AperTO - Archivio Istituzionale Open Access dell'Università di Torino

Flavescence dorée phytoplasma titre in field-infected Barbera and Nebbiolo grapevines

This is the author's manuscript

Original Citation:

Availability:

This version is available <http://hdl.handle.net/2318/152809> since 2017-05-28T20:51:58Z

Published version:

DOI:10.1111/ppa.12068

Terms of use:

Open Access

Anyone can freely access the full text of works made available as "Open Access". Works made available under a Creative Commons license can be used according to the terms and conditions of said license. Use of all other works requires consent of the right holder (author or publisher) if not exempted from copyright protection by the applicable law.

(Article begins on next page)

UNIVERSITÀ DEGLI STUDI DI TORINO

This is the accepted version of the following article: [5. Roggia C., Caciagli P., Galetto L., Pacifico D., Veratti F., Bosco D. and Marzachì C. 2014. Flavescence dorée phytoplasma titre in field-infected Barbera and Nebbiolo grapevines. Plant Pathology 63: 31-41],

which has been published in final form at [10.1111/ppa.12068]

**Flavescence dorée phytoplasma titre in field-infected
Barbera and Nebbiolo grapevines**

Journal:	<i>Plant Pathology</i>
Manuscript ID:	PP-13-018.R1
Manuscript Type:	Original Article
Date Submitted by the Author:	n/a
Complete List of Authors:	Roggia, Chiara; Istituto di Virologia Vegetale, CNR, Caciagli, Piero; Istituto di Virologia Vegetale, CNR, Galetto, Luciana; Istituto di Virologia Vegetale, CNR, Pacifico, Davide; Istituto di Virologia Vegetale, CNR, Veratti, Flavio; Istituto di Virologia Vegetale, CNR, Bosco, Domenico; University of Torino, Di.Va.P.R.A., Entomologia e Zoologia Applicate all'Ambiente, Marzachi, Cristina; Istituto di Virologia Vegetale, CNR,
Topics:	epidemiology
Organisms:	phytoplasmas & relatives
Other Keywords:	

SCHOLARONE™
Manuscripts

FDP-titre in grapevines with different susceptibility to GYs

Flavescence dorée phytoplasma titre in field-infected Barbera and Nebbiolo grapevines

C. Roggia^{1,2}, P. Caciagli¹, L. Galetto¹, D. Pacifico¹, F. Veratti¹, D. Bosco², C. Marzachi^{1*}

¹ Istituto di Virologia Vegetale, CNR, Strada delle Cacce 73, 10135 Torino, Italy

² University of Torino, Di.Va.P.R.A., Entomologia e Zoologia Applicate all’Ambiente, Via
Leonardo da Vinci 44, Grugliasco (TO), Italy

***Corresponding author:** Cristina Marzachi; c.marzachi@ivv.cnr.it

Keywords: grapevine yellows, FD, susceptibility, Vitis vinifera, phytoplasma multiplication, real
time PCR quantification

16

17 **Abstract**

18

19 Flavescence dorée phytoplasma (FDP) titre in the two red grapevine cultivars Barbera and Nebbiolo
20 was measured over the vegetative seasons of two consecutive years in two vineyards of Piemonte
21 Region (north-western Italy), with a double absolute quantification of FDP cells and grapevine
22 DNA in real time PCR. The relationships of pathogen concentration to cultivar susceptibility and
23 symptom severity were investigated. FD titre was always higher in Barbera than in Nebbiolo
24 infected vines, and this difference was significant at early and late summer samplings of 2008 and
25 at early summer sampling of 2009. A seasonal trend in FD concentration (low in spring, high in
26 early summer and intermediate in late summer) was conserved for Barbera and Nebbiolo in both
27 years and vineyards. Considering both cultivars and years from both vineyards, a significant
28 positive correlation between FD concentration and symptom severity was found in the spring
29 samples. Regarding the FDP strains (-C or -D), no differences in pathogen titres were detected for
30 both cultivars. Similarly, the presence of another grapevine yellows phytoplasma, Bois Noir, a
31 subgroup 16SrXII-A phytoplasma, in mixed infection with FDP strains had no effect on FDP
32 concentration. These results demonstrate for the first time that grapevine cvs with different
33 susceptibility to FDP support different pathogen titres.

34

35 **Introduction**

36

37 Phytoplasmas are wall-less plant pathogenic bacteria of the class Mollicutes associated with
38 diseases of numerous plant species (Bertaccini, 2007). They are phloem-limited and transmitted by
39 phloem-sucking leaf-hoppers, plant-hoppers and psyllids (Marzachi et al., 2004), (Weintraub &
40 Beanland, 2006). Phytoplasmas are classified on the basis of the highly conserved 16S rRNA

sequence into more than 30 groups and within the '*Candidatus* Phytoplasma' genus (Firrao et al., 2004), (Zhao et al., 2010).

