

EUPHRESCO Kick-off Meeting, PHYTFIRE WP3

Population analysis via MALDI-TOF ID

What goes on during fire blight treatment?

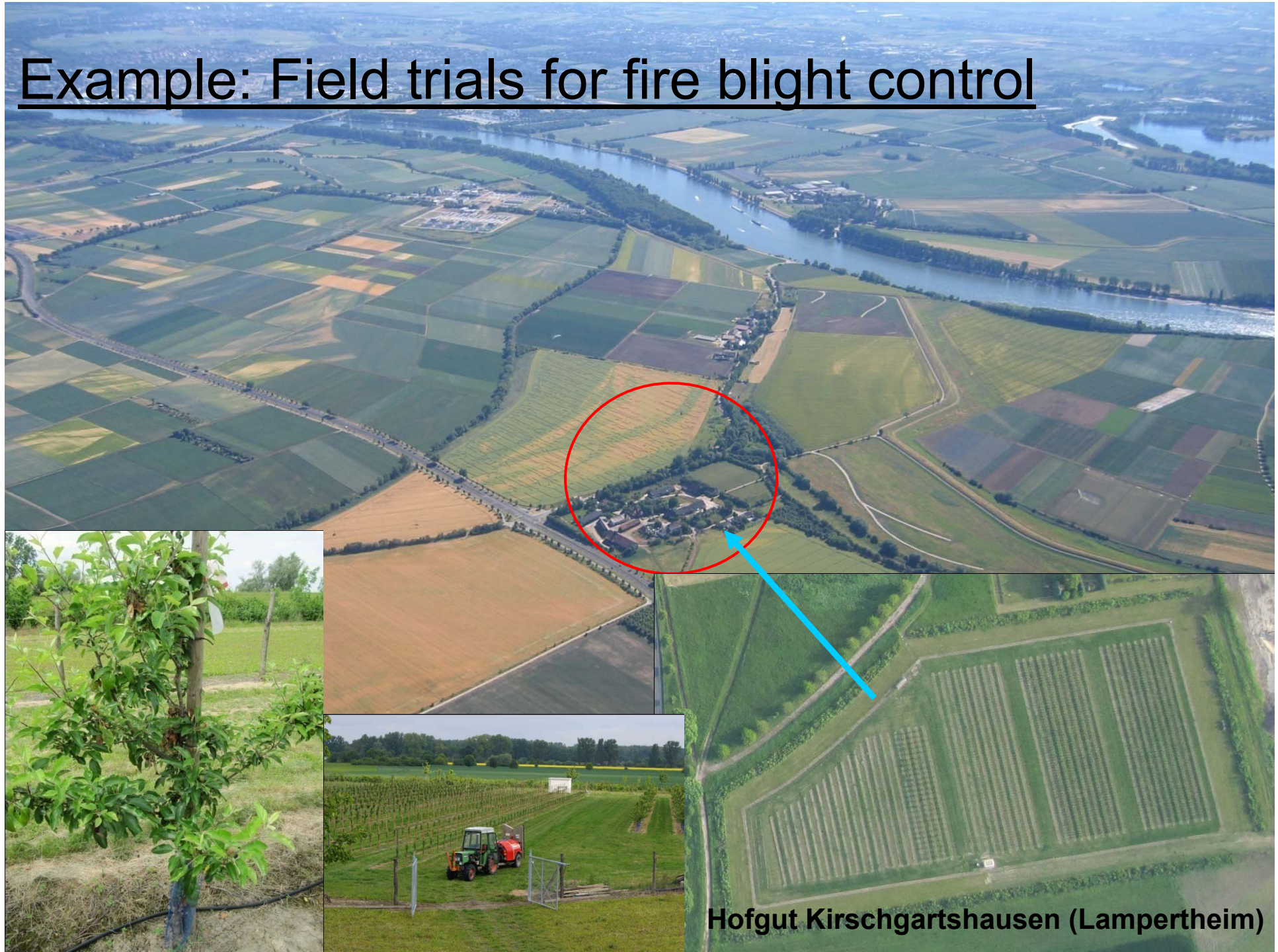
Dr. Annette Wensing, JKI Dossenheim, Germany



PHYTFIRE



Example: Field trials for fire blight control



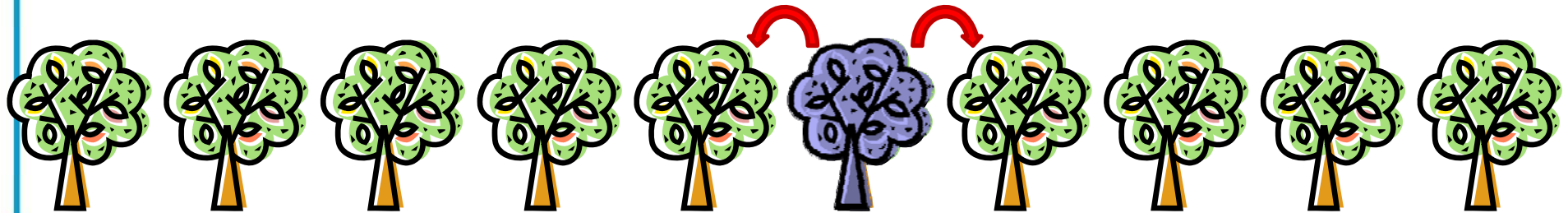
Hofgut Kirschgartshausen (Lampertheim)

Field trials



Setup (EPPO PP 1/166(3))

