



## *Phytophthora infestans* tracking on a European scale

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*+ industrial sponsors*



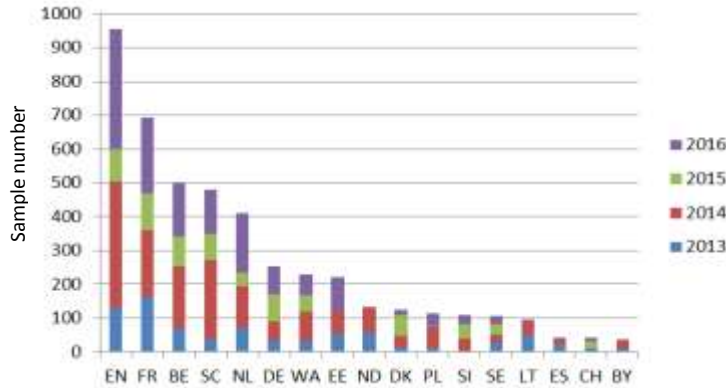
## Euroblight aims/themes

- **Tracking (global) population dynamics of potato blight pathogens:**  
Change = Trouble
- **Stewardship of host resistance genes and active ingredients** - develop and adopt innovative and sustainable control strategies on regional scales
- **Education, advocacy and communication** - taking into account different scales and stakeholders



## Sample statistics

- 2013-2016 - 5250 samples from 34 countries genotyped
- Northwestern Europe most frequently sampled

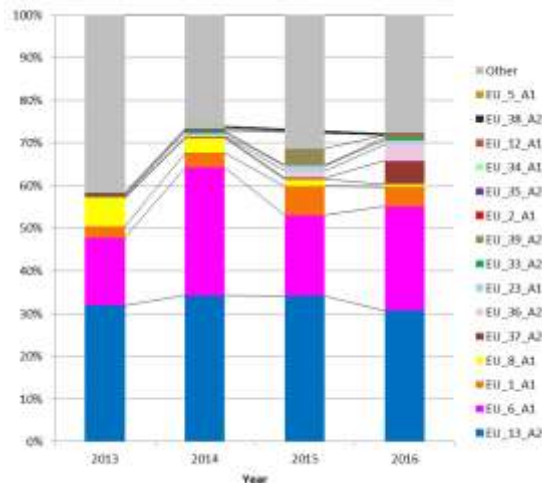


- EN – England
- FR – France
- SC – Scotland
- BE – Belgium
- NL – the Netherlands
- DE – Germany
- WA – Wales
- ND – Northern Ireland
- SE – Sweden
- PL – Poland
- DK – Denmark
- ES – Spain
- SI – Slovenia
- BY – Belarus
- CH – Switzerland
- EE – Estonia
- LT – Latvia



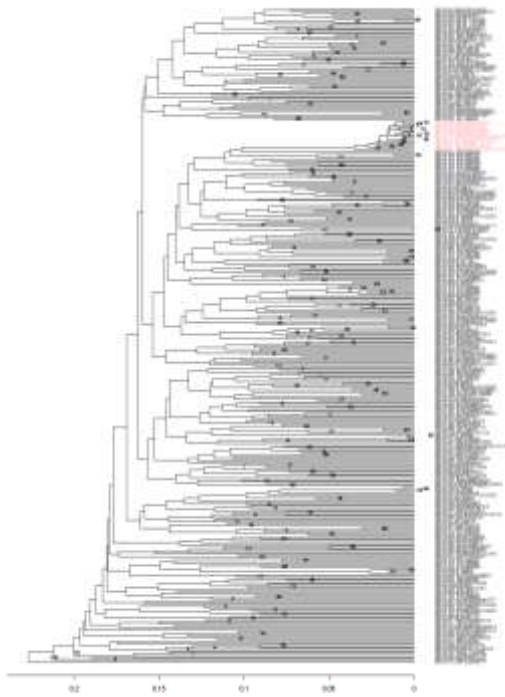
Countries with >30 samples shown

## EU *P. infestans* genotype change

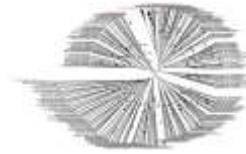


- Around 70% of EU population clonal
- EU\_13\_A2 & EU\_6\_A1 dominant
- EU\_33\_A2 'green 33' very low frequency
- **EU\_36\_A2 & EU\_37\_A2 novel clones increased in 2016**
- 20-30% of population each year genetically diverse 'Other' group probably from oospores. The *P. infestans* "gene pool"