Flavescence dorée (FD) is an epidemic, economically important, quarantine disease of grapevine in France, Italy and Spain (Boudon-Padieu, 2003). The disease is associated with a phytoplasma belonging to the 16SrV (Elm yellows) taxonomic group (Lee et al., 2000). FD epidemics are caused by phytoplasma strains belonging to three phylogenetic strain clusters, according to multilocus sequence analysis of *map*, *uvrB-degV* and *secY* loci (Arnaud et al., 2007). In Italy the disease is caused by phytoplasma isolates belonging to clusters 2 and 3, and in Piemonte (north-western Italy) FD strains within the highly variable cluster 3 (FD-C) are the most prevalent (Martini et al., 2002), although isolates of the less variable cluster 2 (FD-D) are also present. The leafhopper *Scaphoideus titanus* Ball is the specific vector of the different FDP strains to grapevine under natural conditions (Mori et al., 2002, Papura et al., 2009, Schvester et al., 1963). Recently, a role as potential vector of FDP from alternative hosts to grapevine has been suggested for the planthopper *Dictyophara europea* (L.) (Filippin et al., 2009).

FD-infected grapevines usually show symptoms the year after inoculation (Morone et al., 2007), although longer latencies have been reported (Osler et al., 2002). Yellowing, downward curling of leaves, fruit abortion, stunting, and lack of lignification of new shoots are among the most important symptoms (Caudwell, 1983, Caudwell, 1990). Infected grapes dramatically reduce their production, but following the first year of symptom expression, a spontaneous and cultivar-dependent remission of symptoms may occur, and asymptomatic plants usually do not contain detectable FD phytoplasma (Morone et al., 2007).

Control of the disease relies mainly on compulsory insecticide treatments to reduce vector population and roguing of infected plants. No resistance is available to FD, but cultivars with different susceptibility to the disease have been reported (Kuzmanovic et al., 2008). Barbera and Nebbiolo are two traditional and economically important grapevine cultivars of Piemonte for the production of red wines. Barbera is highly susceptible to FD and shows severe symptoms starting

with growth reduction at the beginning of the vegetative season, while symptoms on Nebbiolo vines are milder, especially at the beginning of the vegetative season (Morone et al., 2001). In this study we describe a new approach to real time quantification of FD phytoplasma in grapevine which was used to determine a) relationships between cultivar susceptibility and pathogen concentration in the plant, b) relationships between pathogen presence/concentration and symptom severity, c) FD phytoplasma titre over the vegetative season in order to identify the best timing for sampling and detection.

Materials and methods

Vineyard description

Two vineyards located in Piemonte (north-western Italy) were chosen: Cocconato (Asti province) and Monteu Roero (Cuneo province). Barbera and Nebbiolo cvs are planted at the Cocconato vineyard in 1999 on SO4 rootstock. Nebbiolo cv was planted at Monteu Roero vineyard in 1998 also on SO4 rootstock. Both vineyards were subjected to conventional agronomic and phytosanitary practices. An electronic meteorological station was present in Cocconato between the neighbouring Barbera and Nebbiolo vineyards. For the Monteu Roero vineyard, climatic data were collected at a station located in a neighbouring municipality (Canale) in the same exposure and altitude conditions. Mean month temperature (°C) and number of days in each month with minimum temperature below 0°C (freezing days) were calculated from data collected at both stations.

Establishment of a correct sampling procedure

To establish a correct sampling procedure, five FD-infected Barbera and Nebbiolo plants were selected in the Cocconato vineyard at the growth stage 17 – 19 (12 to 16 leaves separated; Coombe, 1995). Symptomatic and asymptomatic shoots were separately labeled on each plant and samples

93 consisting of three basal, three median and three apical leaves (a total of nine leaves) were
94 separately collected from each shoot. Total DNA was extracted from each sample, and the presence
95 of FD phytoplasma was assessed by nested PCR as detailed below. FD phytoplasma titre was then
96 measured in each sample as described below.

