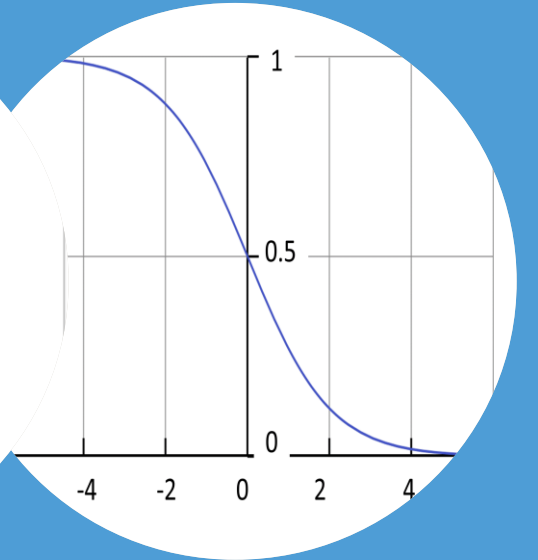


Estimating the rate of spread of Olive Quick Decline Syndrome, caused by *Xylella fastidiosa* ssp. *pauca*, in Puglia

April 30, 2021, Lia Hemerik

David Kottelenberg, Maria Saponari, Wopke van der Werf



Spreading of *X. fastidiosa* into Italy

- In 2000 an epidemic of PD occurred in the United States in southern California
 - Spread by an invasive species of sharpshooters
- In 2013 *X. fastidiosa* was first identified in Europe in Gallipoli (Apulia, Italy)
 - Subspecies *pauca*
 - Causing olive quick decline syndrome (OQDS)



Location of Gallipoli

Map data ©2019 GeoBasis-DE/BKG (©2009), Google Inst. Geogr. Nacional

Olive Quick Decline Syndrome (OQDS)

- Leaf scorching
- Desiccation of twigs and branches
- After a few years, the tree dies
- Large economic losses
 - Olive and olive oil are main sources of income for the region
 - No export of plants
- Cultural impact
 - Family orchards
 - Centuries old trees



Olive tree suffering from OQDS

Vector: Spittlebug (*Philaenus spumarius*)

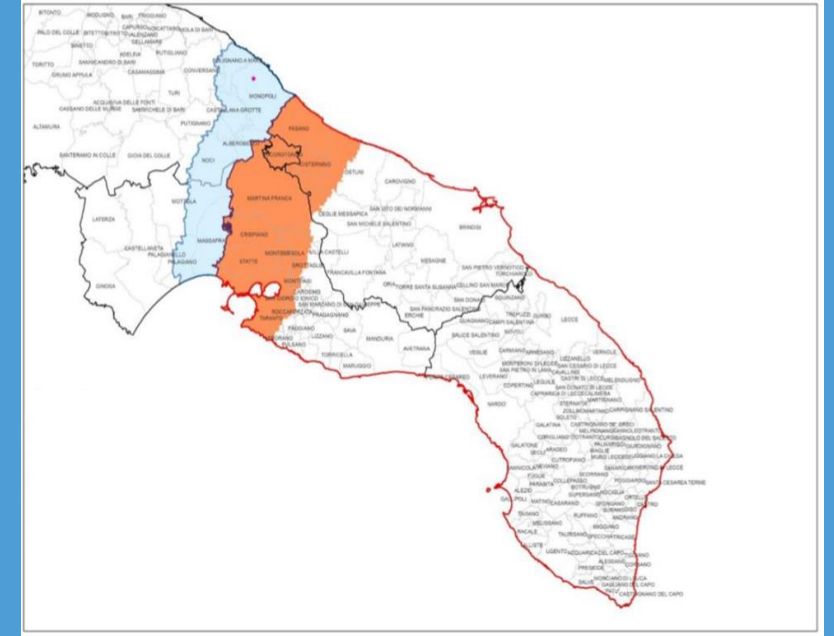
- Can be found throughout most of Europe
- Can survive in many climate types
- Female migrating behavior and dispersal cause short range spread of *X. fastidiosa*
- Possibly hides in clothing or vehicles, causing long range spread of *X. fastidiosa*