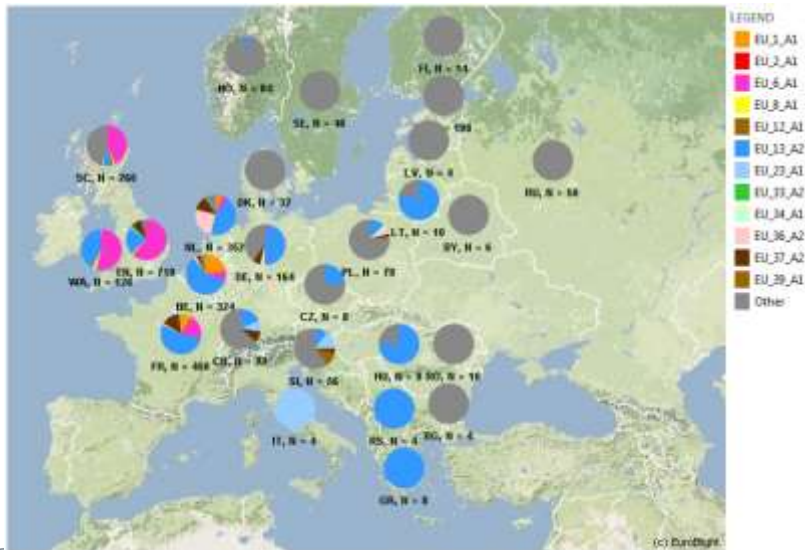


## 2016 – ‘Others’

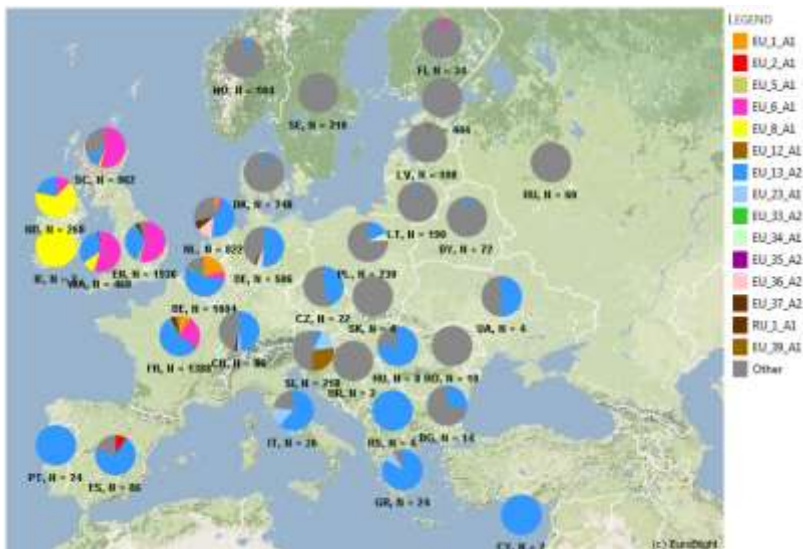


- Considerable genetic diversity in the sexual ‘Other’ population
- Minor genetic variation within asexual clones.