97

98 *Grapevine samples collection*

99 Samples were collected in 2008 and 2009 only from symptomatic grapevines. Each sample
100 consisted of three basal, three median and three apical leaves from one symptomatic shoot, for a
101 total of nine leaves. In spring samplings, symptomatic shoots were severely stunted and dwarfed,
102 and therefore the nine leaves were collected from two or three stunted shoots. Barbera samples were
103 collected at Cocconato, while Nebbiolo samples were collected at both Cocconato and Monteu
104 Roero, as not enough FD-infected plants were present at Cocconato. For sample collection,
105 vineyards were surveyed three times a year: late spring (May – early June), early summer (July –
106 early August) and late summer (September). During each survey, symptomatic grapes were labelled
107 (Margaria et al., 2009) and scored for symptom severity. The same vines were sampled in both
108 years provided that they were symptomatic. New symptomatic vines were sampled to replace
109 recovered vines sampled in the previous year. In spring, symptom severity of each infected plant
110 was rated on a scale of 1 to 3 according to the percentage of symptomatic shoots on the fruit bearing
111 branch (1: up to 30%; 2: more than 30 % and up to 65%; 3: more than 65% and up to 100%).
112 During early and late summer surveys, symptom severity of each infected grapevine was rated as A,
113 B, C or D, according to the proportion of the plant canopy showing phytoplasma-specific symptoms
114 (leaf downward rolling, sectorial leaf and vein reddening, bunch drying, premature leaf dropping,
115 presence of brown pustules on canes, and, for late summer surveys, lack of cane lignification): A:
116 plants lacking FD-specific symptoms, but not looking healthy; B: up to 30 % FD-specific
117 symptoms; C: above 30 % and up to 60 %; D: above 60 %.

118

119 *Total DNA extraction and phytoplasma diagnosis*

120 Total DNA was extracted from 1.5 g of leaf veins following a phytoplasma enrichment protocol
121 (Marzachi et al., 1999) and dissolved in 100 µl of sterile double distilled water (SDW). DNA
122 concentration of extracts was quantified with a NanoDrop (ND-1000 Spectrophotometer, NanoDrop
123 Technologies, Inc., Wilmington, DE, USA). Polymerase chain reaction (PCR) was employed for
124 phytoplasma detection. Universal primer pair P1/P7 was used in direct PCRs as previously
125 described (Deng & Hiruki, 1991; Schneider et al., 1995). Reaction products were diluted 1:40 in
126 SDW and used as templates in nested reactions driven by primers R16(I)F1/R1 or R16(V)F1/R1 for
127 the specific detection of 16S groups I and V, respectively (Lee et al., 1994). Amplicons were
128 separated by electrophoresis in 1% agarose gels buffered in 0.5× TBE (90 mM Tris-borate, 2 mM
129 EDTA) and visualized under UV-light after staining with ethidium bromide. To identify Bois Noir,
130 a group 16SrXII (Stolbur group) phytoplasma infection, two µl aliquots of fragments amplified with
131 R16(I)F1/R1 primers were digested for 2 h with 1U of endonuclease *MseI* (Invitrogen, Carlsbad,
132 CA) at 37°C, according to the manufacturer's recommendations. Digestion products were separated
133 in 5% polyacrylamide gels buffered in 1× TBE along with a 1 kb plus DNA size marker (Gibco
134 BRL, Paisley, UK) and visualized by UV-light after staining with ethidium bromide. A Piemonte
135 strain of FDP (FD-C), acquired from grapevine by *Scaphoideus titanus* and transmitted to *Vicia*
136 *faba* L., a French strain of FDP (FD-D), kindly provided by Dr. E. Boudon-Padieu and graft-
137 maintained in periwinkle in the collection of the Istituto di Virologia Vegetale, CNR, and a
138 Sardinian strain of Stolbur phytoplasma from tomato (T2_92), also maintained in periwinkle
139 (Minucci & Boccardo, 1997), were used as reference isolates and positive controls in PCR and
140 RFLP experiments. A healthy Barbera grapevine from *in vitro* propagation, maintained in an insect-
141 proof greenhouse, was employed as negative control in PCR experiments.

142

143 *Characterization of Flavescence dorée phytoplasma isolates*

144 Characterization of Flavescence dorée isolates was done by restriction analysis of the partial 16S-
145 23S rDNA operon and *secY* gene. In the first case, fragments amplified with P1/P7 primer pair
146 (Deng & Hiruki, 1991; Schneider et al., 1995) were diluted 1:40 with SDW, and re-amplified with
147 primers M1/B6 as previously described (Martini et al., 1999). For the partial amplification of the
148 *secY* gene, total grapevine DNAs were amplified in direct PCR with primers FDf2/ FD9r (Angelini
149 et al., 2001), and then, following 1:40 dilution with SDW, amplified with primers FD9f3/FD9r2
150 (Martini et al., 2002). Amplicons were digested for 2 h with one unit of *TaqI* restriction
151 endonuclease (Invitrogen) at 65°C. Digestion products were separated in 5% polyacrylamide gels in
152 TBE along with a 1 kb plus DNA size marker (Gibco BRL, Paisley, UK) and visualized as detailed
153 above.

