

Implementation of the detection protocol for *Xanthomonas euvesicatoria* in pepper seeds

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The disease

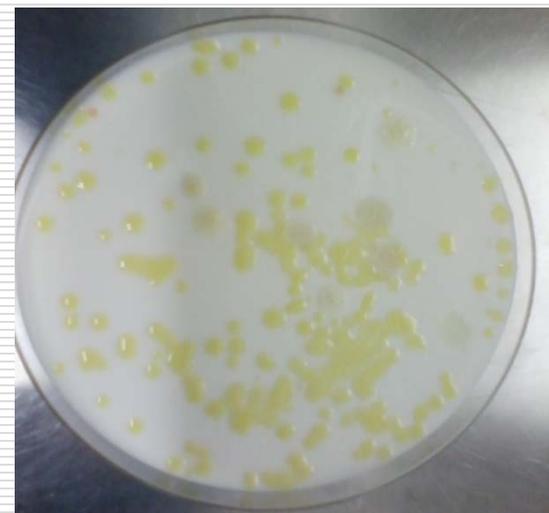
- ❑ Bacterial spot is a worldwide disease, mainly affecting tomato and pepper.
- ❑ Symptoms may affect all aerial parts and are particularly severe on pepper.



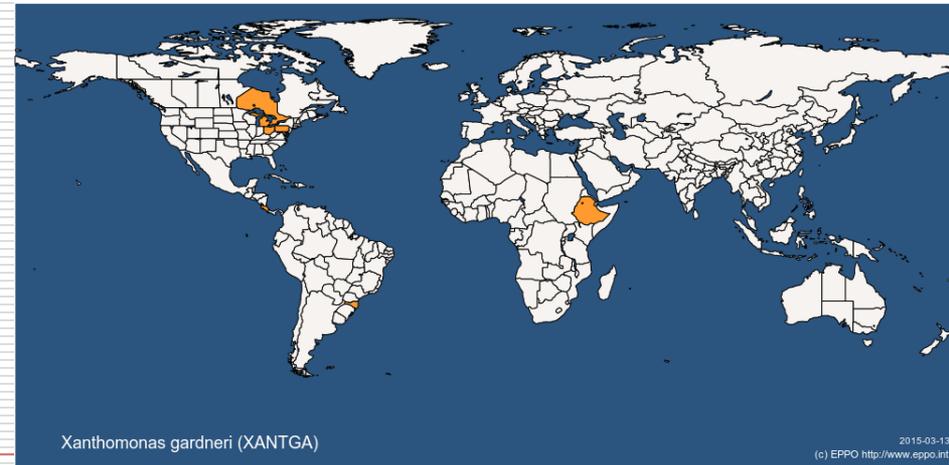
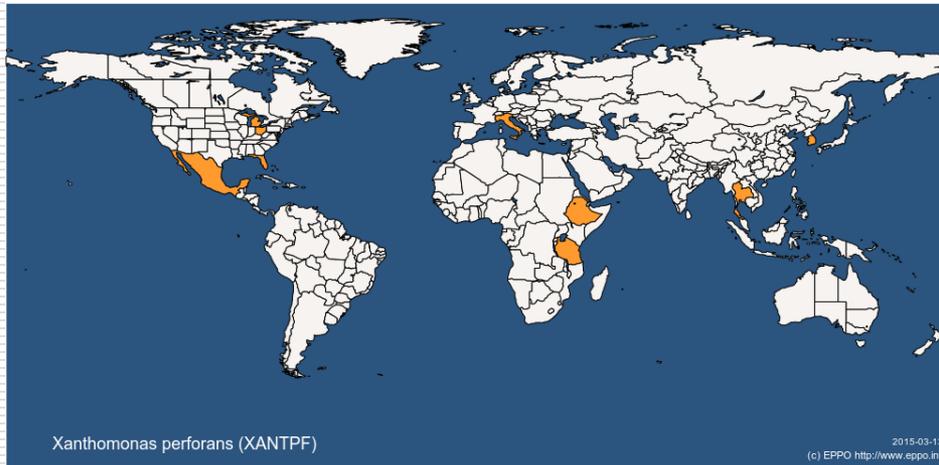
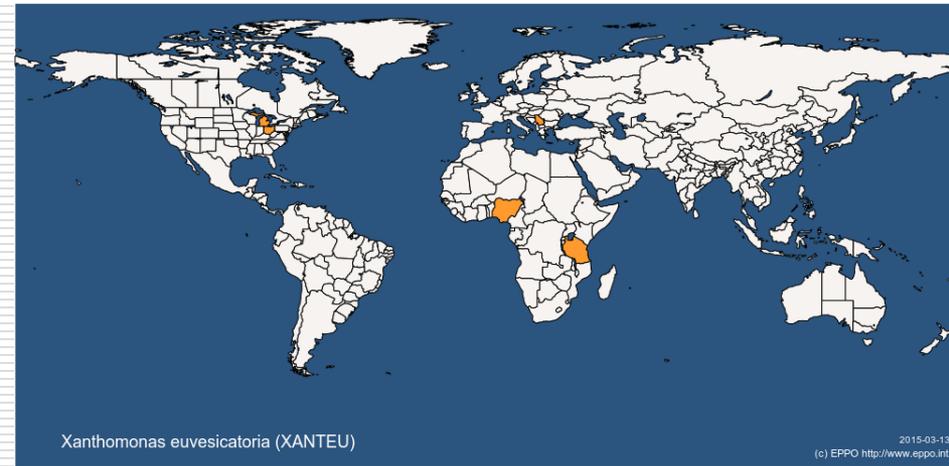
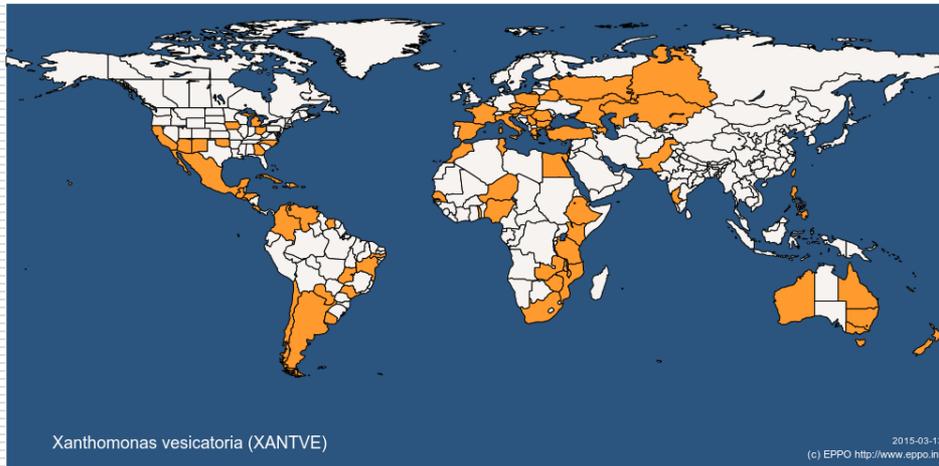
Source:
APS

