiosα: a case study in Corsica Accepted: 3 October 2018 Accepted: 3 October 2018 Published online: 23 October 2018 Published online: 23 October 2018





- DNA extraction (by specimen)
- 2 step PCR approach
  - first PCR: targeted region is amplified using specific primers flanked by tails
- These tails allow for a second PCR reaction to add Illumina adaptors and indexes to multiplex samples
  - MiSeq sequencing + Bioinformatics to get id.
    - monitore the spatio-temporal prevalence of Xf within populations of Ps in Corsica
    - 27 populations of 30 specimens each (more coming).
    - 4 PCR1 replicates per insect to avoid false negatives

3. High-throughput sequencing



3. Bioinformatics





High-throughput sequencing of multiple amplicons for barcoding and integrative taxonomy

Accepted: 03 January 2017 Published: 06 February 2017

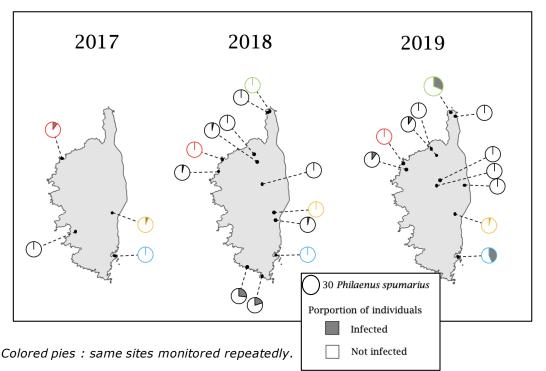
Received: 21 July 2016

Perrine Cruaud<sup>1</sup>, Jean-Yves Rasplus<sup>1</sup>, Lillian Jennifer Rodriguez<sup>1,2</sup> & Astrid Cruaud<sup>1</sup>





Preliminary results

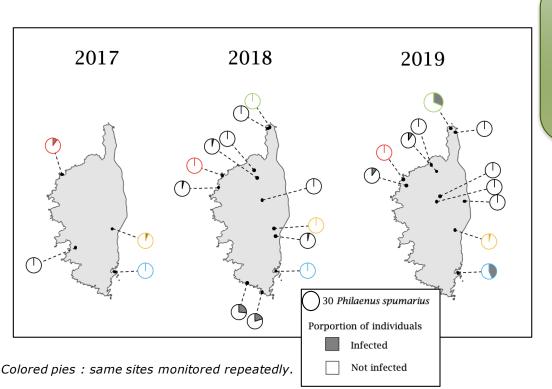


- Xf throughout Corsica
- Xf present in presumably uncontaminated areas based on plant monitoring
- No spatial, temporal or vegetationrelated pattern of prevalence emerge from our first results.





Preliminary results



- *Xf* introduction likely predates 2015
- ecological resilience of Corsican ecosystems linked to plant diversity and lack of monoculture farming?
- Sentinel insects are good complement to plant survey and their use could be generalized in EU



DOI: 10.5281/zenodo.4682793



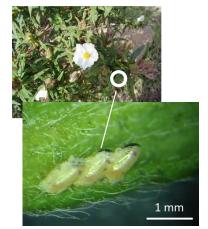
#### **PHILAENUS SPUMARIUS: POSSIBLE CONTROL?**

Overlooked research field: biological control of Ps could be an environmental friendly lever to help lower vector density

Collection of (ca 1100) eggs of Ps throughout Corsica,

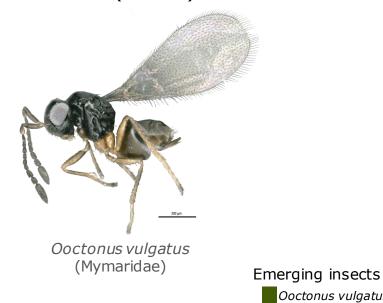
Daily monitoring of emergence

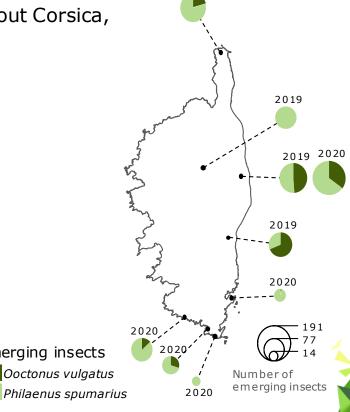
Variable parasitism rates (0-69%)





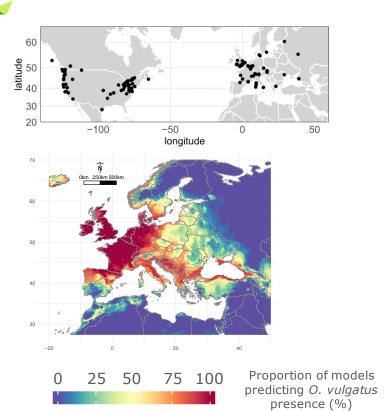






#### **PHILAENUS SPUMARIUS: POSSIBLE CONTROL?**

Occurrences + SDM → O. vulgatus occurs or is likely to occur in many EU regions where P. spumarius also occurs



Provided that host specificity is confirmed and that mass rearing is possible, *O. vulgatus* could contribute to IPM of *P. spumarius*, and more generally, of Xf pathosystem



DOI: 10.5281/zenodo.4680103

Peer Ooctonus vulgatus (Hymenoptera, Mymaridae), a potential biocontrol agent to reduce populations of Philaenus spumarius (Hemiptera, Aphrophoridae) the main vector of Xylella fastidiosa in

Europe

Xavier Mesmin<sup>1,2</sup>, Marguerite Chartois<sup>2</sup>, Guénaëlle Genson<sup>2</sup>, Jean-Pierre Rossi<sup>2</sup>, Astrid Cruaud<sup>2</sup> and Jean-Yves Rasplus<sup>2</sup>







3<sup>rd</sup> European Conference on Xylella fastidiosa and XF-ACTORS final meeting







### Thanks for your attention,

## Keep morale up, Big up for the organization & technical staff!

Astrid Cruaud<sup>1</sup>, Marguerite Chartois<sup>1</sup>, Pauline Farigoule<sup>1,2</sup>, Martin Godefroid<sup>1</sup>, Xavier Mesmin<sup>1</sup>, Ileana Quiquerez<sup>3</sup>, Sabrina Borgomano<sup>3</sup>, François Casabianca<sup>4</sup>, Guénaëlle Genson<sup>1</sup>, Anne-Alicia Gonzalez<sup>1</sup>, Laetitia Hugot<sup>3</sup>, Maxime Lambert<sup>1</sup>, Éric Pierre<sup>1</sup>, Sylvain Santoni<sup>5</sup>, Jean-Claude Streito<sup>1</sup>, Jean-Pierre Rossi<sup>1</sup> and Jean-Yves Rasplus<sup>1</sup>

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