154

155 | *Quantification of Flavescence dorée phytoplasma DNA*

156 FD phytoplasma titre in infected grapevines was measured as number of FD phytoplasma cells per
157 ng of grapevine DNA. To determine the number of FD phytoplasma cells, specific primers,
158 targeting a portion of the *secY* gene of FD phytoplasma were designed using Primer3 software
159 (Rozen & Skaletsky, 2000), and used in an absolute quantification real-time PCR-based assay. For
160 the quantification of grapevine DNA, specific primers, targeting the 18S rDNA gene of grapevine
161 were designed. The chosen primers were subjected to BLASTN analysis on GenBank through the
162 NCBI database to test their specificity. To confirm the exclusive presence of the expected specific
163 amplicon, 5 µL of real time PCR products were analyzed by electrophoresis in 1.5% agarose gel
164 buffered in TBE. Standard curves for the absolute quantification of FD phytoplasma and grapevine
165 DNA were obtained by dilution of plasmid p26SecYFD, containing the appropriate *secY* gene
166 target sequence from FD-C phytoplasma, and of total DNA extracted from healthy grapevine grown
167 from seed, respectively. Three quantities of p26SecYFD (0.5 ng, 5 pg, and 5 fg), were used,
168 corresponding to $1.17 \cdot 10^8$, $1.17 \cdot 10^6$, $1.17 \cdot 10^3$ cells of FDP. For the absolute quantification of
169 grapevine DNA, three quantities of total healthy grapevine DNA (5 ng, 0.05 ng, and 0.005 ng) were

7

used. Grapevine sample DNAs were diluted in SDW to a final concentration of 1 ng/μl, and used (5 μl) as template in real time assays together with Platinum SYBR Green qPCR Supermix UDG (Life Technologies, Carlsbad, CA, USA) and specific primer pairs at a final concentration of 300 nM, in a volume of 25 μL. The PCR was performed in 96-well plates in a BioRad iCycler (BioRad). Cycling conditions were as follows: one cycle at 95°C for 5 min, then 40 cycles at 95°C for 15 s, and 61°C for 1 min. For each sample, plant- and FDP-DNAs were amplified in the same plate with their corresponding reaction mixture, together with the three standard dilutions of healthy host DNA (Hg) and p26SecYFD. Each sample was run in triplicate in the same plate. For each PCR, water instead of DNA was used as negative control. Threshold levels, threshold cycles, and standard curves were automatically calculated by the BioRad iCycler software, version 3.06070. Per-well baseline cycles were determined automatically. Specificity of the reaction was tested by running a melting curve analysis of the amplicons following each quantification reaction.

Data analyses

For the analyses of phytoplasma concentration, raw data (expressed as log values) were used and the concentration of FDP in each plant was expressed as the difference between the logarithm of the number of FDP cells and the logarithm of the ng of grapevine DNA (DLog). FDP titres in Barbera and Nebbiolo grapevines were analysed by Genstat 11th edition rel. 11.1 (VSN International Ltd, Hemel Hempstead, U.K.). T-test and ANOVA were used to compare titre of FDP in samples, while the Mann-Whitney U test was used to compare symptom severity. Spearman's Rank correlation analysis was used to study relationships between symptom severity and FDP titre in spring, early and late summer samples of the two years.

Results

196 *Symptom severity of infected grapevines and temperature parameters of the vineyards*

197 The number of samples collected and symptom severity ratings of grapevine cultivars Barbera and
198 Nebbiolo surveyed in vineyards during the spring, early summer and late summer of 2008 and 2009
199 are summarized in Table 1.

200 Overall, 59 Barbera samples were collected in the two surveys of 2008 (n = 41) and 2009 (n = 18).

201 [Details on the number of plants showing symptoms of the different categories in the two years are](#)
202 [presented in Table 1. During the spring surveys of the two years, 18 samples were collected, 15 in](#)
203 [2008 and three in 2009. During the early summer surveys, 24 samples were collected from](#)
204 [symptomatic Barbera vines, 16 in 2008, and eight in 2009. For early summer samplings, 4 in 2008](#)
205 most plants showed class B and C symptoms, and in 2009 most of them showed class D symptoms
206 (Table 1). [During the late summer surveys, 17 samples were collected from symptomatic Barbera](#)
207 [vines, ten in 2008, and seven in 2009. For late summer samplings, 4 in 2008, most of the plants](#)
208 showed class B and C symptoms, and in 2009 most of them showed class D symptoms (Table 1).

209 Overall, 60 samples were collected from Nebbiolo grapevines. [As for the Barbera plants, details on](#)
210 [the number of Nebbiolo vines showing symptoms of the different categories in the two years are](#)
211 [presented in Table 1. During the spring surveys of the two years, 8 samples were collected, five in](#)
212 [2008 and three in 2009. During the early summer surveys of the two years, 28 samples were](#)
213 [collected, 15 in 2008 \(11 from Monteu Roero and 4 from Coccoconato, respectively\), and 13 in 2009](#)
214 [\(9 from Monteu Roero and 4 from Coccoconato\). For early summer surveys, 4 in 2008, none of the](#)
215 infected plants showed symptoms of class D, and in 2009 none of the plants showed symptoms of
216 class A (Table 1). [During the late summer surveys of the two years, 24 samples were collected from](#)
217 [symptomatic vines, 14 in 2008 \(10 from Monteu Roero and 4 from Coccoconato\), and 10 in 2009 \(7](#)
218 [from Monteu Roero and 3 from Coccoconato\). For late summer surveys, 4 in 2008 most of the plants](#)
219 showed symptoms of classes B and C, and in 2009 about 50 % of the infected grapevines showed
220 symptoms of class D (Table 1).

