



Transmission dynamics of European stone fruit yellows on thirteen *Prunus* species in controlled conditions

P. Ermacora, L. Carraro, F. Ferrini, M. Martini, N. Loi

Dipartimento di Biologia e Protezione delle Piante,
University of Udine, Via delle Scienze 208, 33100 Udine,
Italy

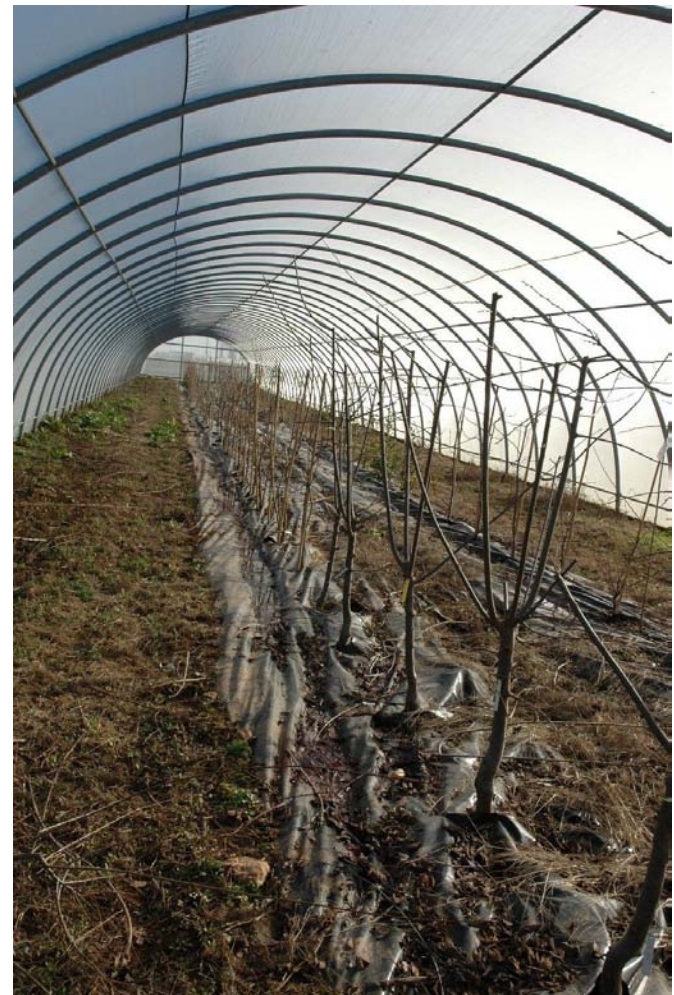
Aim of the study

Improve knowledges about susceptibility and sensitivity of several *Prunus* species to ESFY infection, by using the vector *Cacopsylla pruni* in semi-controlled conditions and analyzing their host-feeding preference

Materials and Methods



130 test plants belonging to 13 different *Prunus* species



Materials and Methods

Prunus amygdalus (Almond)

P. armeniaca (Apricot)

P. avium (Sweet Cherry)

P. cerasifera (Myrobalan Plum)

P. cerasus (Sour cherry)

P. domestica (European plum)

P. mahleb (Mahaleb cherry)

P. laurocerasus (Cherry laurel)

P. padus (Bird cherry)

P. persica (Peach)

P. salicina (Japanese plum)

P. spinosa (Blackthorn)

P. tomentosa (Nanking Cherry)

Materials and Methods

***C. Pruni* capture**

- Reimmigrant *C. pruni* were collected in spring 2007 and 2008 in a plum orchard located in a high ESFY disease pressure area in Friuli - Venezia Giulia Region (Italy).
- Samples of insects were submitted to nested PCR
- Insects were immediately released in the screenhouse.