The causal agent

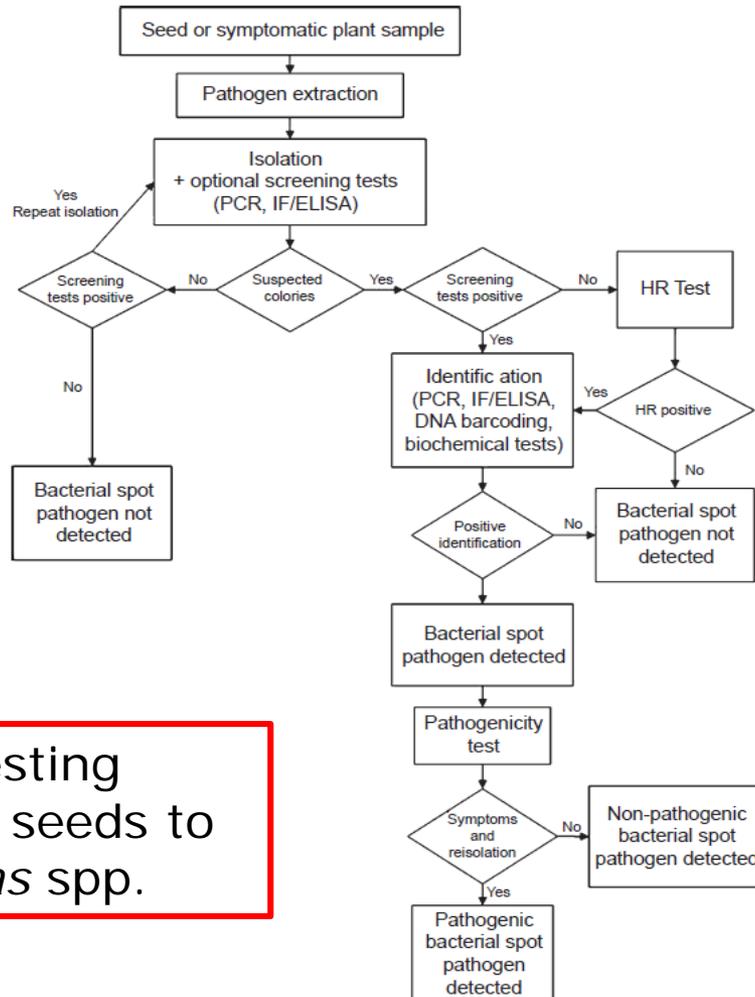
- The causal agent was formerly known as *Xanthomonas campestris* pv. *vesicatoria*.
- Now reclassified into four species:
 - *Xanthomonas vesicatoria*
 - *Xanthomonas euvesicatoria*
 - *Xanthomonas perforans*
 - *Xanthomonas gardneri*
- All four species are regulated



Pathogens distribution



The EPP0 diagnostic protocol



Flow diagram for testing tomato and pepper seeds to detect *Xanthomonas* spp.