221 In the Cocconato vineyard, spring symptoms were more severe in Barbera than in Nebbiolo plants
222 in both years ($P=0.991$) (Table 2). FD symptoms in Barbera and Nebbiolo infected plants at
223 Cocconato were more severe in 2009 than in 2008 (Barbera, early summer: $P=0.999$; late summer:
224 $P=0.996$; Nebbiolo, early summer: $P=0.975$; late summer: $P=0.990$). In 2008, early summer
225 symptoms at Cocconato were more severe on Barbera than on Nebbiolo ($P=0.999$). In the same
226 year, early summer symptoms on Nebbiolo plants were more severe in Monteu Roero vineyard
227 compared to Cocconato ($P=0.999$). The presence of BN phytoplasma in mixed infection with FDP
228 in few plants had no effect on symptom expression on both Barbera and Nebbiolo (Supplementary
229 table 1). Also FDP infection of the plant with -C or -D isolates did not influence symptom
230 expression on both Barbera and Nebbiolo (Supplementary table 1).

231 In the Monteu Roero vineyard, mean winter (November to March) and spring (April and May)
232 temperatures in 2008 were 4.8°C and 16.1°C, and in 2009 they were 4.1°C (winter) and 14.7°C
233 (spring), with 63 (2008) and 79 (2009) freezing days during the winter time. In the Cocconato
234 vineyard, mean winter and spring temperatures in 2008 were 6.7°C and 17.1°C, and in 2009 they
235 were 5.6°C (winter) and 15.5°C (spring), with 41 and 49 freezing days recorded in 2008 and in
236 2009, respectively.

237
238 *Phytoplasma detection and Flavescence dorée strain characterization*

239 Table 3 reports the number of Barbera and Nebbiolo grapevines positive for the presence of FD and
240 BN phytoplasmas alone or as mixed infection among the total samples collected in the surveys of
241 2008 and 2009. In spring surveys of 2008, almost all symptomatic Barbera samples were singly
242 infected with FDP, and most Nebbiolo plants were doubly infected with FDP and BNP. In spring
243 2009, all symptomatic plants were singly infected with FDP, irrespective of cv identity and
244 vineyard. In early summer surveys of 2008 and 2009, most symptomatic Barbera and Nebbiolo
245 samples were infected with FDP only, although BNP and FDP-doubly infected grapes were found
246 in both cvs and vineyards. In 2008 and 2009 late summer surveys, most symptomatic Barbera and

247 Nebbiolo vines were infected with FDP in single infection; few FDP and BNP mixed infections
248 were also found in symptomatic plants sampled in late summer surveys of both years. BN
249 phytoplasma was never detected as single infection in grapevines analyzed in any survey during
250 each of the two years.

251 Overall, in the Cocconato vineyard, only FD-C type phytoplasmas (16S/secY; n = 17) were found
252 in infected Nebbiolo plants sampled in the two years. FD-C phytoplasmas were detected in most
253 Barbera samples collected in all surveys (n = 56), and three FD-D isolates were detected in 2008
254 surveys. FDP isolates from one Nebbiolo and six Barbera vines from the Cocconato vineyard could
255 not be assigned to a type isolate. In the Monteu Roero vineyard, FD-C (n = 25) and -D (n = 11)
256 phytoplasmas were both present in the surveys of 2008, and 2009. FDP isolates from six grapes
257 from this vineyard could not be properly assessed.

258

259 *Quantification of Flavescence dorée phytoplasma*

260 The following primers were designed for the absolute quantification of FDP and grapevine DNA:
261 FdSecyFw/FdSecyRv (5' TGCCTTATGTTACTGCTTCT 3' / 5' TAATAATGATGGGGATTCAA
262 3'), and Vitis18SF1/Vitis18SR1 (5' ATGATTAACAGGGACAGTCG3' /
263 5' GGTATCTGATCGTCTTCGAG3'). BLAST analysis of primers FdSecyFw/FdSecyRv, flanking
264 a 185 bp region of the FD-70 isolate *secY* gene (accession number: AM238512), showed their
265 specificity for the target gene of several "*Ca. P. vitis*" isolates and phytoplasmas belonging to other
266 16Sr-V subgroups. These primers did not match any Stolbur or "*Ca. P. asteris*" related sequences,
267 therefore preventing aspecific amplification from two phytoplasmas commonly infecting grapevine.