4 blocks with 10 trees each per variant



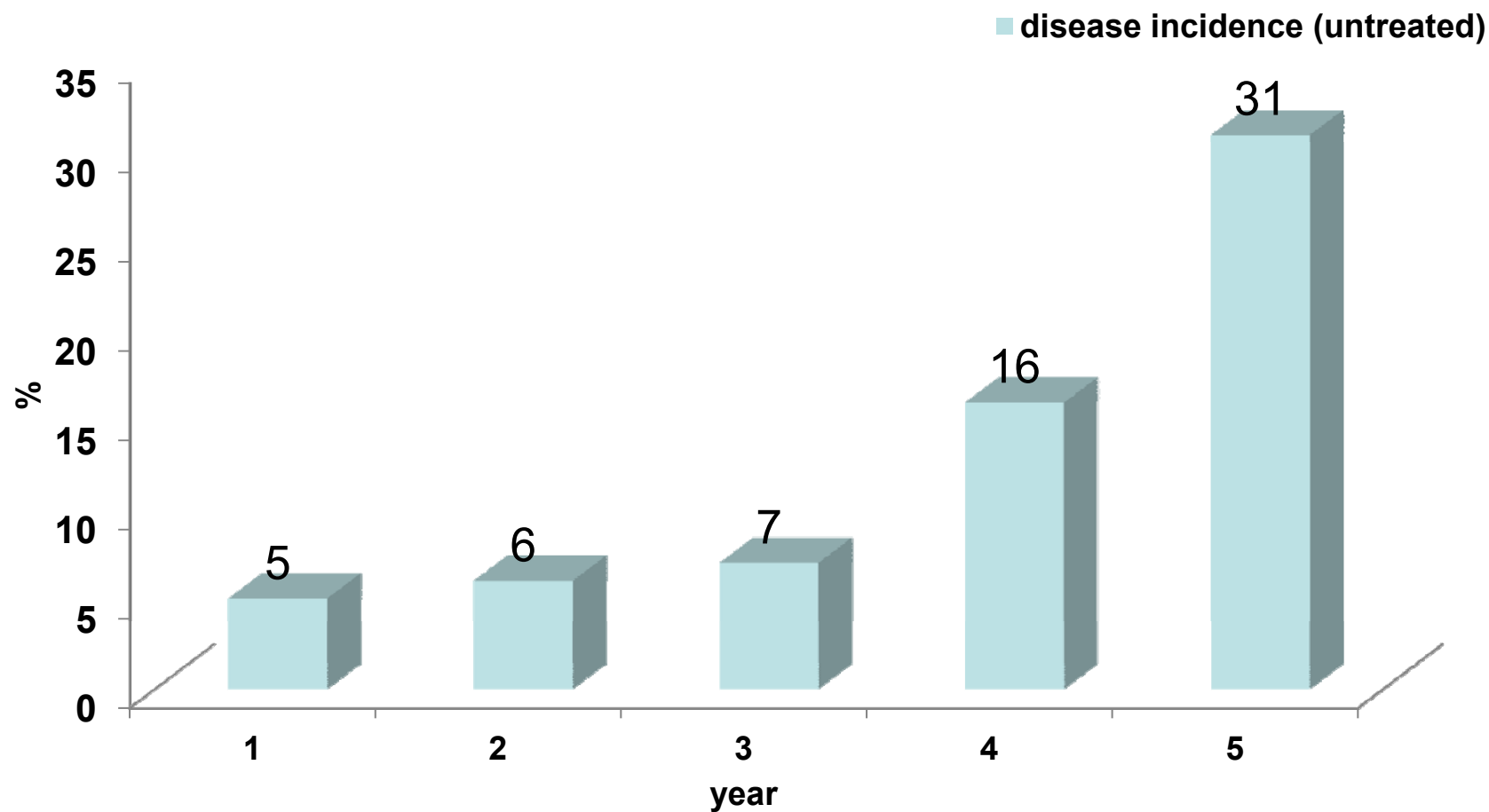
artificial inoculation of one tree/block (10^8 cfu/ml)