Materials and Methods

'*Candidatus* Phytoplasma prunorum' detection

Sampling: 1g of leaf midribs collected in September

DNA extraction according Doyle & Doyle protocol

Direct PCR: P1/P7 primers

Nested PCR: f01/r01 primers

Agarose gel electrophoresis and ethidium bromide staining

RFLP with *Bsa* I, *Ssp*I

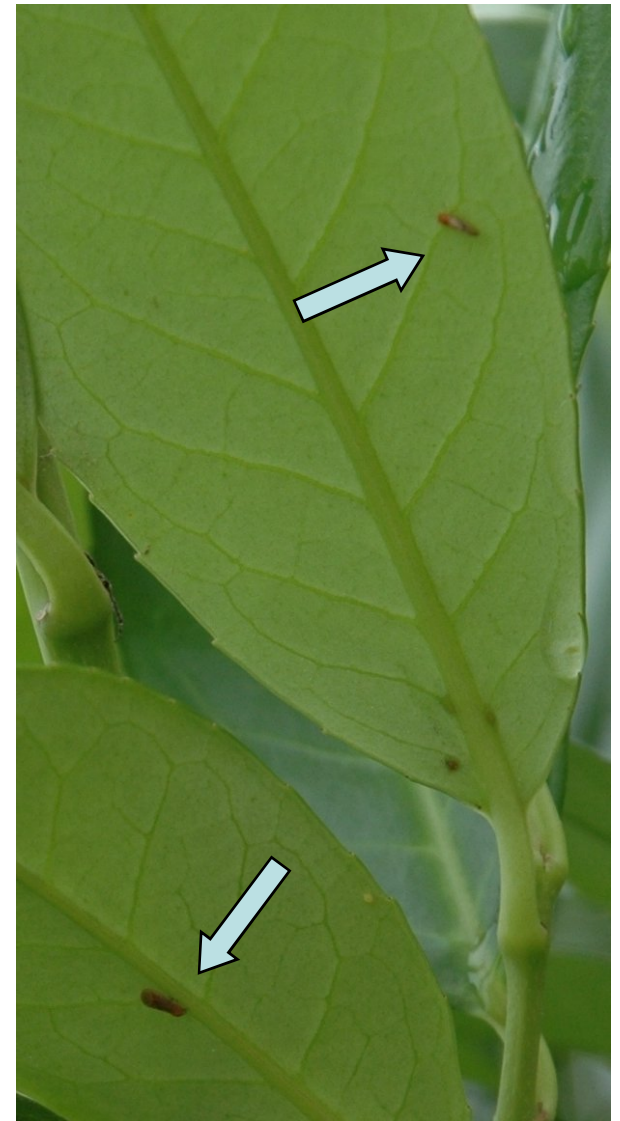
Results

***C. Pruni* capture and infectivity**

- 2400 reimmigrant adults of *C. pruni* captured and released in 2007 and 4700 in 2008.
- In 2007, 16% of insect samples tested positive in PCR
- In 2008, 21% of insect samples tested positive in PCR

Results

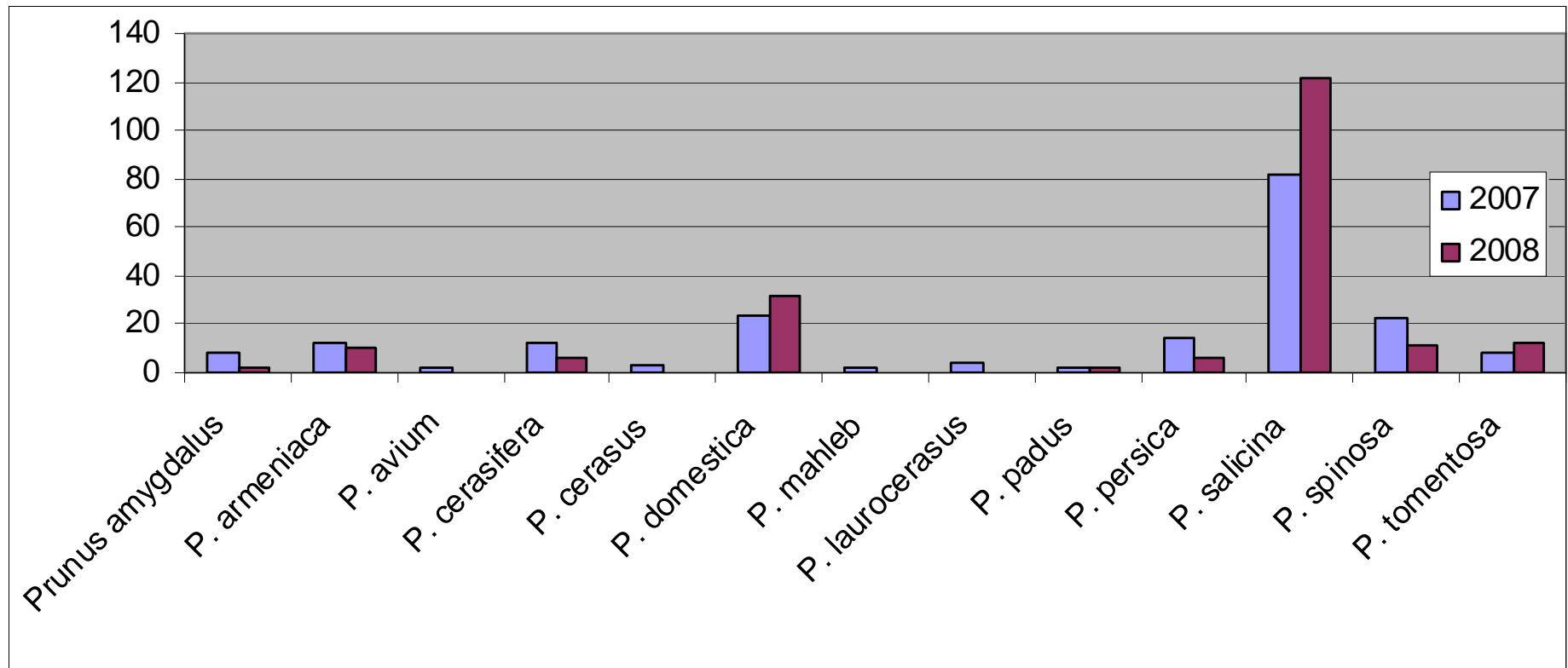
Reimmigrant *C. pruni* distribution on *Prunus* spp.