268 The FdSecyFw/FdSecyRv amplified a phytoplasma-specific amplicon of the expected size only
269 from FD-C and FD-D infected periwinkle DNA, and gel electrophoresis confirmed the absence of
270 nonspecific products (not shown). Melting curve analysis of the amplicon obtained with
271 FdSecyFw/FdSecyRv showed a unique melting peak at 76.5°C (Fig. 1, Panel **BC**). The
272 Vitis18SF1/Vitis18SR1 primers amplified a grapevine-specific amplicon of the expected size (152

bp) and gel electrophoresis confirmed the absence of nonspecific products (not shown). Melting curve analysis of the amplicon showed a unique melting peak at 85.5°C (Fig. 1, Panel D). Serial dilutions of FDP plasmid DNA ranging from 2.143×10^7 to 2.143×10^3 copies of plasmid were tested in triplicate and the CT values were plotted against the copy number. Serial dilution of healthy grapevine DNA ranging from 50 ng to 0.05 ng were also tested in triplicate and the CT values were plotted against the DNA concentration. The linear correlation coefficient (r) between the CT and the logarithm of the DNA copy number was repeatedly greater than 0.995 for the amplification of both targets (Fig. 1, panels A and C). PCR efficiencies of both curves were constantly around 108% and 103% (plant and phytoplasma, respectively).

Formatted: Superscript

Establishment of a correct sampling procedure

Results of this preliminary experiment are presented in Table 4. FD phytoplasma was detected by nested PCR in all samples from Barbera and Nebbiolo symptomatic shoots, confirming the infected status of the analyzed plants. FD phytoplasma was also detected in three samples from asymptomatic shoots of two Barbera plants, while it was never detected in asymptomatic shoots of infected Nebbiolo plants. FD phytoplasma titre was measured in all samples from symptomatic shoots of Barbera and in two nested PCR positive sample from the two symptomatic shoots of one Nebbiolo plant. FD phytoplasma titre in symptomatic shoots of the five Barbera plants ranged from about 39 to 1180 cells/ng grapevine DNA, and it was above the quantification threshold in one of the three nested PCR positive samples from asymptomatic Barbera shoots. In Barbera, the variance among FDP titre within shoots of each plant was lower than that among plants (Table 5), therefore, collection of leaves from one symptomatic shoot per plant was set as sampling protocol for successive quantification experiments.

Flavescence dorée phytoplasma concentration in Barbera and Nebbiolo grapevines

Despite the low number of FD-D phytoplasma infected plants over the two years, tests indicated that no significant differences were recorded in the concentration of FD-C and -D phytoplasmas within each sampling of the different cvs (Table 6). Further tests also showed that the presence of BNP had no effect on FDP concentration (Table 7). FDP concentration in Barbera grapevines sampled in spring, early and late summer of 2008 and 2009 are illustrated in Fig. 2. Mean FDP concentration in Barbera samples was 242 cells/ng grapevine DNA in spring 2008, and 2 cells/ng grapevine DNA in spring 2009. In early summer samplings, FDP titre was 8,856 and 8,735 cells/ng grapevine DNA in 2008 and in 2009, respectively; in late summer, FDP concentration was 6,420 cells/ng grapevine DNA in 2008 and 671 in 2009. Due to significant differences in the variance, FDP titres in spring and early summer samplings were analyzed separately from those of late summer. Significant differences were recorded in FDP titre between sampling seasons and years. FDP concentration was consistently higher in 2008 than in 2009 (3.299 vs 2.456 DLog; $N_1 = 31$; $N_2 = 10$; S.E.D. = 0.2799; least significant difference at 0.05 level = 0.5672), and the difference was mainly due to phytoplasma titre in spring, since differences between FDP titre in 2008 and 2009 early summers were not significant (least significant difference at 0.05 level between two years: spring = 0.0618; early summer = 0.6108). FDP concentration was also significantly higher in 2008 late summer sampling compared to 2009 ($P = 0.072$). A seasonal trend in FDP concentration (low in spring, high in early summer and intermediate in late summer) was conserved in Barbera vines sampled in both years (Fig. 2).

FDP titre in Nebbiolo vines (Fig. 2) was significantly different according to year (2008 and 2009) and sampling season (spring, early and late summer), while it was similar in both vineyards (Table 8). Mean FDP titre in Nebbiolo vines was 134 cells/ng grapevine DNA in spring samples, 1,417 in early summer samples and 489 in late summer samples of 2008. In 2009, mean FDP titre in spring samples was 2 cells/ng grapevine DNA, in early summer 545 cells/ng grapevine DNA and in late summer samples 334. All data were then pooled to analyze differences between years and sampling seasons. Overall, in 2008, FDP in Nebbiolo was more abundant than in 2009 (2.837 vs 2.310 DLog;