Spittlebug

Models

- Many models have been developed
 - Spatially explicit spread model (White et al., 2017)
 - SIR model (Soubeyrand et al., 2018)
 - Network model (Strona, Carstens & Beck, 2017)
- No models analyzing the shape of the invasion front or rate of spread
 - Even though data is available
- Expert knowledge before study: in one year time 90% of newly infected plants will be within 5.2 km from previously infected area (95% CI of 0.7 – 14 km)

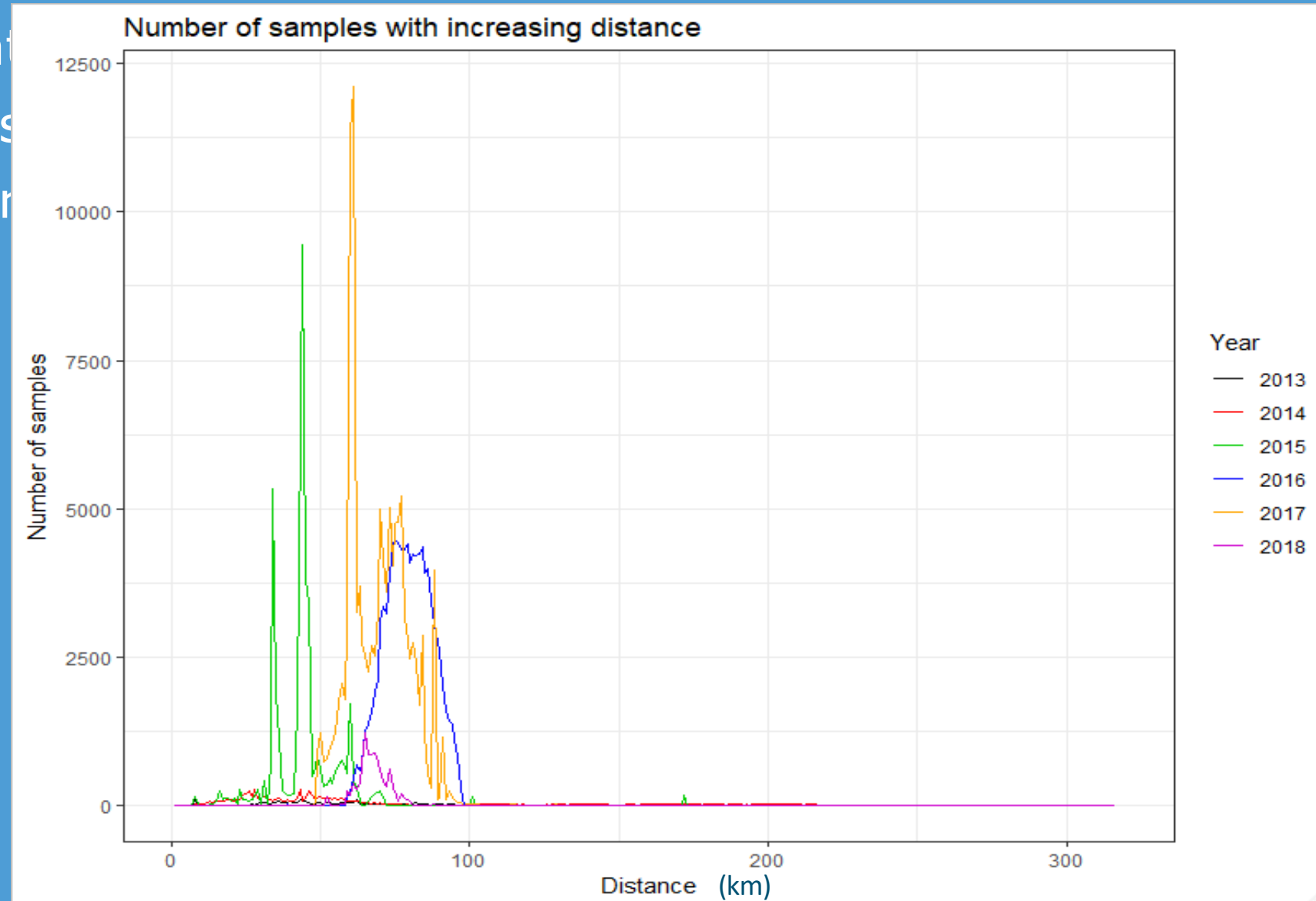


Demarcated zones in Puglia, Italy (European Union, 2019).

Sampling data

- Spatio-temporal data
- Very heterogeneous
- Until 2016, no clear

Locations of sampling in Puglia



Research questions and Data

- What is the shape of the travelling invasion front of *Xylella fastidiosa* in Puglia?
- What is the rate at which this front moves through Puglia?
- What is the width of the front?
- The columns
 - no: number, lon: longitude, lat: latitude
 - Result
 - 0: *Xylella* is absent (these I call negatives)
