

Characterization of *Pseudomonas syringae* pv. *actinidiae* genetic diversity in Portugal

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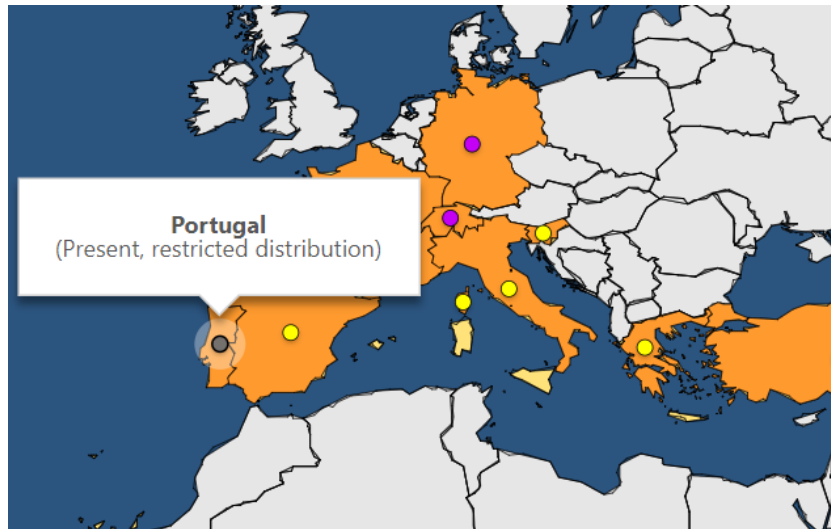
IX International Symposium on Kiwifruit

Porto, 8 September 2017





First report: 2010



Previous studies revealed that Psa isolates analysed from Entre Douro e Minho region were identical to the **biovar 3** (Renzi *et al.*, 2012; Moura *et al.*, 2015)

Portuguese Psa population from Entre Douro e Minho region presents **some degree of genetic variability** (Moura *et al.*, 2015).

New Disease Reports (2010) 22, 10. [doi:10.5197/j.2044-0588.2010.022.010]



New Disease Reports

First report of bacterial canker of *Actinidia deliciosa* caused by *Pseudomonas syringae* pv. *actinidiae* in Portugal

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Identificação e caracterização de *Pseudomonas syringae* pv. *actinidiae* (Psa) na Região do Entre Douro e Minho (Portugal)

Identification and characterization of *Pseudomonas syringae* pv. *actinidiae* (Psa) in Entre Douro e Minho region (Portugal)

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Recibido/Received: 2013-03-05
Aceptado/Accepted: 2013-05-20

Phytopathologia Mediterranea (2012) 51, 2, 402–409

Short Notes

Widespread distribution of kiwifruit bacterial canker caused by the European *Pseudomonas syringae* pv. *actinidiae* genotype in the main production areas of Portugal

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S03.04

Bacterial canker of kiwi caused by *Pseudomonas syringae* pv. *actinidiae* in Portugal – Disease Importance and pathogen characterization

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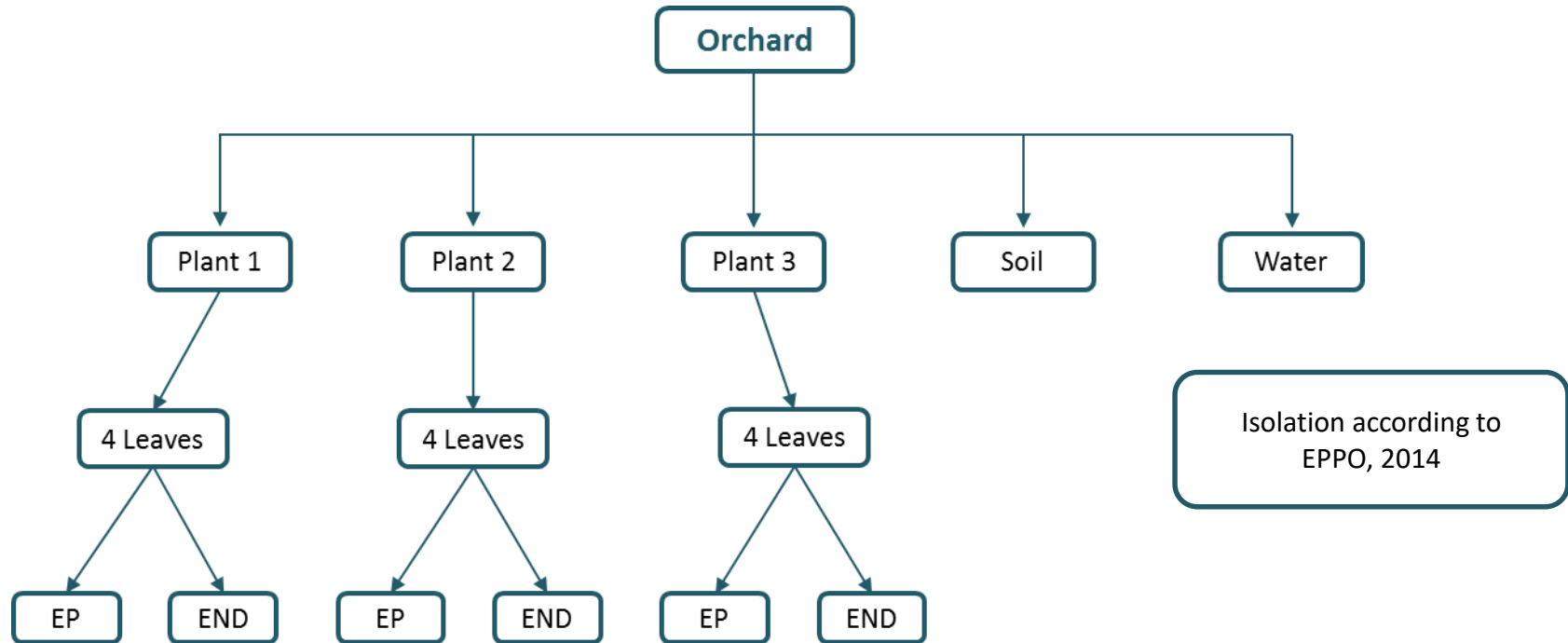
Objective: Analyse the diversity of Psa populations in Portugal.