324 $N_1 = 32$; $N_2 = 26$; S.E.D. = 0.1819; least significant difference at 0.05 level = 0.3651), and the
 325 difference was mainly due to the concentration of the spring samples, since differences between the
 326 remaining seasons were not significant (least significant different at 0.05 level between single
 327 seasonal samplings of the two years: spring = 1.0078; early summer = 0.5229; late summer =
 328 0.5909). There was no difference in the FDP titre of early and late summer samples of the two years
 329 (least significant difference at 0.05 level between early and late summer 2008 = 0.5345 and between
 330 early and late summer 2009 = 0.5805), but a seasonal trend in FDP concentration (low in spring,
 331 high in early summer and intermediate in late summer) was conserved for Nebbiolo in both years
 332 and vineyards (Fig. 2).
 333 FDP concentration was always higher in Barbera than in Nebbiolo infected vines, and this
 334 difference was significant at early and late summer samplings of 2008 (early summer: $P=0.008$; late
 335 summer: $P=0.002$) and at early summer sampling of 2009 ($P < 0.001$) (Fig. 2).
 336

337 *Flavescence dorée phytoplasma concentration and symptom severity*

338 Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples
 339 showed a significant, positive correlation between FDP concentration and symptoms severity
 340 ($r=0.412$; S.E.=0.367; $N=25$; $DF=23$; $t=1.89$; $P=0.071$ Spearman's correlation coefficient $r=0.412$;
 341 adjusted for ties = 0.367; $N=25$; $DF=23$; $t=1.89$; $P=0.071$), and so did the Barbera samples of 2008
 342 (Spearman's correlation coefficient $r=0.389$; $N=25$; $DF=23$; t approximation 1.72 ; $P=0.099$), but
 343 not in 2009, ~~pooling both early and late summer samplings ($r=0.389$; S.E.=0.377; $N=25$; $DF=23$;~~
 344 ~~$t=1.72$; $P=0.099$)~~, and the Nebbiolo samples of 2009 (Spearman's correlation coefficient $r=0.430$;
 345 adjusted for ties $r=0.373$; $N=22$; $DF=20$; $t=1.98$; $P=0.087$), but not in 2008, ~~pooling both early and~~
 346 ~~late summer samplings ($r=0.448$; S.E.=0.397; $N=23$; $DF=21$; $t=1.98$; $P=0.061$)~~. Early and late
 347 summer samples were analyzed together for correlations between FD concentration (as Log[FD])
 348 and symptom severity as the two samplings were evaluated on the same severity scale. No other
 349 significant correlation was detected.

Overall, there was no highly significant correlation between FDP concentration and symptoms severity.

Discussion

Flavescence dorée infection in 2008 and 2009 in the two vineyards of Coconato and Monteu Roero was caused mainly by Flavescence dorée type C phytoplasma (FD-C). This isolate is stably present in the area since the early outbreak of the epidemics in 1998, and it is the most represented one in FD-infected vines of Piemonte region (Marzachi et al., 2001). FD-D type phytoplasmas were detected sporadically in infected Barbera and Nebbiolo grapevines collected in spring, early and late summer of both years. Grapevines showing early symptoms were mostly infected with FD-C phytoplasma, as reported in other grapevine-growing areas (Angelini et al., 2006), and this isolate was detected throughout the vegetative season. Bois noir disease is also present in Piemonte vineyards (Marzachi et al., 2001), and BNP was found only in mixed infections with FDP in samples from the two vineyards. As we only sampled symptomatic grapevines in the two highly FD-infected locations, the presence of BNP in asymptomatic plants cannot be ruled out.

~~The FDP type (C or D) had no effect on symptom severity of the infected vines of both cultivars. Also, the presence of BNP in mixed infection, a rare occurrence, had no effect on symptom expression. Under these experimental conditions, a competition effect between BNP and FDP cannot be ruled out.~~

For two consecutive years, infected Barbera vines showed more severe spring symptoms than Nebbiolo ones. This observation coincides with the fact that Barbera is known as a highly FD-sensitive cv. Both cvs showed more severe symptoms in 2009 than in 2008, and in 2009 lower winter and spring mean temperatures and more freezing days were recorded at both localities than in 2008. Low spring temperatures have an immediate effect on grapevine in delaying bud burst