Objective of our studies

- Implement the diagnostic protocol for pepper seeds.
 - In particular, focusing on *X. euvesicatoria*
- Compare two different DNA extraction methods.
- Compare ELISA diagnostic results with PCR and Direct Isolation on agar media.
- Suggest implementation of the DP

Material and Methods

- During the analytical season 2014 on pepper seeds in Novi Sad (Serbia), 13 seeds lots found *Xeuv* positive
 - Test applied: DAS-ELISA (Loewe)
- The second lab sample was taken for:
 - Direct isolation on YGCA
 - Preparation of seed extracts for PCR
 - Heat shock
 - DNeasy Plant Mini Kit columns (Qiagen)

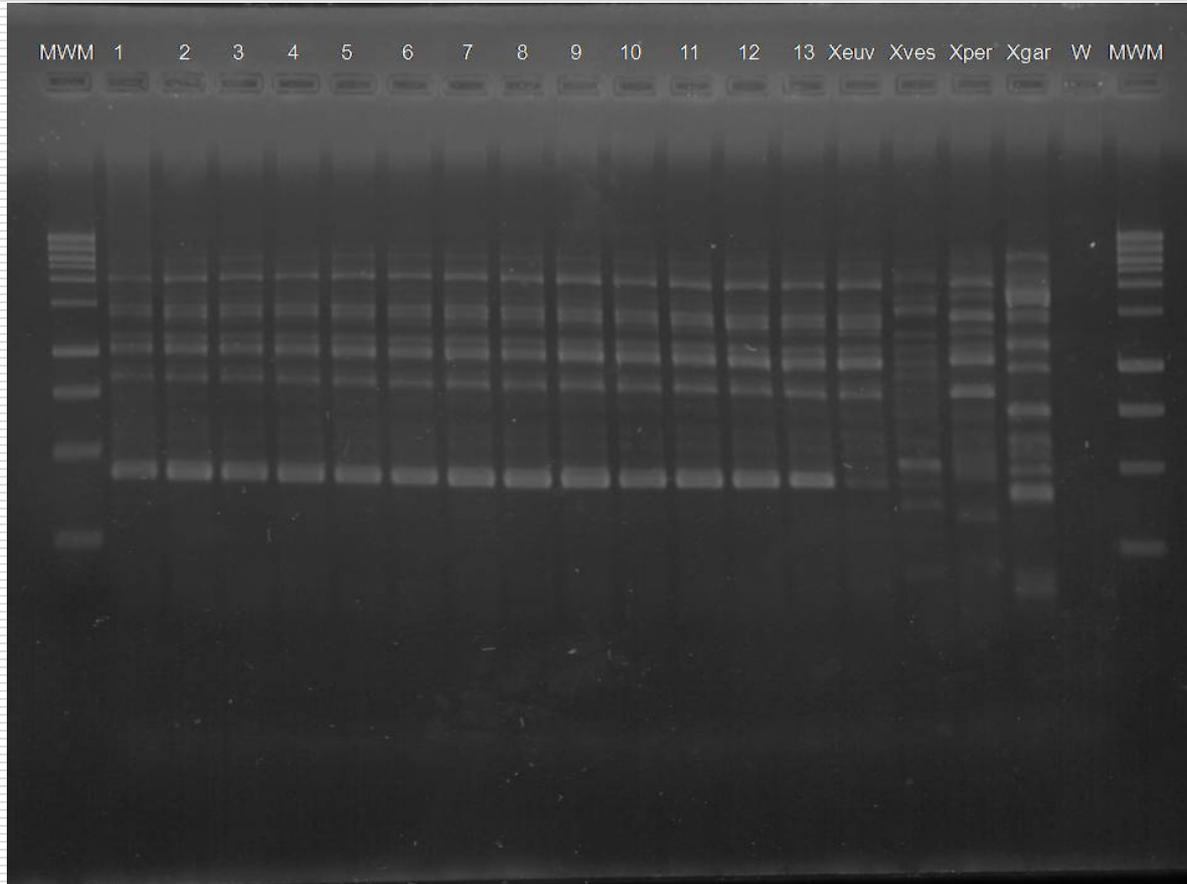
Material and Methods

- PCR protocol applied on seed extracts:
 - EPPO
 - Primer pairs (Koenraad *et al.*, 2007)
- For the identification of putative *Xeuv* colonies obtained on YGCA:
 - BOX, REP, ERIC
 - Analysis of genetic profiles:
 - UPMGA Cluster analysis (GelCompar 4.1, Applied Maths, Kortrijk, Belgium)
 - Pearson's correlation coefficient