 - 1: *Xylella* is present (these I call positives)
 - Date, Year, Month
- 409,515 rows
 - 298,230 rows after removal of NA's and missing coordinates

	no	lon	lat	result	date	year	month
1	1	17.95389	40.50889	0	2013-11-13	2013	11
2	2	17.92639	40.52167	0	2013-11-13	2013	11
3	3	17.97722	40.54639	0	2013-11-13	2013	11
4	4	18.02444	40.48750	0	2013-11-13	2013	11
5	5	18.04250	40.48694	0	2013-11-13	2013	11
6	6	18.02056	40.52417	0	2013-11-13	2013	11
7	7	18.06889	40.53000	0	2013-11-13	2013	11
8	8	18.08194	40.51389	0	2013-11-13	2013	11
9	9	18.04611	40.53167	0	2013-11-13	2013	11
10	10	17.86722	40.53806	0	2013-11-13	2013	11

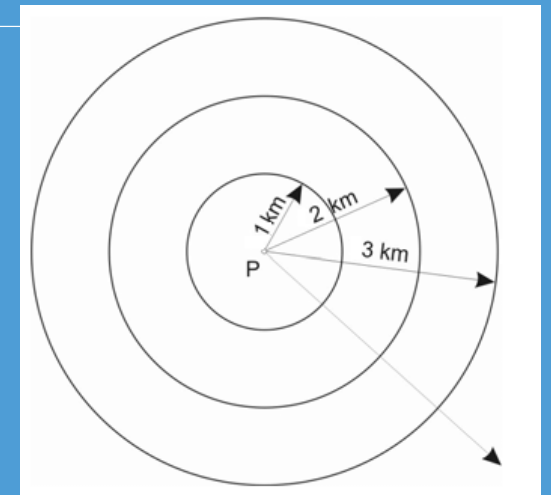
Methods: Rate of Spread

- Eight datasets (four types of information sharing & two temporal cut-offs)

- Data transformation:

Calculate for each calendar year or year from April to April, and in each concentric ring

- number of positives,
- number of measurements,
- proportion of positives (= number of positives / number of measurements)



Adapted from Kloch & Krynski, 2008.