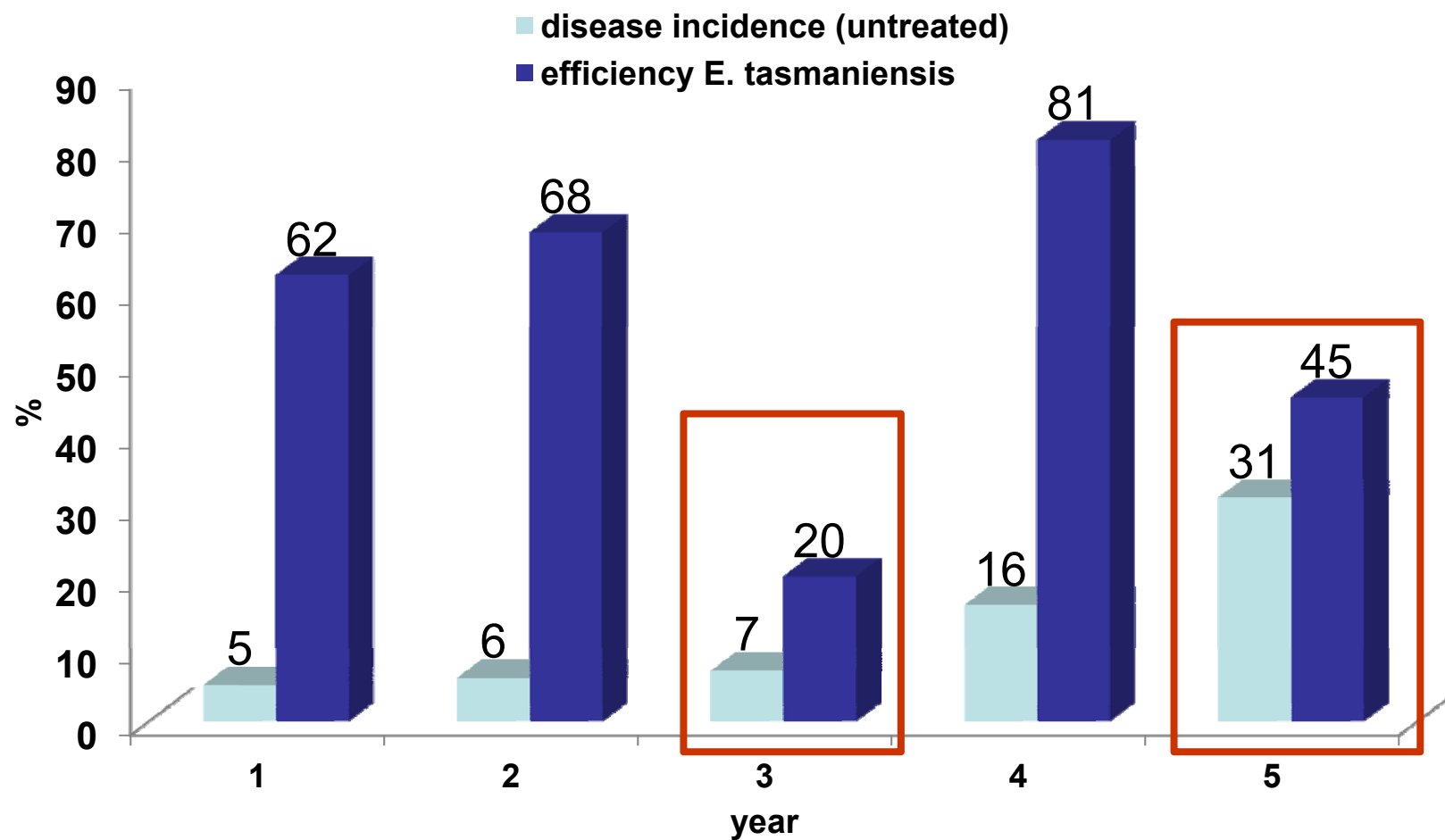
secondary infection of neighboring trees via natural vectors

→ **defined conditions**

Differences in disease incidence



Differences in efficiency



Objective:

What happens on the flower?

Application

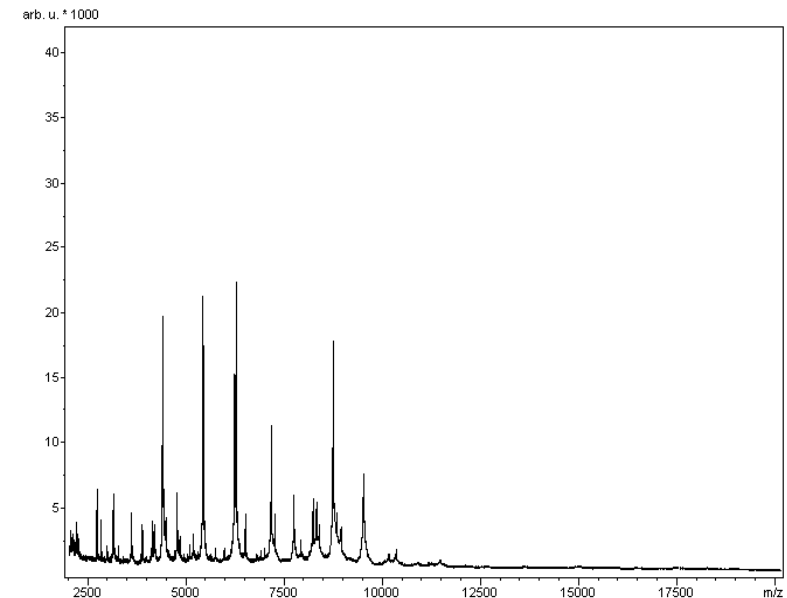
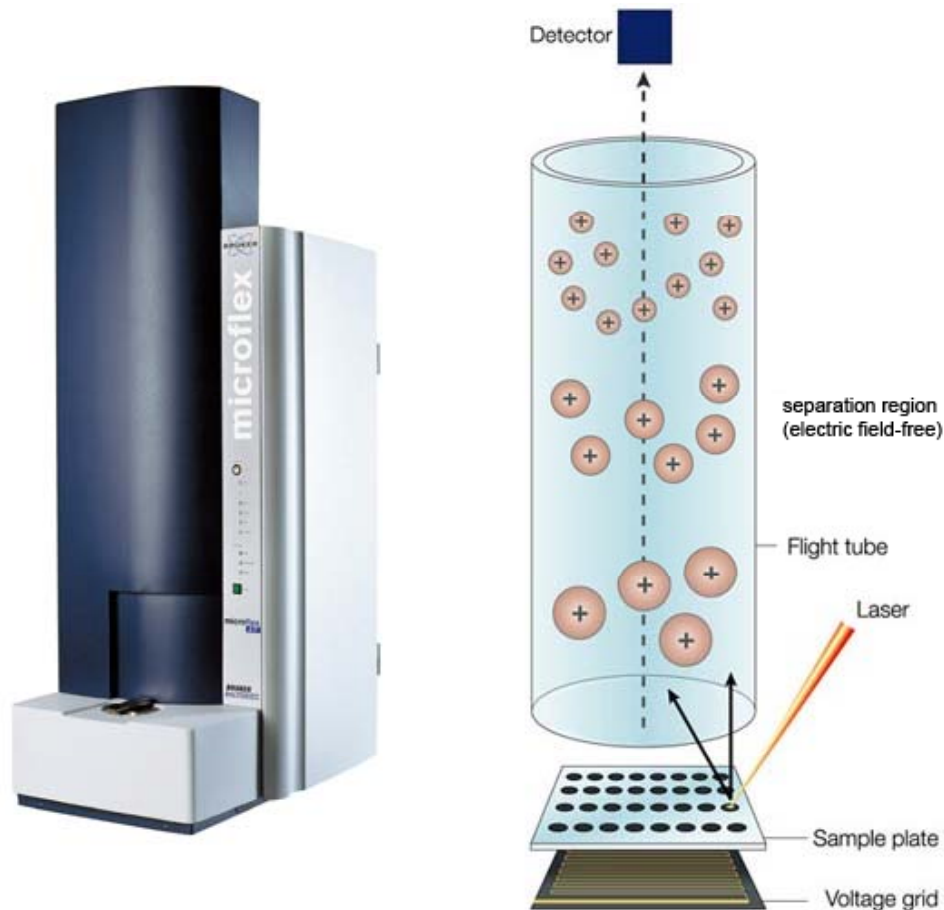
?