Orchard	Localization	Cultivar	Age (years)	First detection of Psa	Psa disease severity degree
A	Viana do Castelo	<i>A. deliciosa</i> cv. "Hayward"	7	2010	1
B	Guimarães	<i>A. deliciosa</i> cv. "Erika"	5	2015	2
C	Albergaria-a-Nova	<i>A. deliciosa</i> cv. "Hayward"	16	2016*	1
D	Montemor-o-Velho	<i>A. deliciosa</i> cv. "Hayward"	4	2015	3
E	Montemor-o-Velho	<i>A. deliciosa</i> cv. "Hayward"	30	2016	2

*This study

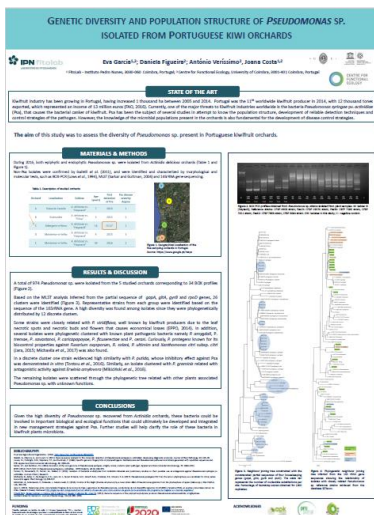
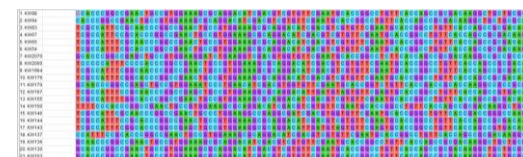
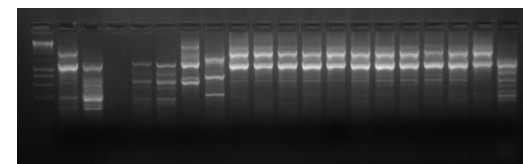
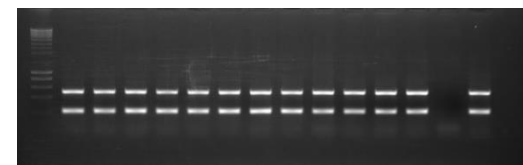
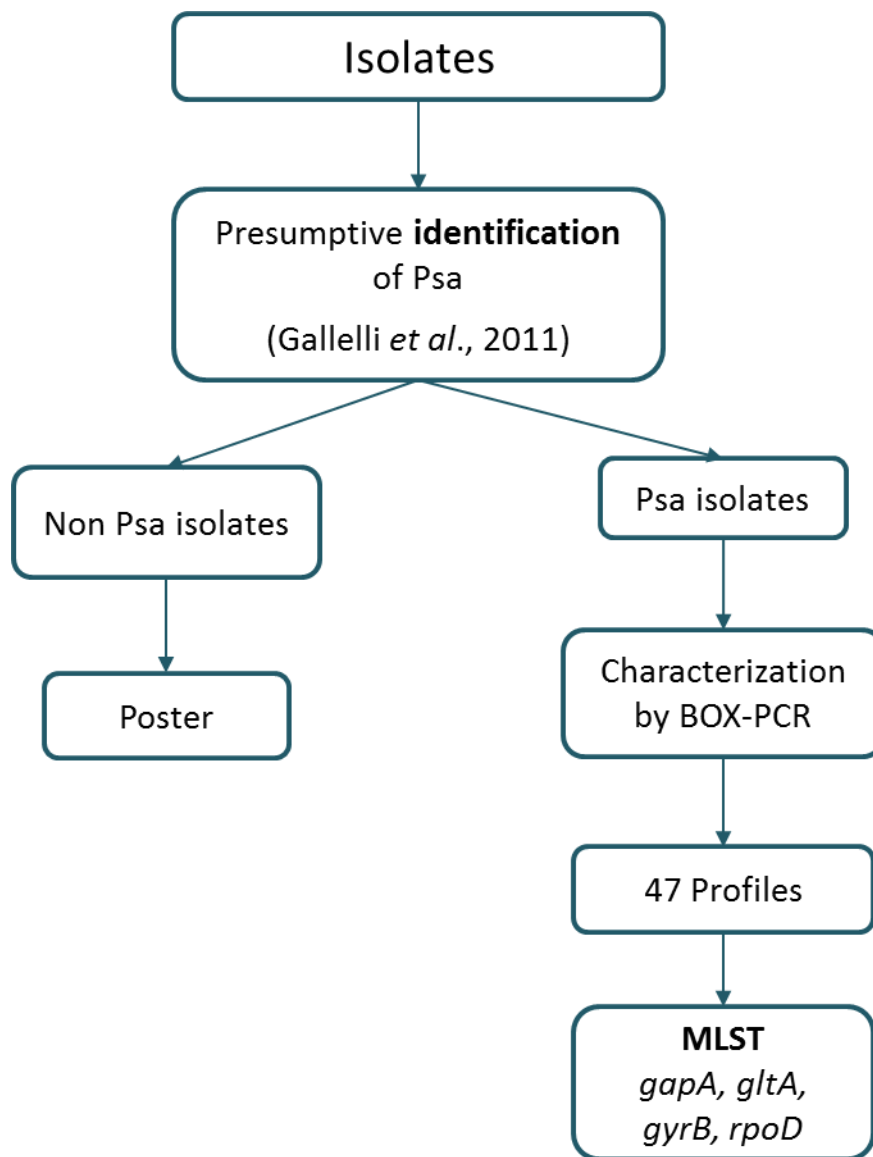
Spring and Autumn
2016





