

Online resource 1:

Vector transmission and epidemiology of ‘*Candidatus Phytoplasma pyri*’ in Austria

Journal of Plant Diseases and Protection

Monika Riedle-Bauer¹, Caroline Paleskić^{1,2}, Christina Schönhuber², Martina Staples¹, Günter Brader^{2*}

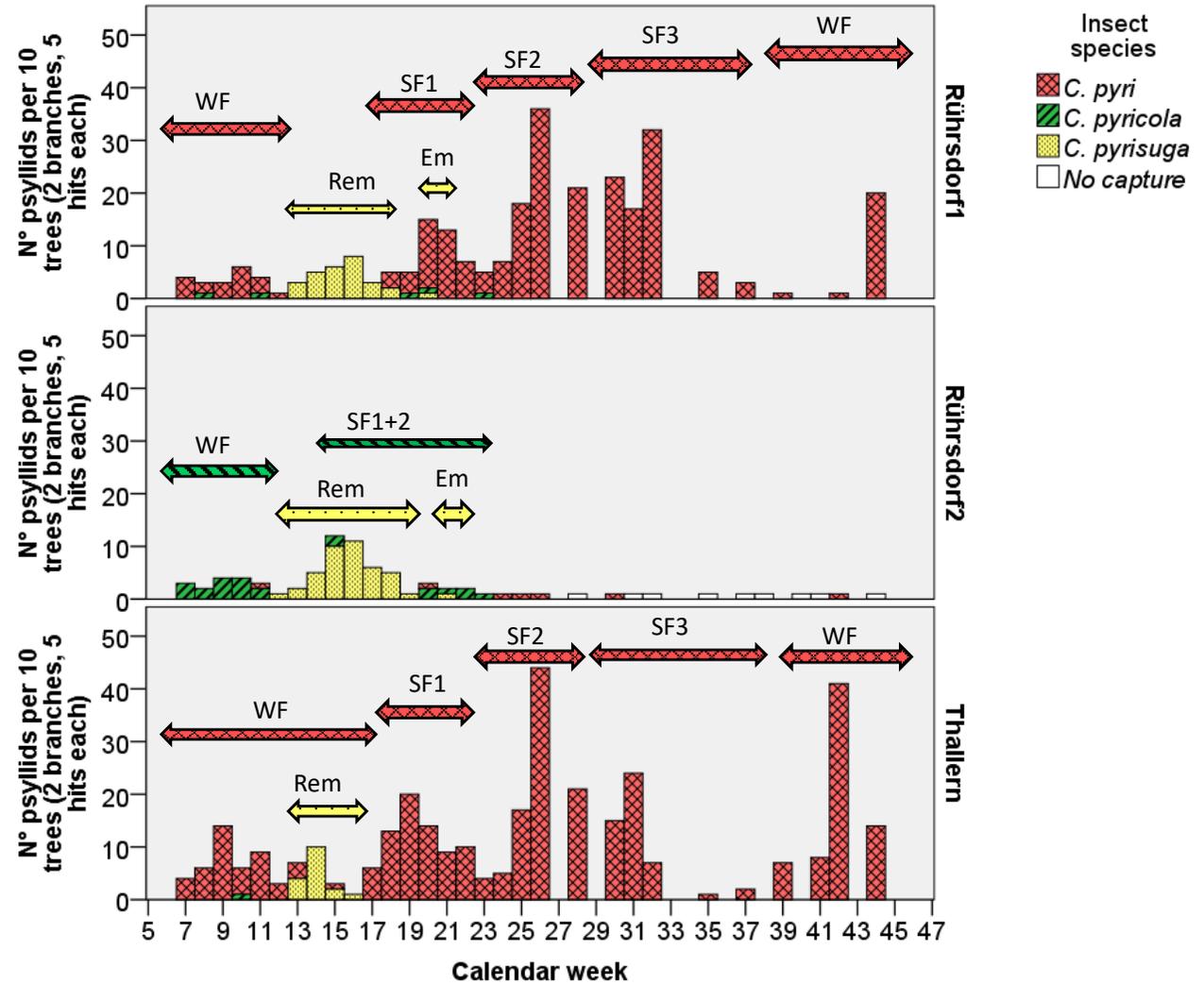
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Captures of pear psyllids at the locations Rührsdorf 1, Rührsdorf 2 and Thallern, Lower Austria, in the course of the year 2014. Numbers represent total captures of insects on 10 trees (2 branches with 5 hits per tree).

WF-winterform individuals, SF1,2,3, Summerform individuals presumably 1st, 2nd, 3rd generation, Rem-remigrants, Em-Emigrants.