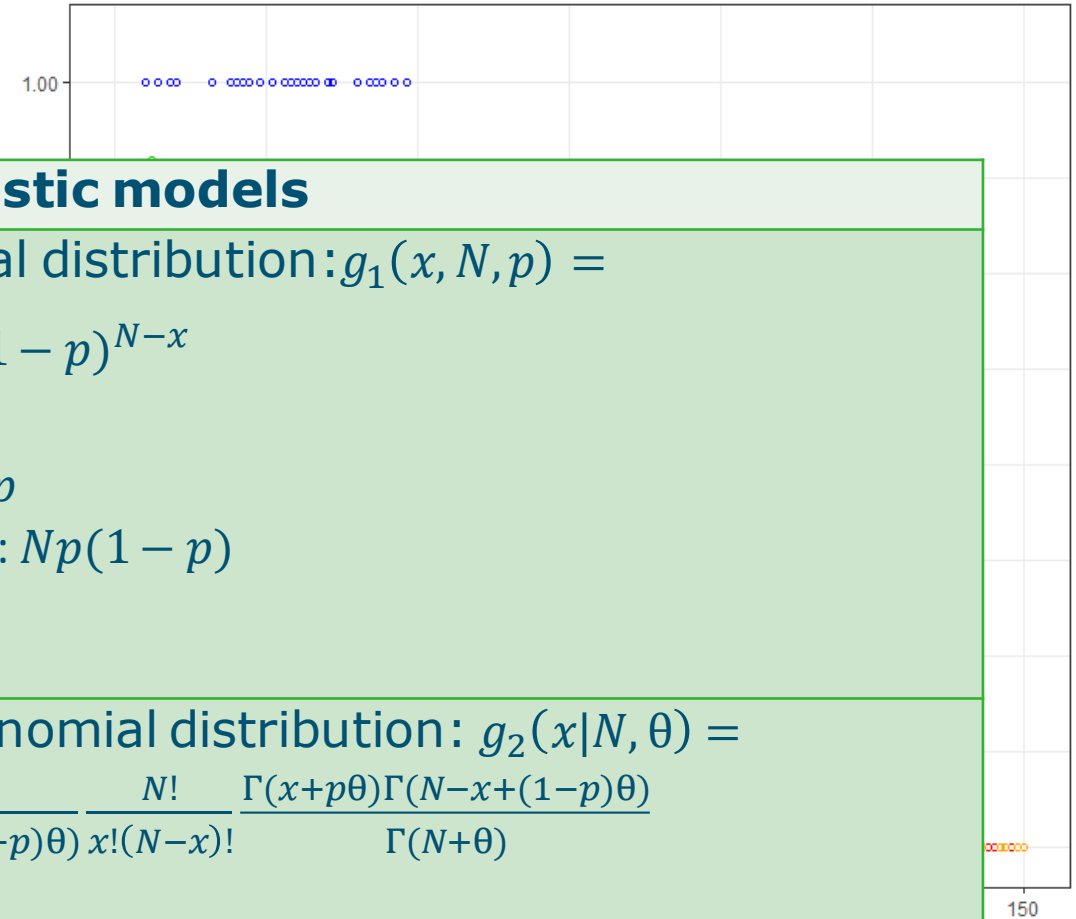
Concentric circles around a central point with 1 km widths.

	dist	pos	n	prop
1	1	0	0	NaN
2	2	0	0	NaN
3	3	0	0	NaN
4	4	0	0	NaN
5	5	0	0	NaN
6	6	0	2	0.00000000
7	7	0	9	0.00000000
8	8	14	127	0.11023622
9	9	0	11	0.00000000
10	10	0	6	0.00000000

The transformed data. Showing the first 10 rows of the 2013 data frame.

Data and models

Proportion of positives with increasing distance



Deterministic models

Negative exponential (NE):

$$f_1(x) = a \exp(-rx)$$

Logistic function:

$$f_2(x) = \frac{1}{1 + \exp(r(x - x_{50}))}$$

Constrained negative exponential function (CNE):

$$f_3(x) = \begin{cases} 1 & \text{if } x < x_{100}, \\ \frac{1}{\exp(-r(x_{100}))} \exp(-rx) & \text{if } x \geq x_{100}. \end{cases}$$

Hill function:

$$f_4(x) = a \left(1 - \frac{x^n}{h^n + x^n} \right)$$

Stochastic models

Binomial distribution: $g_1(x, N, p) =$

$$\binom{N}{x} p^x (1-p)^{N-x}$$

Mean: Np

Deviance: $Np(1-p)$

Beta-binomial distribution: $g_2(x|N, \theta) =$

$$\frac{\Gamma(\theta)}{\Gamma(p\theta)\Gamma((1-p)\theta)} \frac{N!}{x!(N-x)!} \frac{\Gamma(x+p\theta)\Gamma(N-x+(1-p)\theta)}{\Gamma(N+\theta)}$$