Spring and Autumn

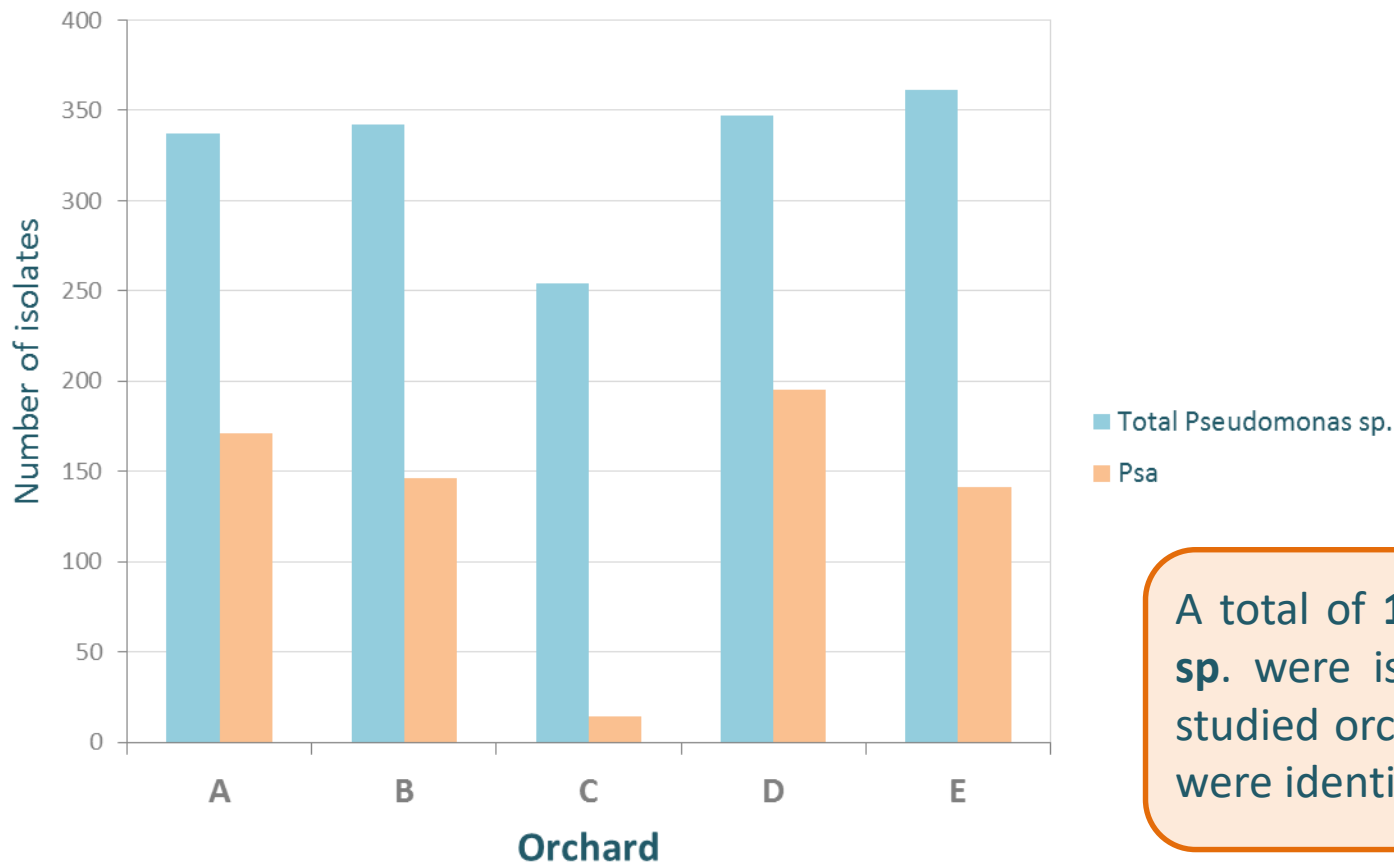






Objective 1

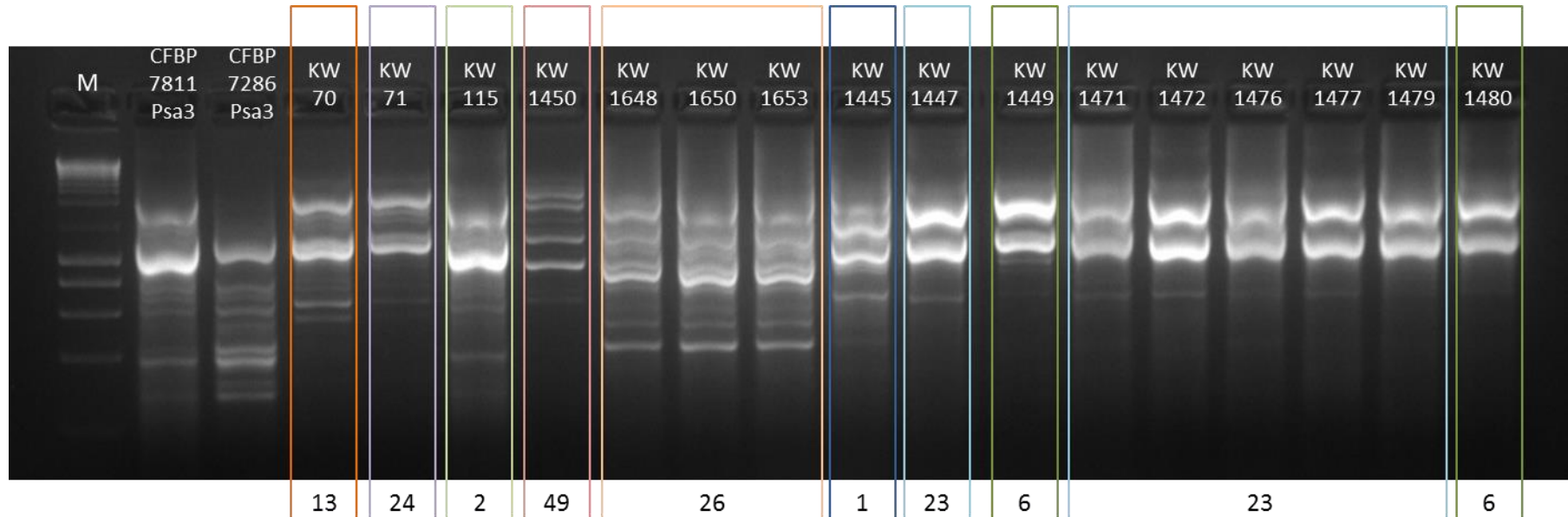
Isolate the maximum number of *Pseudomonas* sp. from kiwifruit orchards.