Result

- screening for specific organisms
- evaluation of population composition

MALDI-TOF

Matrix assisted laser dissorption/ionisation time of flight analysis



Claydon et al., 1996; Holland et al., 1996;
Krishnamurthy et al., 1996; Keys et al., 2004

www.jki.bund.de

Protein fingerprinting

Erwinia amylovora CFBP1232

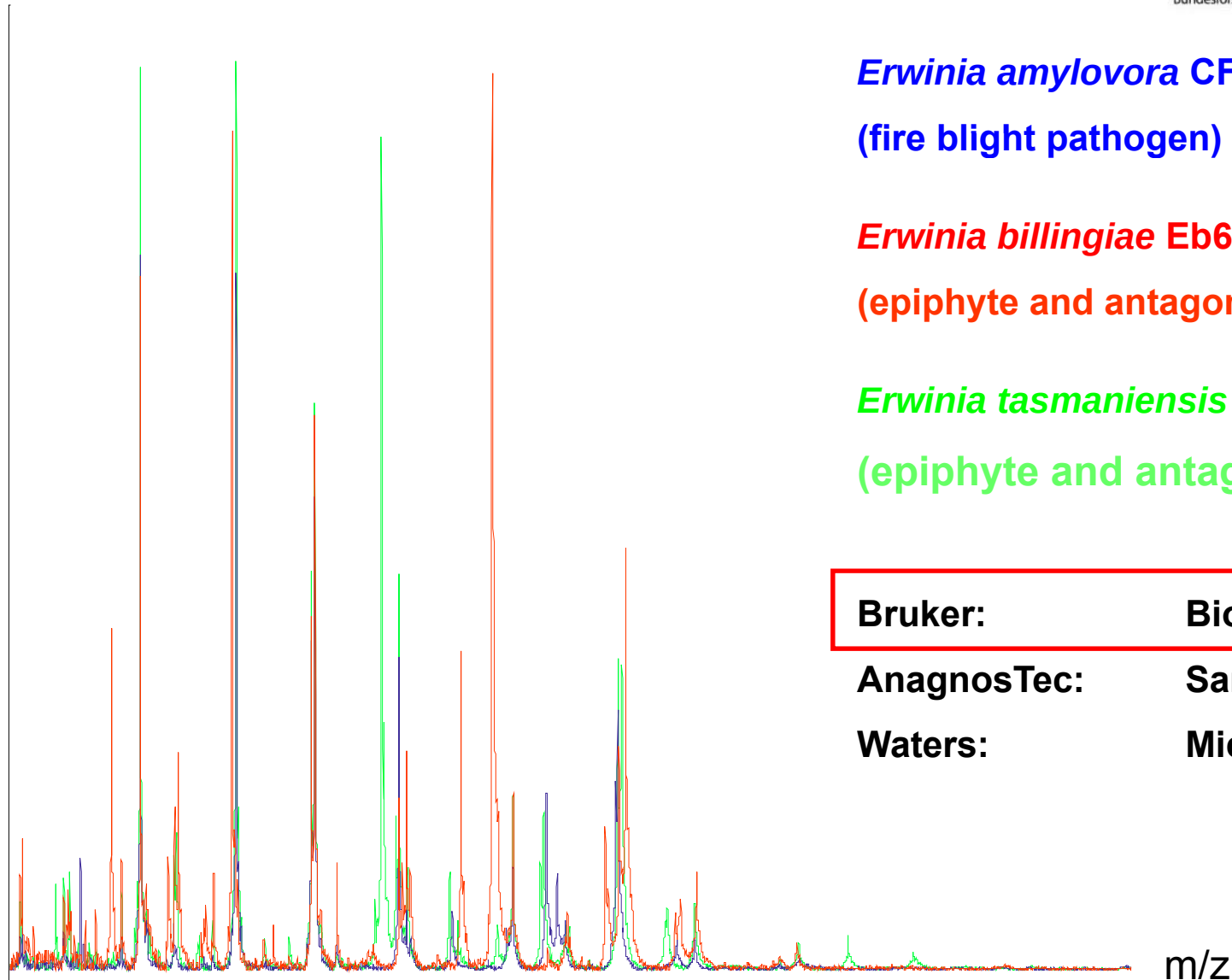
(fire blight pathogen)

Erwinia billingiae Eb661

(epiphyte and antagonist)

Erwinia tasmaniensis Et1/99

(epiphyte and antagonist)



Bruker:

Biotyper

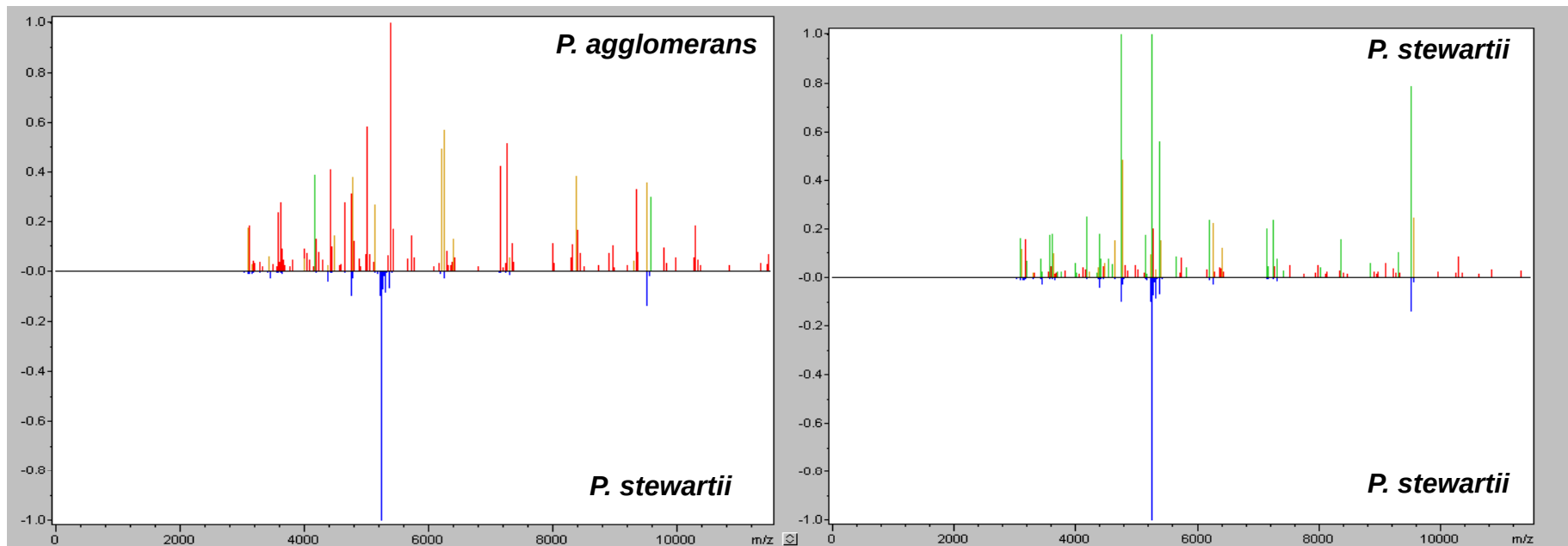
AnagnosTec:

Saramis

Waters:

MicrobeLynx

Reference spectra matching:

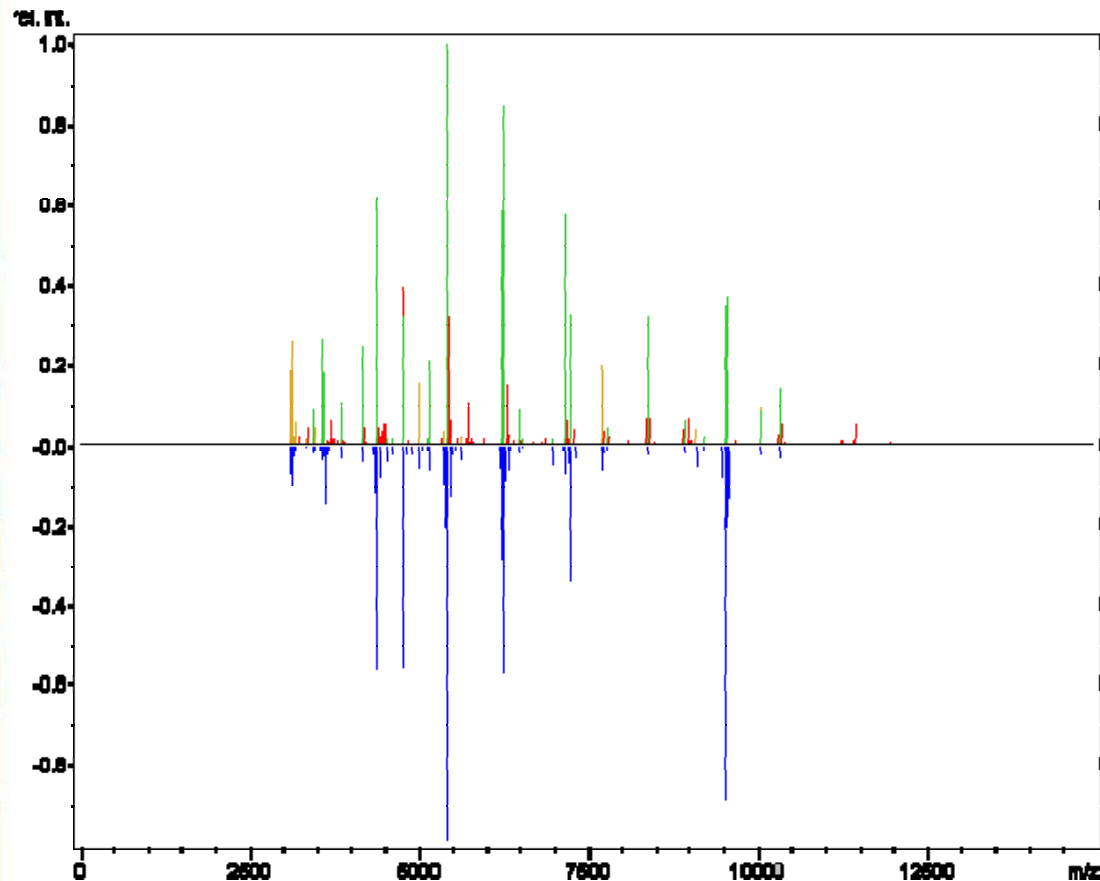


full match = green
partial match = yellow
non-matching = red

Wensing A, Zimmermann S, Geider K (2010)
Appl Environ Microbiol 76: 6248-6256

Reference spectra matching:

Example: German *E. tasmaniensis* isolate



Detected Species	Log(Score)
Erwinia tasmaniensis Et 2_99_PAH	2.200
Erwinia tasmaniensis 1-99 MMG	2.088
Erwinia tasmaniensis Et 1_99_PAH	2.047
Erwinia pyrifoliae 16-96 MMG	1.807
Erwinia amylovora CFBP1232 MMG	1.729
Enterobacter asburiae DSM 17506_DSM	1.672
Erwinia pyrifoliae 1-96 MMG	1.607
Erwinia sp EJP 562_PAH	1.595
Erwinia pyrifoliae DSM 12163 HAM	1.571
Enterobacter cloacae 20105_2 CHB	1.570

Wensing A, Gernold M, Geider K (2012)
J Appl Microbiol 112: 147-158

Isolate identification:

Table 1: Strain identification by Biotyper analysis:

	strain	log score ID
<u><i>E. amylovora</i></u>	CFBP1430	Ea 2.1
	Ea273	Ea 2.0
	Ea1/79	Ea 2.2
	Ea7/74	Ea 2.0
	Tp3	Ea 2.2
	EaRW1/06	Ea 2.2
	Ea775	Ea 2.0
	OR6	Ea 2.1
	Ea4/82	Ea 2.2
	EaSp1767-3	Ea 2.2
	EaSp1951-5	Ea 2.3
<u><i>E. pyrifoliae</i></u>	Ep1/96	Ep 2.0
	Ep2/96	Ep 2.2
	Ejp557	Ep 2.3
<u><i>E. tasmaniensis</i></u>	Et1/99	Et 2.3
	Et2/99	Et 2.2
	Et4/99	Et 2.4
	Et09-1	Et 2.4
	SM88	Et 2.3
	SM882	Et 2.3
	Esa13	Et 2.3
	Esa41	Et 2.3
	FLA03	Et 2.2
<u><i>E. billingiae</i></u>	Eb660	Eb 2.2
	Eb661	Eb 2.0
	Eb1261	Eb 2.0
	Kae2a	Eb 2.0
	Kae9a	Eb 2.3
	RIPFgo2	Eb 2.3

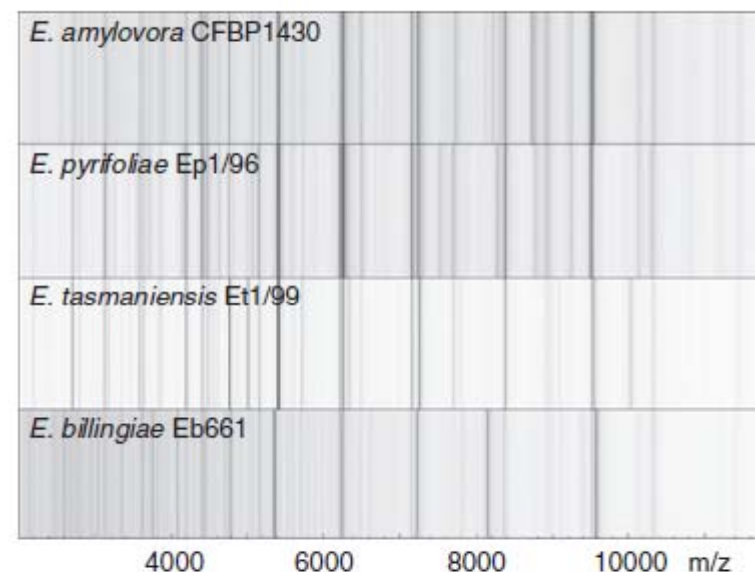


Figure 4 Bar code representation of protein profiles of MALDI-TOF MS spectra. The protein spectra of various *Erwinia* species were visualized in a pseudo-gel representation with peak heights converted to colour intensity.

Wensing A, Gernold M, Geider K (2012)
J Appl Microbiol **112**: 147-158

Method comparison:

**Example fire blight
artificial infected tissue (immature pear):**



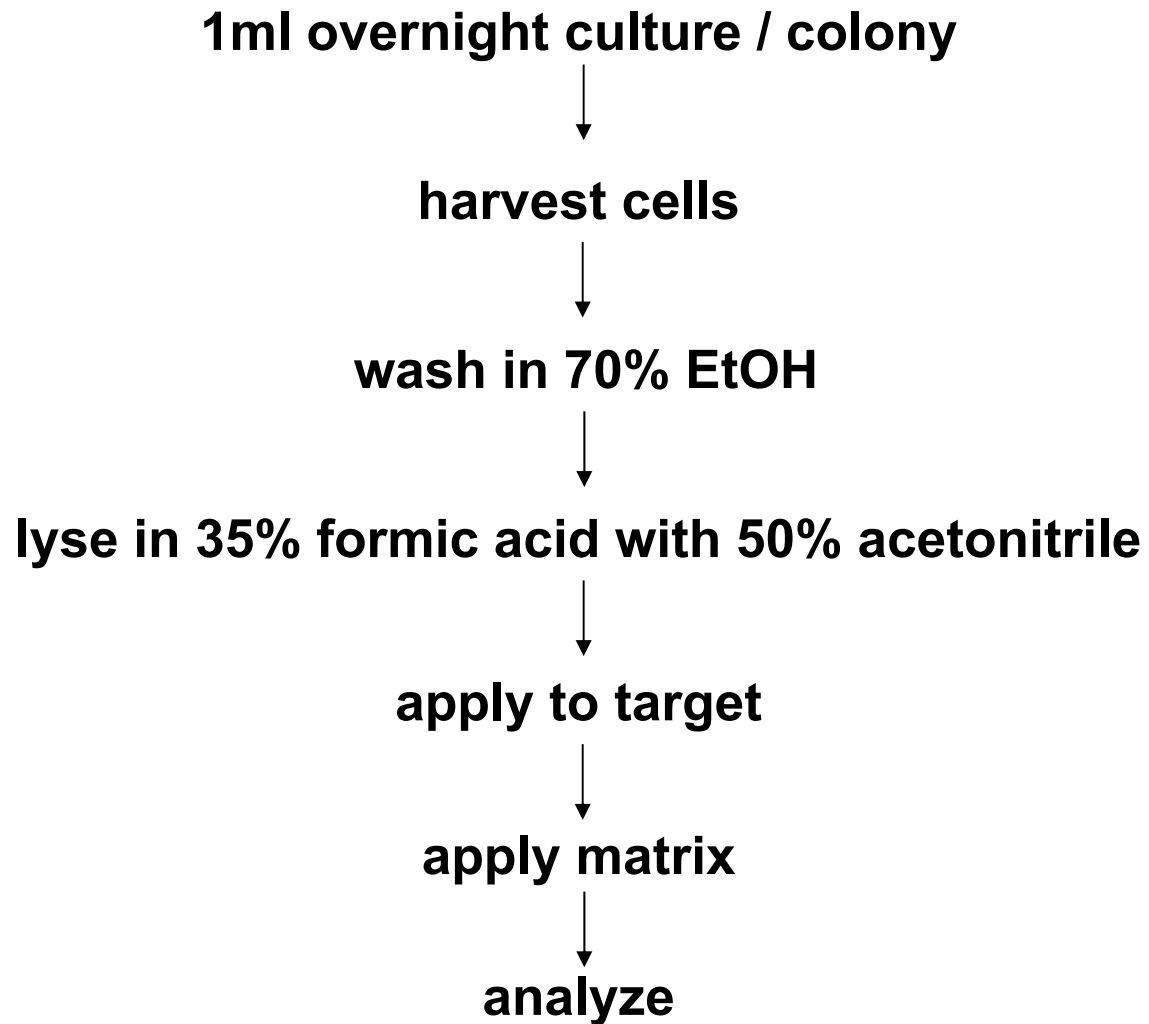
sample with starting symptom formation

Detected Species	Lo...
 <i>Erwinia amylovora</i> CFBP1232 ...	2.020
 <i>Erwinia amylovora</i> 1-79 MMG	1.822
 <i>Erwinia pyrifoliae</i> Ep 16_96_PAH	1.570
 <i>Erwinia tasmaniensis</i> Et 2_99_...	1.512

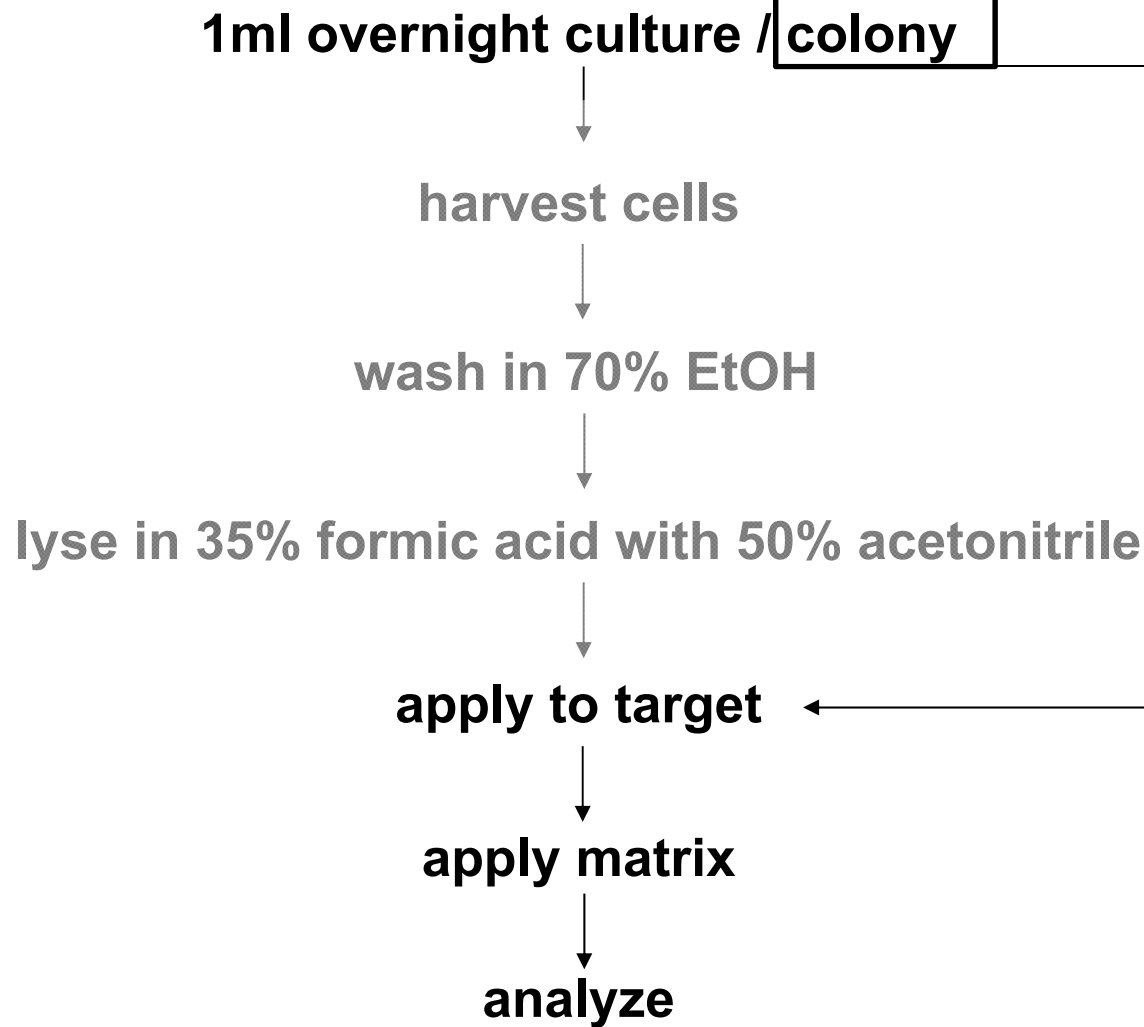
**BUT:
For most samples only mixed spectra,
NOT reliable yet!**