Mean: Np

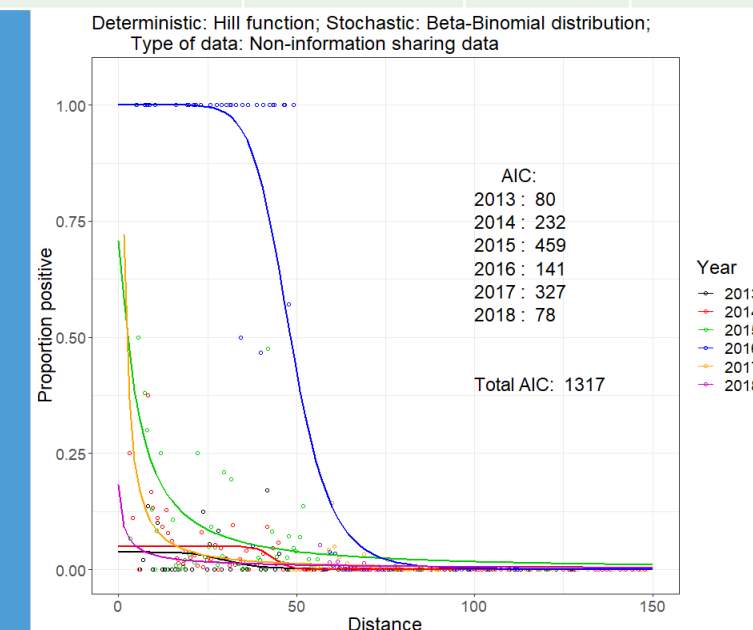
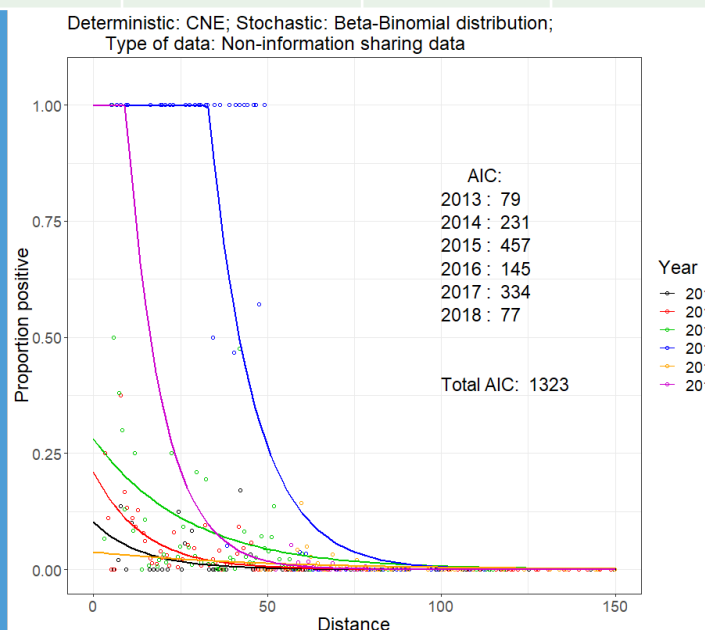
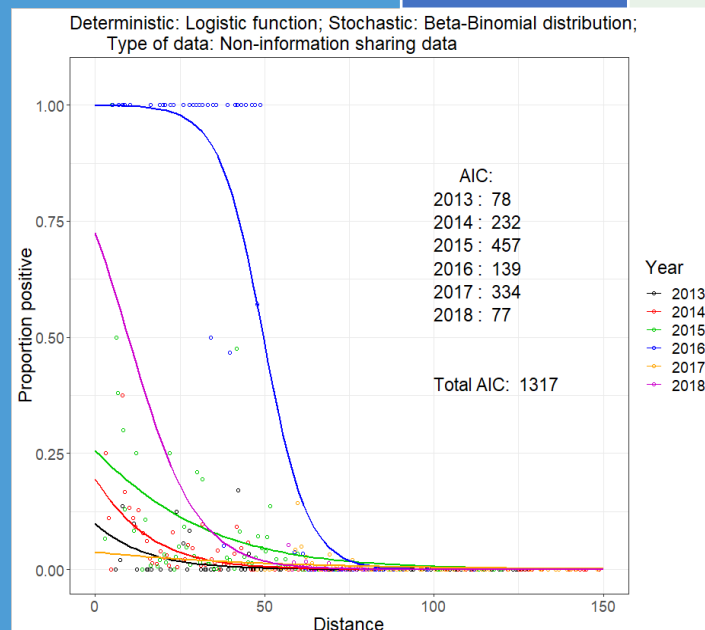
Deviance: $Np(1-p) \left(1 + \frac{N-1}{\theta-1} \right)$