A total of **1641 *Pseudomonas* sp.** were isolated from the 5 studied orchards of which **667** were identified as ***Psa***.

Objective 2

Determine the **diversity of Psa** populations.

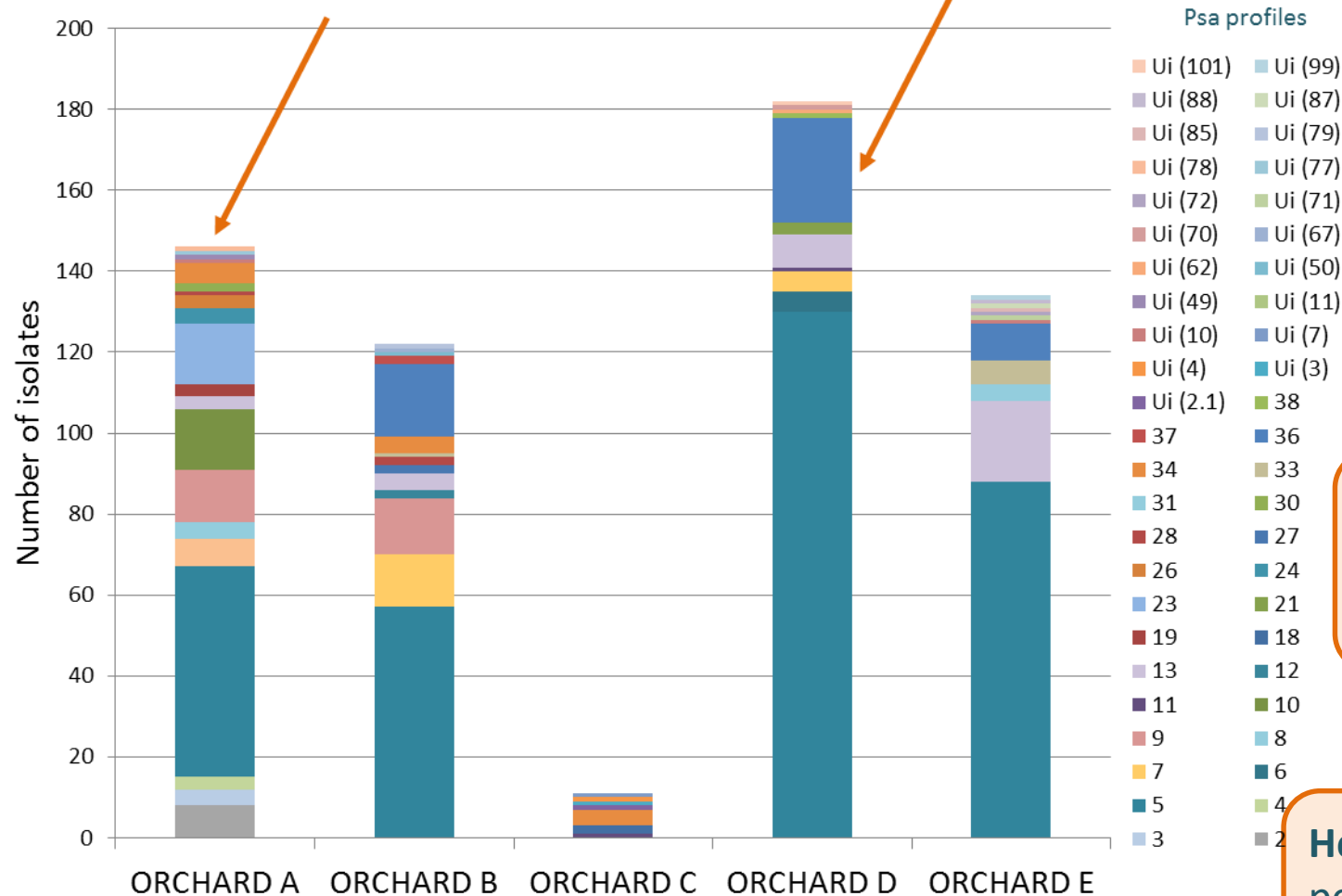


Profiles	Nº of isolates	Profiles	Nº of isolates	Profiles	Nº of isolates	Profiles	Nº of isolates	Profiles	Nº of isolates	Profiles	Nº of isolates
2	8	10	15	24	4	36	53	Ui (49)	1	Ui (78)	1
3	4	11	2	26	3	37	2	Ui (50)	1	Ui (79)	1
4	3	12	2	27	2	38	1	Ui (62)	1	Ui (85)	1
5	327	13	35	28	3	Ui (2.1)	1	Ui (67)	1	Ui (87)	1
6	12	18	2	30	2	Ui (3)	1	Ui (70)	1	Ui (88)	1
7	18	19	3	31	4	Ui (4)	1	Ui (71)	1	Ui (99)	1
8	4	21	3	33	7	Ui (7)	1	Ui (72)	1	Ui (101)	1
9	27	23	15	34	13	Ui (10)	2	Ui (77)	1		

A total of **47** BOX profiles were identified.