## 2016 data

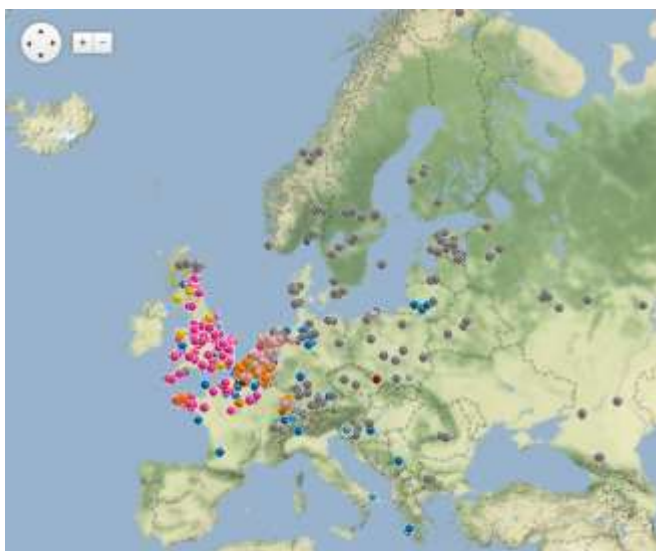


## 2013-2016 summary



• 5251 samples from 34 countries

## 2016 samples



• 1595 samples from 26 countries

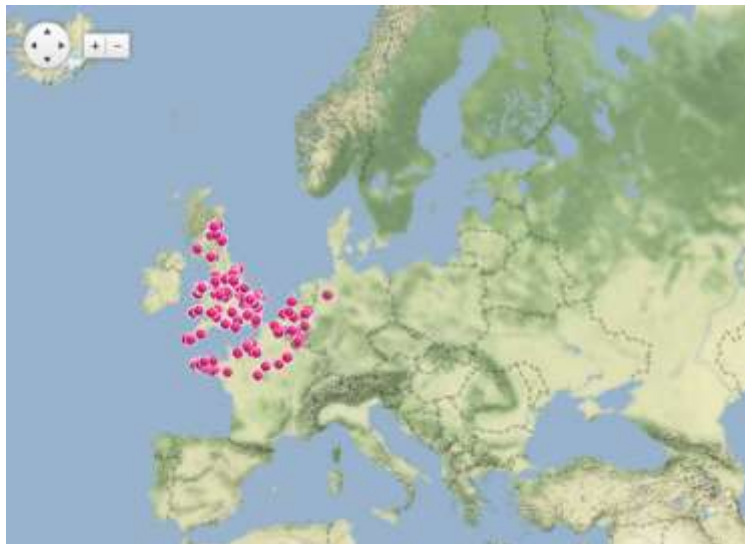
## 2016 samples – EU\_13\_A2



 **EuroBlight**  
A potato late blight network for Europe

 Live map at  
<http://euroblight.net/pathogen-characteristics-and-host-resistance/sampling-sites-and-genotype-maps/>

## 2016 samples – EU\_6\_A1

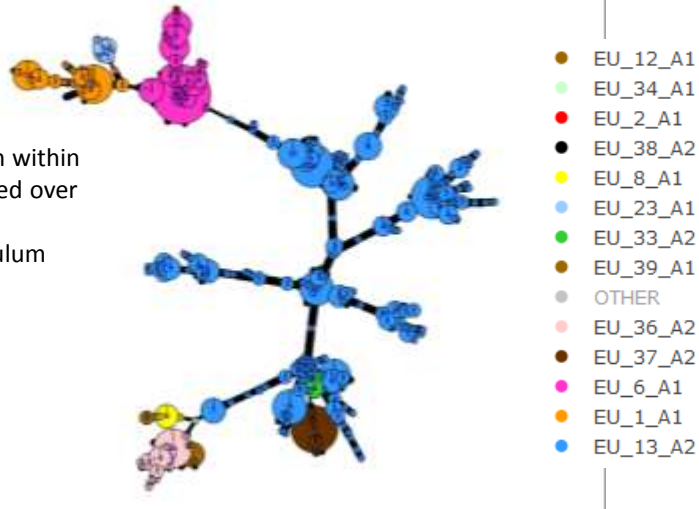


 **EuroBlight**  
A potato late blight network for Europe

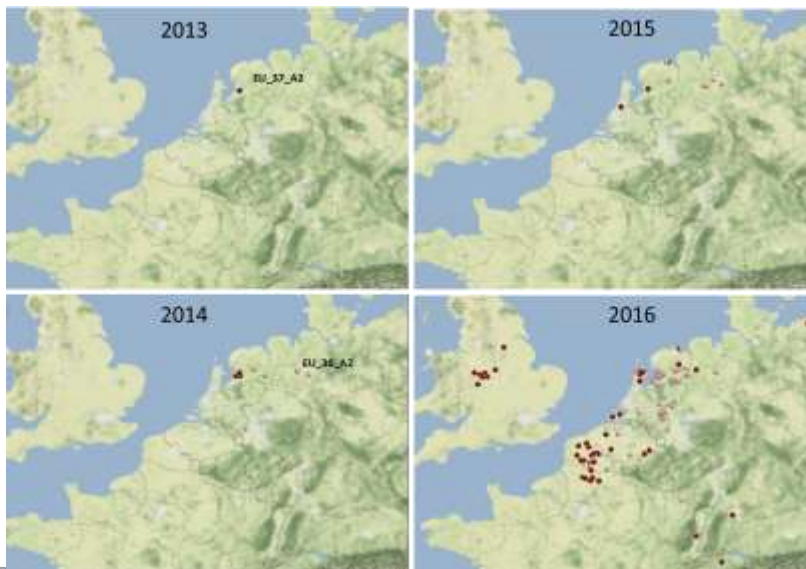
 Live map at  
<http://euroblight.net/pathogen-characteristics-and-host-resistance/sampling-sites-and-genotype-maps/>