Results

***Reimmigrant C. pruni* distribution**

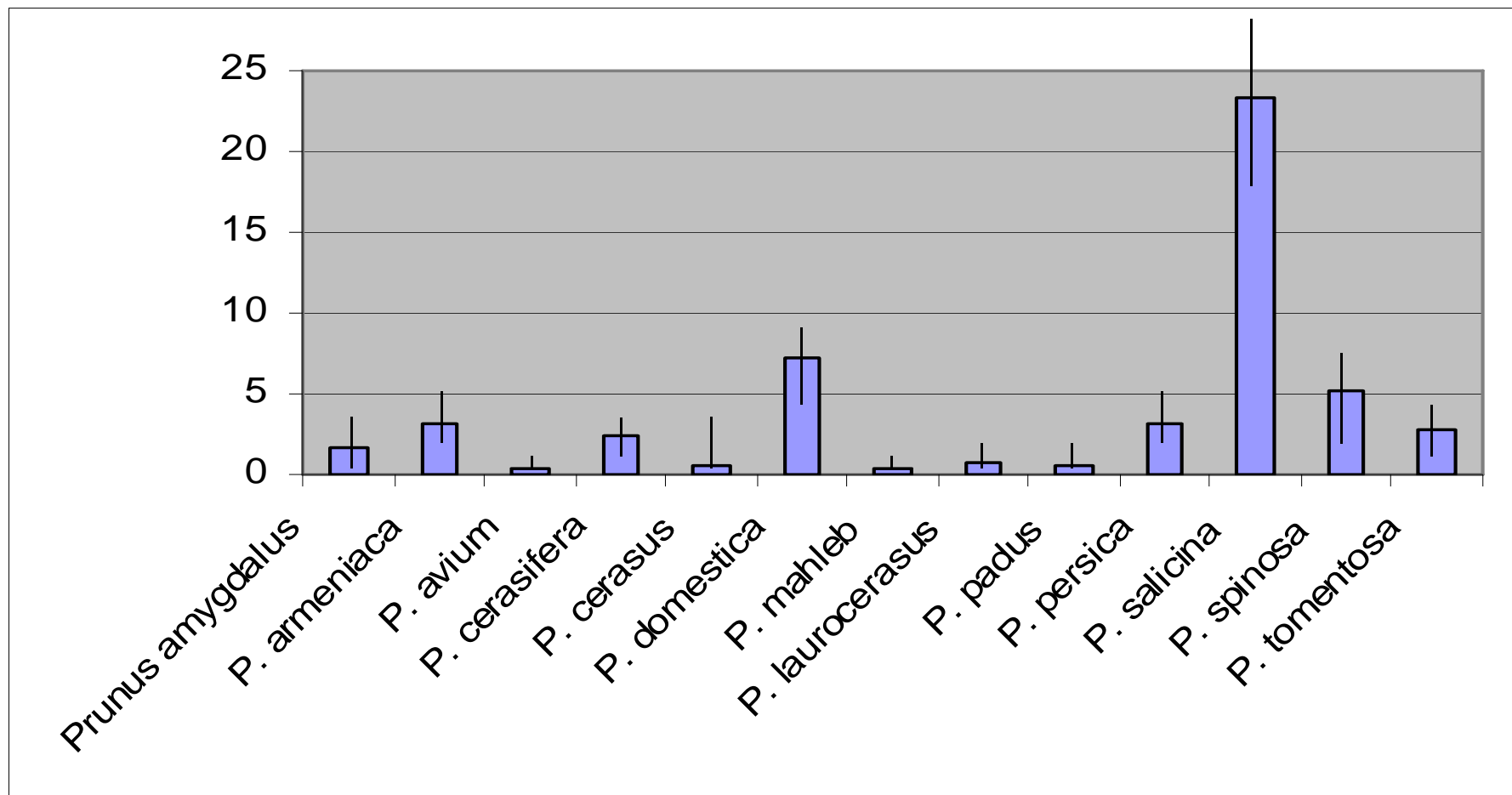
Total number of visual detected *C. pruni* on the 10 plants of each *Prunus* spp. in 2007 and 2008



Results

***Reimmigrant C. pruni* distribution**

Mean number of *C. pruni*/plant 2007/2008



Results

Disease progression

	Number of positive plants by using PCR		
	September 2007	September 2008	September 2009
<i>Prunus amygdalus</i>	0	2	0
<i>P. armeniaca</i>	2	5	4
<i>P. avium</i>	0	0	0
<i>P. cerasifera</i>	1	2	2
<i>P. cerasus</i>	0	0	0
<i>P. domestica</i>	0	1	0
<i>P. mahleb</i>	0	1	0
<i>P. laurocerasus</i>	0	0	0
<i>P. padus</i>	0	0	0
<i>P. persica</i>	0	0	3
<i>P. salicina</i>	5	10	10
<i>P. spinosa</i>	0	0	0
<i>P. tomentosa</i>	0	3	3

Conclusions

Prunus salicina is the most attractive species for *C.pruni*

P. salicina: 100% of plants are infected

P. armeniaca: 40% of plants are infected

Work in progress

Experiments by using olfactometers and fly-tunnels

Chemicals involved in host-vector interaction

Behaviour of the infected plants to a re-infection