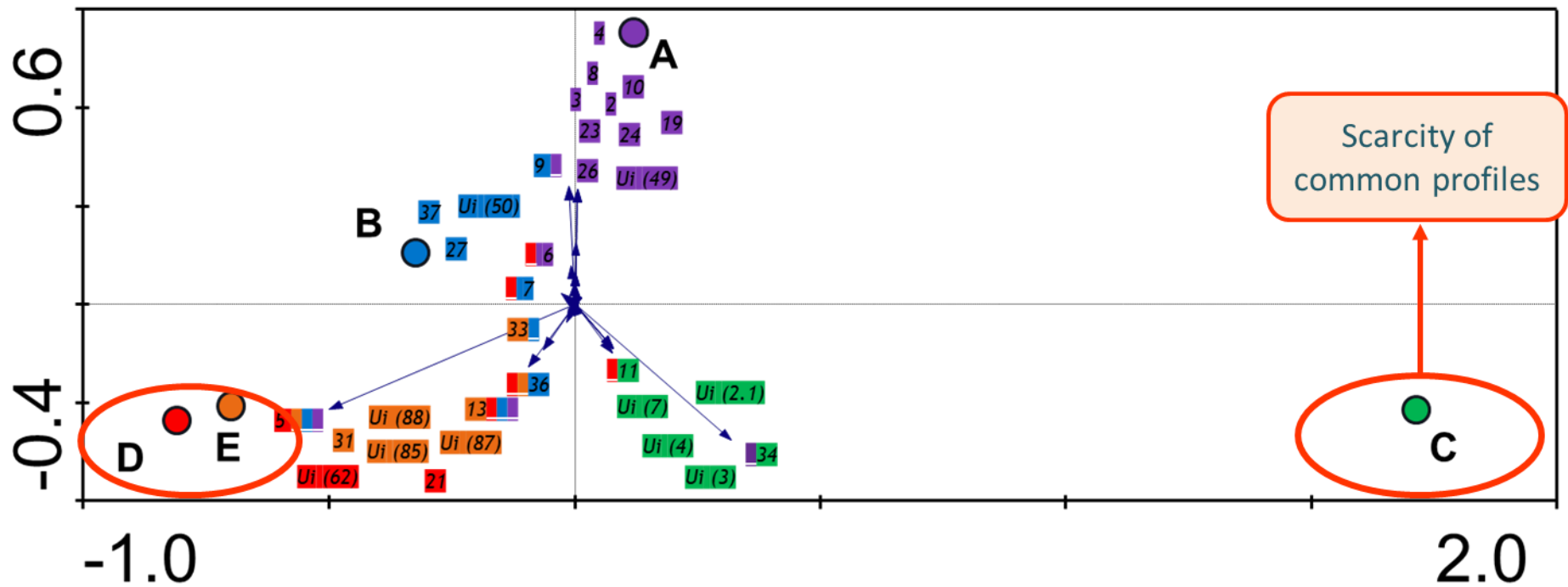
Objective 3

Determine the **diversity** of Psa populations **between orchards**.



Not a single profile was common to all orchards.

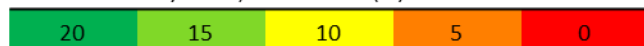
Heterogeneity in Psa population was found between orchards.



Principal component analysis – inter-species correlate – of Psa profiles from each studied orchard. Purple A: total plant isolates from orchard A; Blue B: total plant isolates from orchard B; Green C: total plant isolates from orchard C; Red D: total plant isolates from orchard D; Orange E: total plant isolates from orchard E. Numbers correspond to Psa profiles. Colours identify Psa profiles that have more weight in each condition.

	A	B	C	D	E
A					
B	12.50				
C	3.33	0.00			
D	5.26	17.39	7.14		
E	4.35	12.90	0.00	18.75	

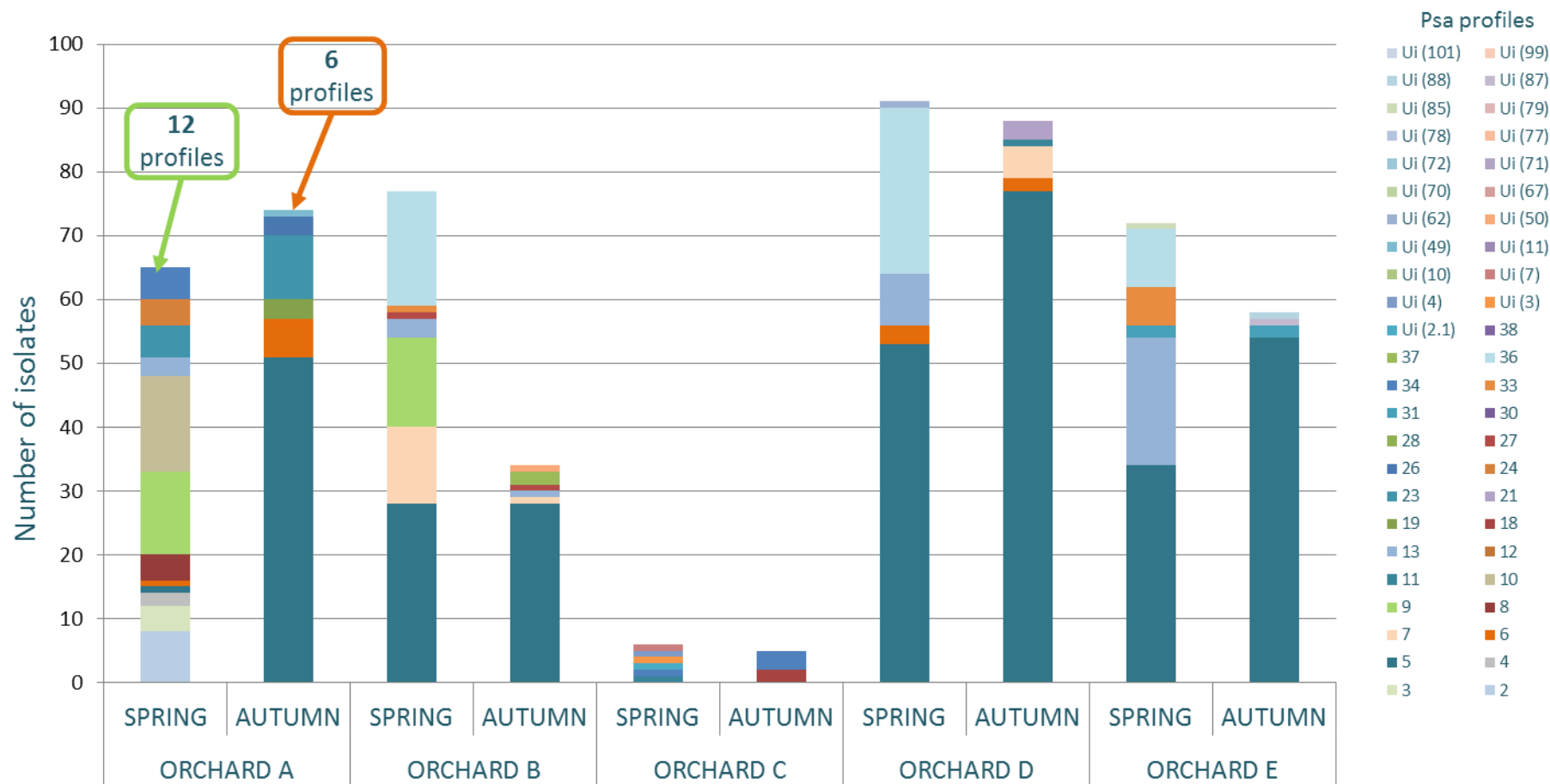
Jaccard similarity index, colour scale (%)