Results

AIC

	Binomial distribution				Beta-binomial distribution			
Year	Negative exponential	Logistic	CNE	Hill	Negative exponential	Logistic	CNE	Hill
2013	159	159	159	159	78	78	78	80
2014	344	344	344	356	232	232	232	232
2015	8853	8847	8847	8133	457	457	457	459
2016	587	239	239	265	221	139	145	141
2017	4673	4442	4442	4612	334	334	334	327
2018	87	87	87	90	108	77	77	78
Total	14703	17227	14169	13603	1430	1317	1323	1317



Determining the rate of spread

■ Logistic function: $pp_l = \frac{1}{1 + \exp(r(x - (x_{50} + ct)))}$

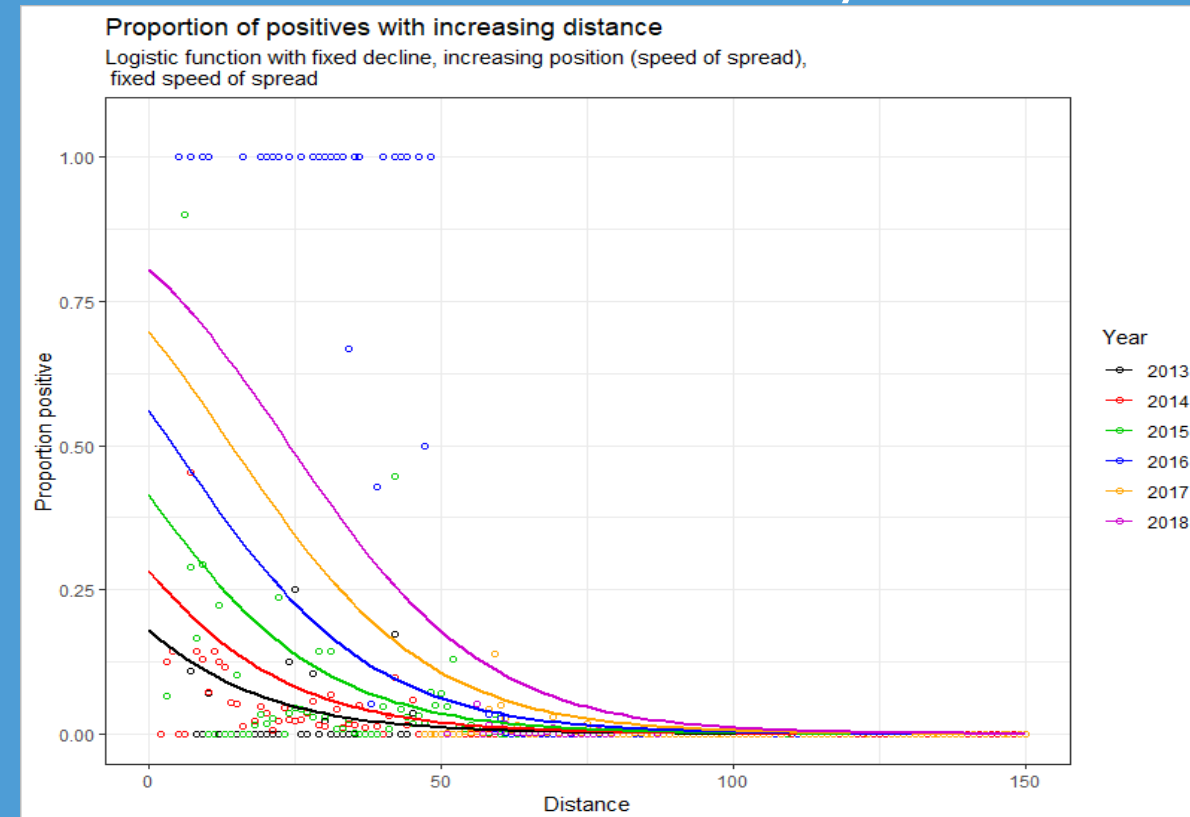
c : rate of spread in km/year
 t : time in years since 2013