Online resource 2:

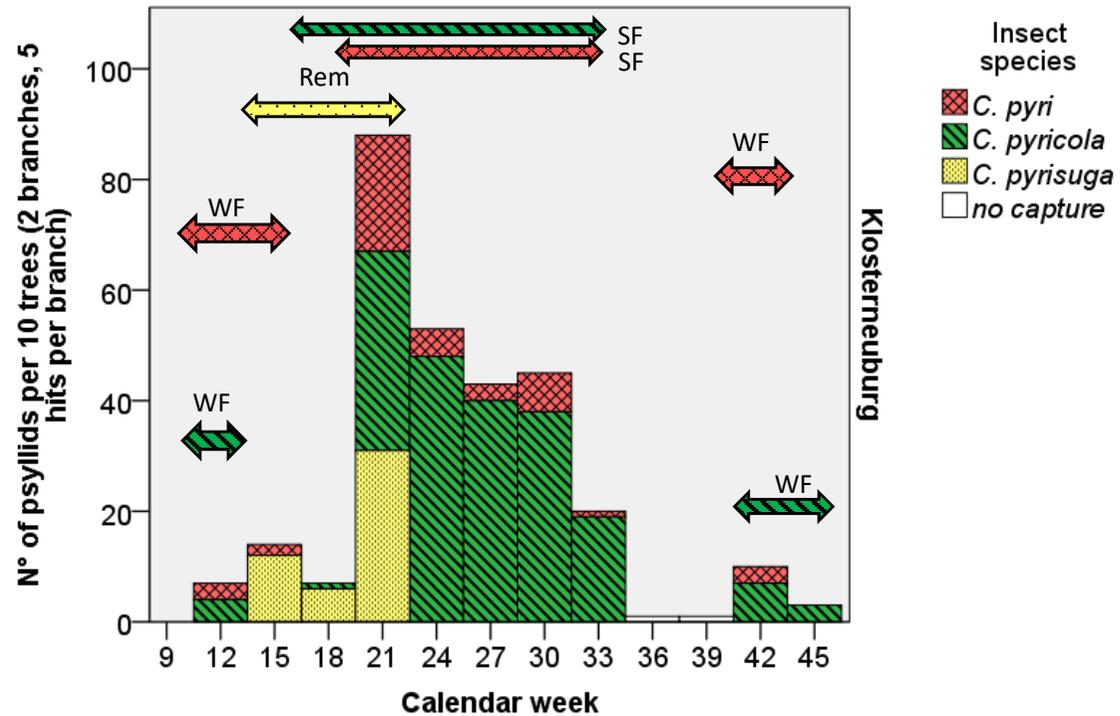
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Captures of pear psyllids in Klosterneuburg in 2012. Numbers represent total captures of insects on 10 trees (2 branches with 5 hits each per tree).



Online resource 4:

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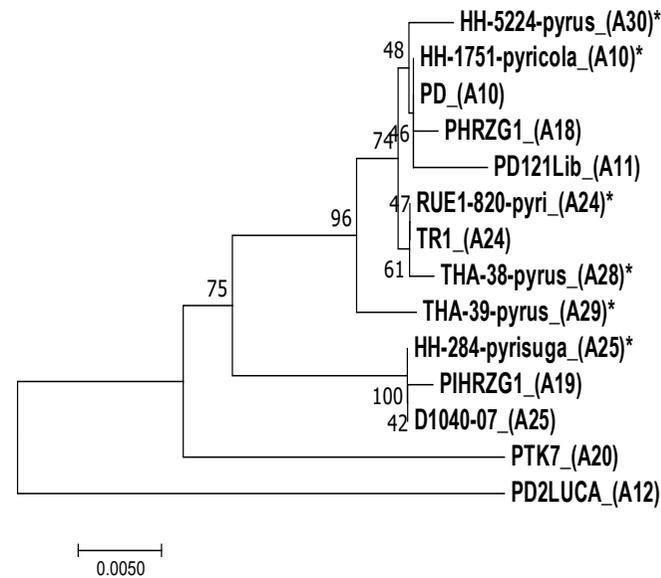
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Phylogenetic relationship of identified *aceF* (a) and *imp*
(b) strains compared to strains of NCBI database. The
optimal tree with the sum of branch length = 0.0988 (a)
and 0.3276 (b) are shown. The percentage of replicate
trees in which the associated taxa clustered together in
the bootstrap test (1000 replicates) are shown next to
the branches. Strains identified in this study are marked
with an asterisk. In brackets after the strain name the A
(*aceF*) and *imp* (I) nomenclature of Danet et al. (2011)
and, if applicable, the A-K nomenclature of Bohunicka
et al. (2018) is indicated.

a



b