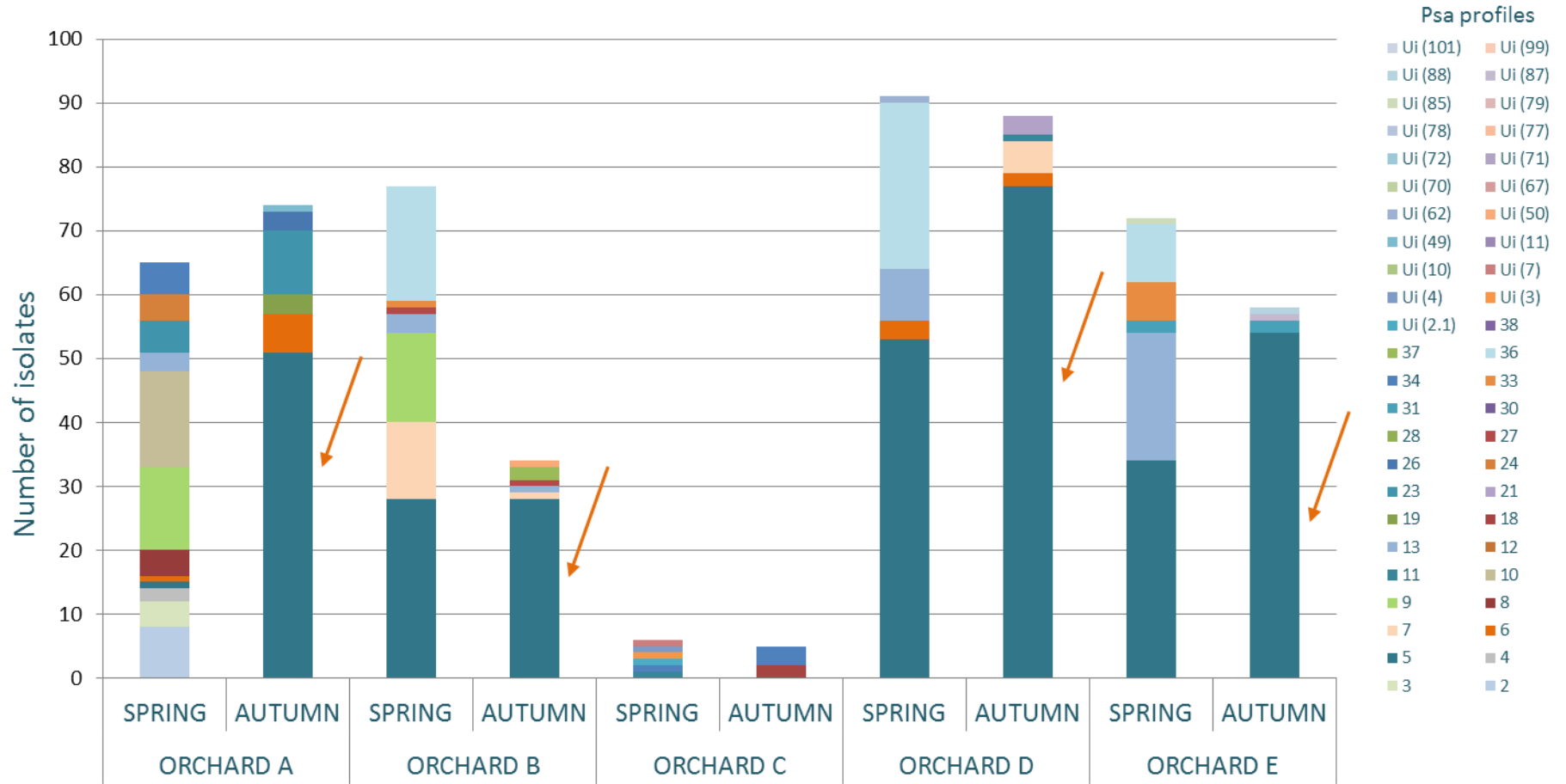
Beta diversity matrix, determined by Jaccard similarity index for the five studied orchards (A, B, C, D and E).

Objective 4

Determine the **diversity** of Psa populations **over time**.



The **diversity** among Psa populations determined in **spring** was **higher** when compared to those determined in autumn.