Assumption:

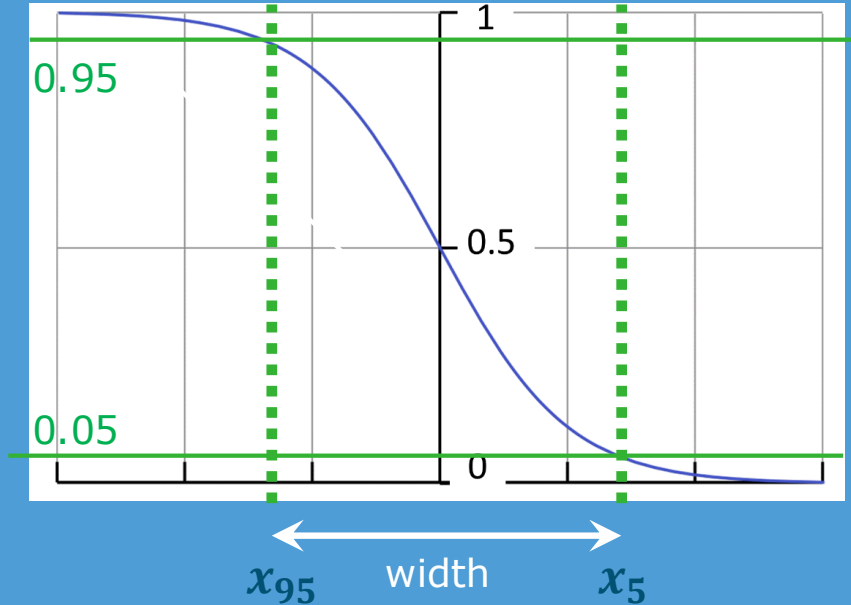
The rate of spread and the shape of the front are constant over the years

Parameters	
r	0.059 km ⁻¹
x_{50}	-25.76 km
c	9.95 km/year

The fitted logistic curves to estimate the rate of spread.



Width of front



Starting with

$$\frac{1}{1 + \exp(r(x_{95} - (x_{50} + ct)))} = 0.95$$

$$\frac{1}{1 + \exp(r(x_5 - (x_{50} + ct)))} = 0.05$$

We get

$$x_5 - x_{95} = \frac{2 \log(95)}{r}$$

Resulting in an estimate of

$$x_5 - x_{95} = 99.8 \text{ km.}$$

$$x_1 - x_{99} = 2 \log(99) / 0.059 = 155.8 \text{ km.}$$

Discussion

- First analysis that estimates the shape of an invasion front
- Non-trivial task because of the difficult data
- Because of this we have simulated the sampling process
- and performed simulations with different assumed points of origin
- First empirical estimate of the rate of spread of an *X. fastidiosa* invasion in Europe 10 km per year (95% CI 7.5-12.5 km per year)
- The width of the front is 100 to 160 km
- This knowledge on the spread of *X. fastidiosa* can aid management decisions in controlling the disease



Thanks to David Kottelenberg, Maria Saponari and Wopke van der Werf

Thank you for your attention!