Wensing A, Gernold M, Geider K (2012)
J Appl Microbiol **112**: 147-158

sample preparation:



sample preparation:



MALDI-TOF Mass Spectrometry Is a Fast and Reliable Platform for Identification and Ecological Studies of Species from Family *Rhizobiaceae*

Laura Ferreira¹, Fernando Sánchez-Juanes¹, Paula García-Fraile², Raúl Rivas², Pedro F. Mateos², Eustoquio Martínez-Molina², José Manuel González-Buitrago^{1,3}, Encarna Velázquez^{2*}

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International Biodeterioration & Biodegradation 69 (2012) 82–86



Whole-cell MALDI-TOF: Rapid screening method in environmental microbiology

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APPLIED AND ENVIRONMENTAL MICROBIOLOGY, Oct. 2011, p. 6858–6866
0099-2240/11/\$12.00 doi:10.1128/AEM.05465-11
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Vol. 77, No. 19

Matrix-Assisted Laser Desorption Ionization (MALDI)–Time of Flight Mass Spectrometry- and MALDI Biotyper-Based Identification of Cultured Biphenyl-Metabolizing Bacteria from Contaminated Horseradish Rhizosphere Soil[†]

Ondrej Uhlik,^{1,2} Michal Strejcek,^{1,3} Petra Junkova,¹ Miloslav Sanda,² Miluse Hroudova,² Cestmir Vlcek,² Martina Mackova,^{1,2} and Tomas Macek^{1,2*}

comparison of samples:

	1	2	3	4	5	6	7	8	9	10	11	12
A												
B												
C												
D												
E												
F												
G												
H												

	D1	D2	D3	D4	D5	D6	D7	D8	D9	D10	D11	D12
Matched Pattern	Score	Score	Score	Score	Score	Score	Score	Score	Score	Score	Score	Score
	Value	Value	Value	Value	Value	Value	Value	Value	Value	Value	Value	Value
D1	2.845	0.127	0.828	1.3	1.234	0.098	0.884	0.53	0.825	0.44	0.825	2.606
D2	0.451	2.845	2.13	1.015	1.162	2.17	2.206	2.199	2.435	2.051	2.18	0.123
D3	0.608	2.227	2.845	0.269	0.431	2.601	2.586	2.478	2.192	2.623	2.464	0.485
D4	1.364	1.388	0.733	2.845	2.354	0.838	1.078	0.902	1.19	0.868	0.856	1.436
D5	1.224	1.089	0.712	2.427	2.845	0.395	0.791	0.172	1.205	0.696	0.691	0.97
D6	0.519	2.161	2.498	0.817	0.524	3	2.465	2.444	2.26	2.485	2.392	0.513
D7	0.868	2.31	2.579	0.659	0.79	2.472	2.845	2.566	2.323	2.584	2.486	0.46
D8	0.791	2.264	2.521	0.884	0.637	2.465	2.553	2.877	2.246	2.538	2.497	0.783
D9	0.793	2.334	2.067	1.045	0.917	2.217	2.22	2.147	3	2.196	2.214	0.732
D10	0.7	2.16	2.666	0.534	0.638	2.589	2.627	2.593	2.211	2.845	2.461	0.698
D11	0.781	2.341	2.523	0.616	0.509	2.399	2.56	2.553	2.336	2.443	2.845	0.076
D12	2.66	0.463	0.716	1.426	1.209	0	0.488	0.324	0.501	0	0.893	2.845



Sampling technique:

Sample Size?



vs.



Sampling technique:

Timeframe?



Protocol development:

- sampling procedure
- culture conditions for selection of isolates

Comparison of systems

Bruker:	Biotyper
AnagnosTec:	Saramis
Waters:	MicrobeLynx

Evaluation with established methods

- specific PCR +qPCR for selected species
- sequencing approaches

Output:

- Protocol development for application of a new high-throughput technique for microbial identification to fire blight settings
- Evaluation of systems, sampling and culturing conditions
- Information on development of *E. amylovora* in the context of a competing microflora
- Information on effect of fire blight treatment on the natural microflora
- Information on development of microbial biocontrol organisms (biosafety)

A close-up photograph of a flowering branch, likely from a fruit tree, showing several pink buds and green leaves. The branch is dark brown and textured. The background is a blurred outdoor setting with a wooden fence and greenery. A semi-transparent rectangular box with a light gray background is centered over the image, containing the text "Thank you!" in a bold, black, sans-serif font.

Thank you!