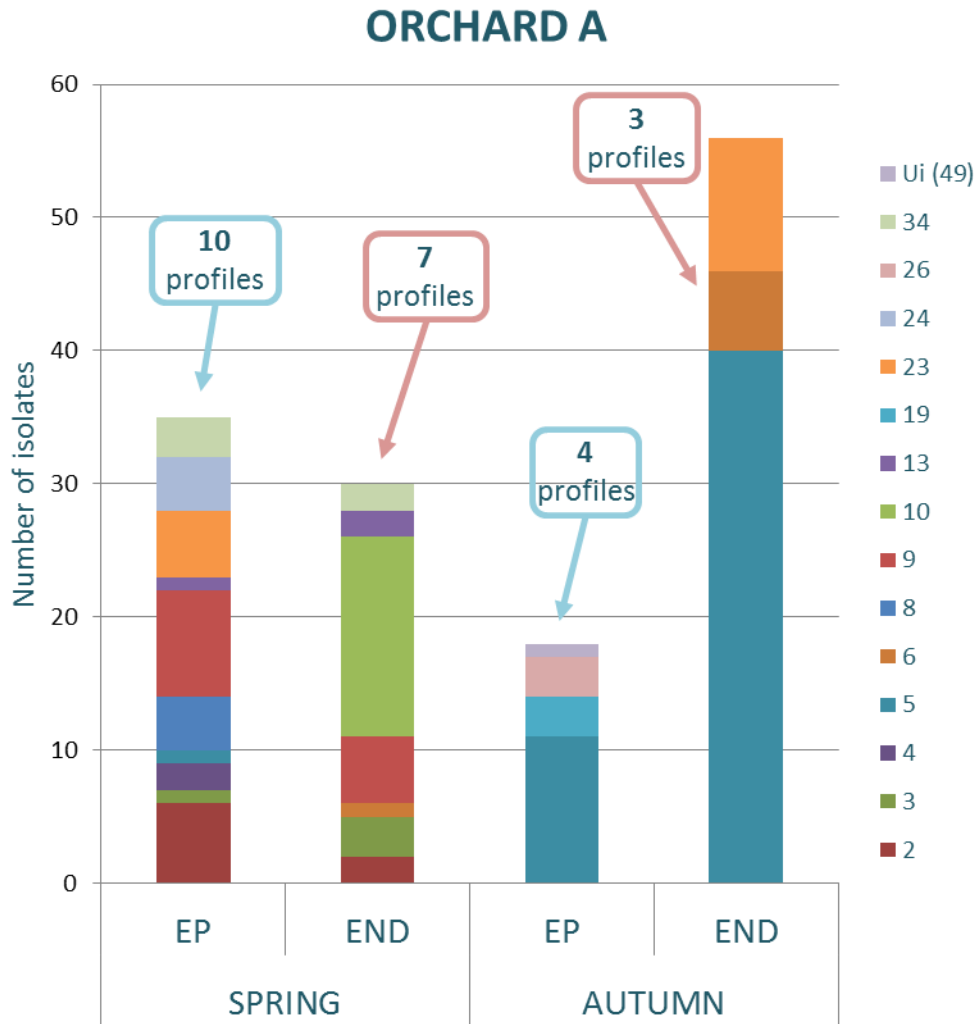


Dominant population
in Autumn:
Psa profile 5

Psa profile 5 could be more
resilient or better adapted to
summer conditions.

Objective 5

Diversity of **endophytic** and **epiphytic** Psa strains over time.



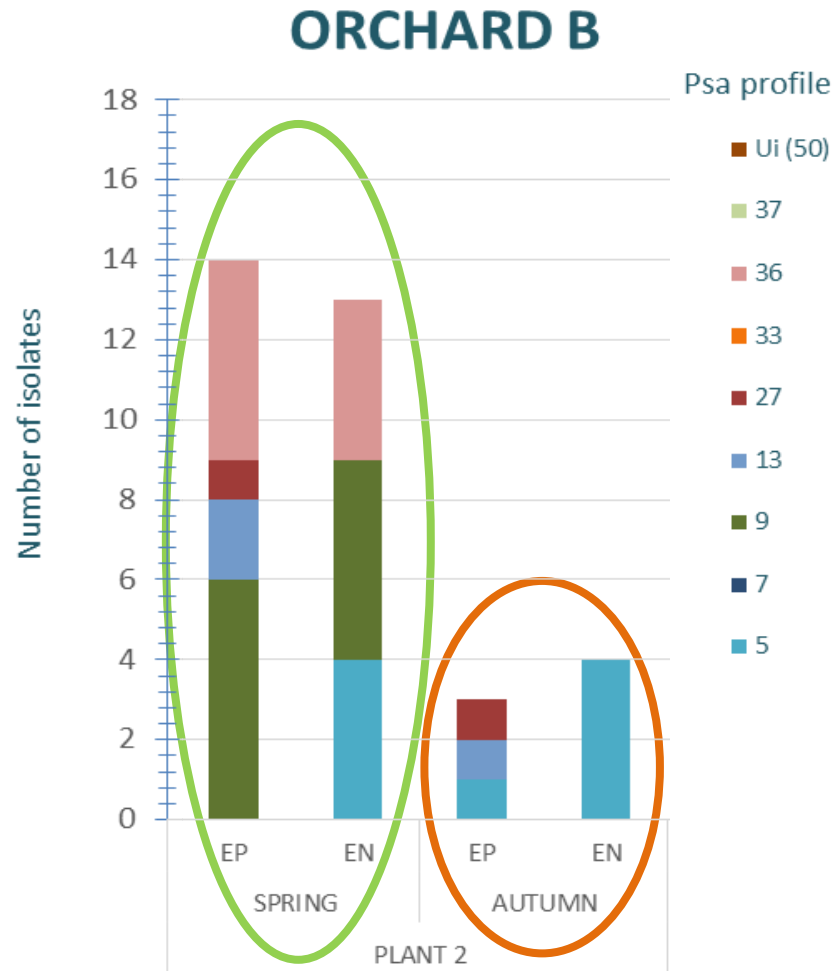
Psa population **heterogeneity** was found **between endophytic and epiphytic** isolates.

Psa population varied **over time**.