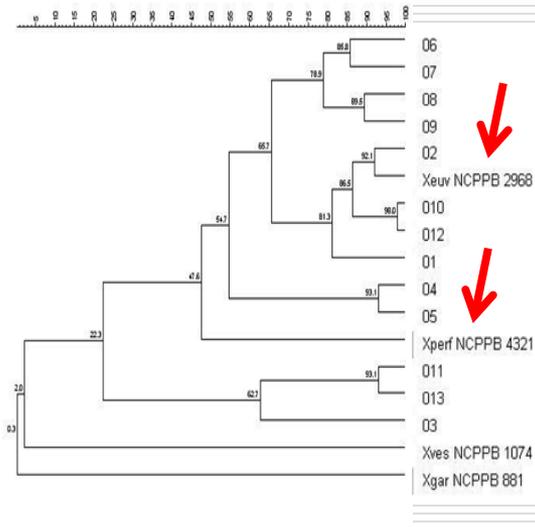
Results

| Samples (varieties or breeding lines) | ELISA | Direct isolation, confirmed by PCR on pure cultures and genotyping | PCR DNA extraction with heating shock | PCR DNA extraction with DNeasy Plant Mini Kit |
|---------------------------------------|-------|--|---------------------------------------|---|
| BL 1 2013/5 | + | - | + | + |
| BL 2 2013/8 | + | - | - | - |
| Novosađanka 1 | + | - | - | + |
| Novosađanka 2 | + | - | + | + |
| Altina | + | - | + | + |
| Anita | + | - | + | + |
| Amphora | + | - | + | + |
| Una | + | - | - | + |
| Matica | + | + | + | + |
| Soraksari | + | + | - | + |
| Vranjeska | + | + | + | + |
| Plamena | + | + | - | + |
| Manpryka - ECW | + | - | - | + |

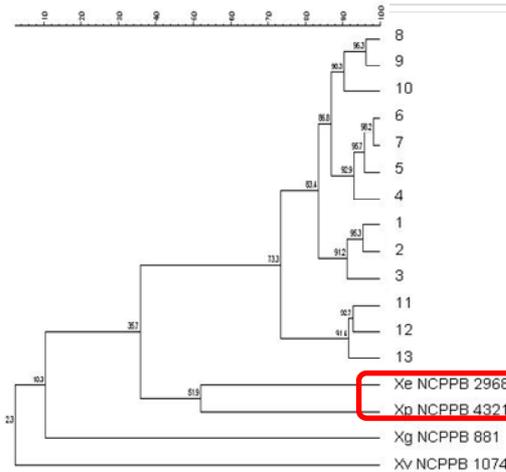
Results – ERIC PCR



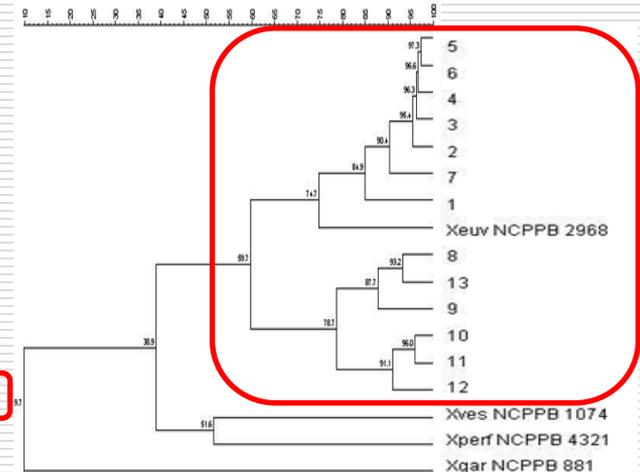
Results - Clustering



BOX



ERIC



REP

Discussion and Comments

□ Comparing results:

- PCR is best done by previous DNA extraction and purification using mini columns
- PCR confirmed most ELISA results
- Direct isolation is not so sensitive

□ Identifying *Xeuv*

- Genotyping not always able to discriminate *Xeuv* from *Xper*
- REP primers are more discriminative than BOX or ERIC

**Thank
you for
your
attention!**