## Genetic diversity 2016 – MSN clones only

- Minor variation within clones generated over time
- Can track inoculum spread



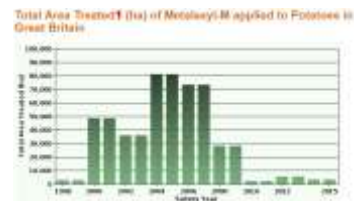
## EU\_36\_A2 & EU\_37\_A2 timeline



Live maps at <http://euroblight.net/pathogen-characteristics-and-host-resistance/sampling-sites-and-genotype-maps/>

## Population change and blight management

- **13 different fungicide groups on market for blight management**
- **Careful use required to reduce selection pressure**
- **EU\_13\_A2** emerged 2004, resistant to metalaxyl. Usage has fallen dramatically in Europe but it is still a good product where resistance is not dominant
- **33\_A2** insensitivity to fluazinam
- **37\_A2** insensitivity to fluazinam



Live maps at

<http://euroblight.net/pathogen-characteristics-and-host-resistance/sampling-sites-and-genotype-maps/>

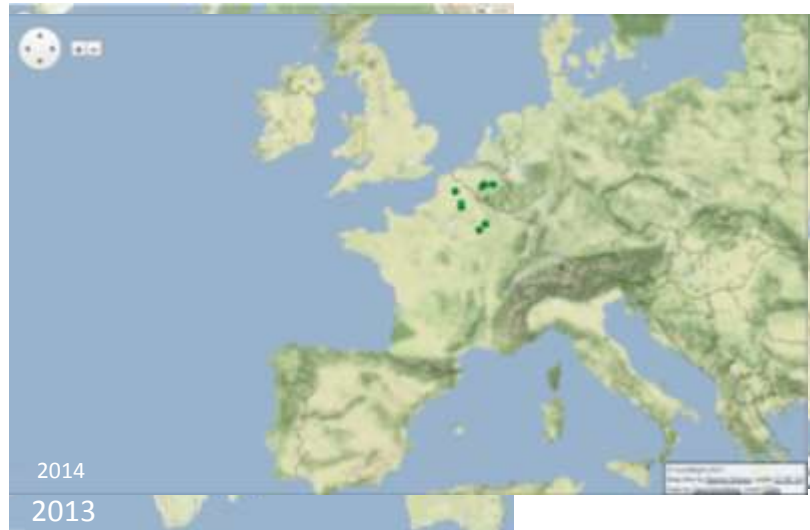
## Green33 in the Netherlands 2008 - 2012



- Green33
  - NL2010: 20%
  - NL2011: 22%
  - NL2012: 6%



## Green33 in recent years



## Press release on EU-37-A2





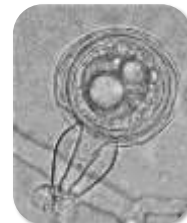
## Clones – key questions

- **Origin – where and how?**
  - Several first reported in NL and DE
  - Presumed recombinant – commonly triploid  
(36\_A2 triploid, 37\_A2 diploid) - (see Li et al., 2017 MPMI)
- **Drivers of spread?**
  - Adaptation: Mutation – Selection: host, fungicide, survival, aggressiveness
  - Chance – bottleneck, right place at right time



## Conclusions

- New insights on pathogen diversity
- **Dominance of a few clones** across large areas of European crops shows that EU growers/industry **share management challenges**
- Surveys needed to inform fungicide use strategies to minimise the risks of product failure – **13\_A2, EU\_37\_A2 and EU\_33\_A2**
- Much **primary inoculum is locally generated** and spread. Better management of inoculum sources would aid management
- A highly diverse gene pool of novel types is present as a result of **sexual oospore formation and derived infections**
- **High genetic diversity increases the risk** of blight management problems - evolving virulence against novel host resistance genes and reduced sensitivity to specific fungicide active ingredients
- Phenotypic traits of existing and novel genotypes being examined in ERA-NET project (IPMBlight2.0)
- Live data mapped on [www.euroblight.net](http://www.euroblight.net)



## Europe sponsors/contributors/collaborators

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