Higher diversity was found among **epiphytic** Psa populations when compared to endophytic Psa populations.

Objective 6

Diversity of **endophytic** and **epiphytic** Psa strains in the **same plant**, **over time**.



Structure of Psa populations **varied over time** in the same plant.

Objective 7

Psa diversity between **North** and **Centre** region.

Index	North		Centre	
	Orchard A	Orchard B	Orchard D	Orchard E
Dmg	3,40	2,33	2,07	2,60
H'	1,43	1,13	0,77	1,11
J'	0,77	0,73	0,57	0,69
1-D	0,67	0,60	0,40	0,56

Higher diversity

Dominant profiles

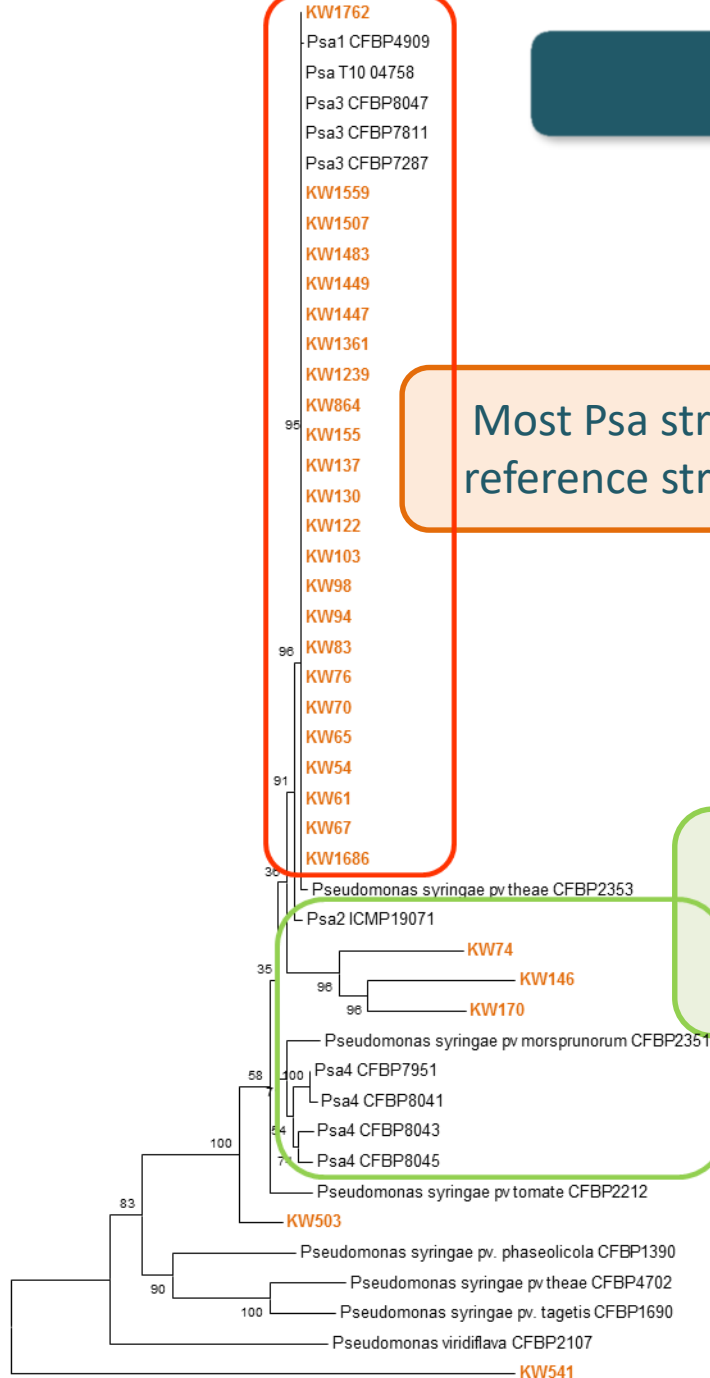
Orchards from the **North** region presented **higher Psa diversity** than the orchards from the Centre region.

Presence of **dominant profiles** was more evident in **Centre** region.

Phylogenetic analysis

Most Psa strains were clustered with reference strains from Psa1 and Psa3.

3 isolates were closely related with Psa2 and Psa4 (*P. syringae* pv. *actinidifoliorum*).



Neighbour joining tree constructed with the concatenated partial sequences of four housekeeping genes (*gapA*, *gltA*, *gyrB* and *rpoD*).



- Psa populations present in Portuguese orchards were **heterogeneous**.
- This **heterogeneity** was found **within orchards** and **between orchards**.
- The **structure** of Psa populations **varied over time** in the same plant.
- The **diversity** among Psa populations determined in **spring** was **higher** when compared to autumn.
- The **decrease in Psa diversity** in autumn was accompanied by an **increase in dominant populations**, such as Psa profile 5, suggesting that this clone could be more resilient or better adapted to summer conditions.
- Orchards from the **Centre region** presented **less Psa diversity** due to the presence of dominant profiles.



ACKNOWLEDGES

Laboratório de Fitossanidade



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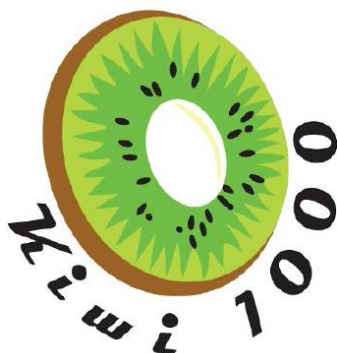


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inscrita na Lista do Património
Mundial em 2013



IPN fitolab

LABORATÓRIO DE FITOSSANIDADE



Trabalho realizado no âmbito da Ação 1.1 Grupos Operacionais “I9K – InovKiwi - Desenvolvimento de estratégias que visem a sustentabilidade da fileira do kiwi através da criação de um produto de valor acrescentado” promovida pelo PDR2020 e co-financiada pelo FEADER, no âmbito do Portugal 2020.



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