(Tomasi et al., 2007), but they also have a long term effect in reducing fruit set (%) and berry number per bunch in a cv-specific manner (Ebadi et al., 1995). Temperature stresses also affect Ca^{2+} homeostasis in grapevine (Wang et al., 2004). Also, low temperatures reduce respiration rate in grapevine (Franck et al., 2011), thus reducing the production of H_2O_2 (Foyer et al., 2009), a molecule probably involved in defense response pathways to phytoplasmas (Musetti et al., 2007). We can speculate that more severe FD symptoms in both cvs in early and late summer assessment of 2009 may be an effect of a long-lasting stress of the plant caused by the occurrence of low winter/spring temperatures in that year. Moreover, in 2008, temperature probes recorded lower winter/spring mean temperatures and more freezing days in the Monteu Roero vineyard compared to Cocconato, and significantly more severe early summer symptoms were recorded in Nebbiolo infected vines in Monteu Roero than in Cocconato. Quantification of FDP-DNA was obtained as the ratio of phytoplasma DNA per ng of grapevine DNA as previously described for “*Ca. P. asteris*” (Marzachi & Bosco, 2005). Real time based PCR assays for diagnosis and characterization of several phytoplasmas, including FDP, have been proposed (Angelini et al., 2007, Bianco et al., 2004, Galetto et al., 2005, Hren et al., 2007, [Mehle et al., 2013](#), Pelletier et al., 2009), but this is [one of the first attempts, together with the one of Prezelj and co-workers \(2012\)](#), to quantify FDP in field-infected vines of different cvs. A double absolute quantification assay was developed and only samples with a threshold cycle value within that of the lowest standard concentration were considered. The quantification of less than 3 phytoplasma cells per ng plant DNA in three spring samples can still be considered reliable according to the MIQE guidelines (Bustin et al., 2009) as it exceeds by three times the most sensitive limit of detection. It is known that phytoplasmas have uneven distribution in the grapevine canopy, but a direct correlation between the presence of phytoplasma specific symptoms and FDP detection was clearly shown as all samples collected from symptomatic shoots of both cvs were positive for the presence of FDP, while all samples from asymptomatic Nebbiolo and most from asymptomatic Barbera were nested PCR negative. [Similar results are reported for other grapevine varieties \(Prezelj et al., 2012\)](#)

where FDP was mainly associated with symptomatic tissues, although it was also detected in some symptomless tissues of highly infected (and symptomatic) plants. Also, the differences between FDP titre in different symptomatic shoots from the same Barbera plant were lower than the differences among plants, therefore confirming that sampling from one symptomatic shoot can provide a reliable and representative evaluation of FDP titre in the infected parts of the plant.

We can suggest that FD-C and -D phytoplasmas multiply at the same level in grapevine, and that the presence of BNP has no influence on the accumulation of FDP in doubly infected plants. Very few quantitative information on plants co-infected by two phytoplasmas are available, however, mild and severe isolates of "*Ca. P. asteris*" show different multiplication rate and movement in the herbaceous host (Sinha, 1983) and, in the case of Ash yellows phytoplasma, strains that differ in aggressiveness have a different multiplication and/or movement following co-inoculation to the laboratory host periwinkle (Sinclair & Griffiths, 2000). In Apple proliferation (AP) infected apple trees, the ratio of mild to severe strains in the phytoplasma population determines the virulence of the infection and shifts in the population composition may alter virulence of multiple strain accession (Seemueller et al., 2011).

In both cvs and in both years, the lowest FDP titre was observed in spring, the highest in early summer and an intermediate one at the end of the vegetative season. A continuous increase of FDP titre up to the end of the vegetative season was reported for other grapevine cvs in Slovenia (Prezelj et al., 2012). Seasonal fluctuations of phytoplasma populations in AP-infected apple trees have been measured with highest levels occurring from December to May (Baric et al., 2011). Despite the occurrence of more severe symptoms throughout the vegetative season of 2009 compared to 2008, at each sampling date, FDP titre was significantly higher in 2008 than in 2009, irrespective of the cv. This difference was most evident in the spring samplings, when as many as hundreds FDP cells were recorded in 2008 in contrast to only as few as 2 FDP cell per ng plant DNA in 2009. Since the lowest FDP titre was consistently measured in both cvs, both years and at both sites in spring, winter temperature together with plant dormancy and pruning, that eliminates most of the infected

plant parts, have a cooperative effect in reducing phytoplasma titre in the plant. Multiplication of pathogen resumed with the vegetative season and the differences in FDP titres between the two years were reduced, although FDP-specific symptom expression was more severe in 2009 than in 2008. We can speculate that colder winter/early spring climatic conditions have a long-lasting stress effect on the grapevines, increasing both their sensitivity to FD and symptom severity even in the presence of a low pathogen load. Also, the hypothesis that the pathogen may multiply less efficiently in the stressed plant, therefore reaching a low titre despite the severe symptoms, cannot be ruled out.

FDP titre in Barbera grapevines was always higher than in Nebbiolo. This is in good correlation with more severe symptoms expressed by FD-infected Barbera vines compared to Nebbiolo ones and provides the most likely explanation for the milder symptoms shown by FDP-infected Nebbiolo. Actually this latter cv sustains a low multiplication of the pathogen. Proteomic and transcriptomic analyses of FDP-infected Nebbiolo plants show that most proteins modulated during infection belong to the “cell rescue, defense and virulence” class (Margaria & Palmano, 2011a). On the other hand, preliminary studies on Barbera cv, show a lower presence of proteins of the ‘defense’ category compared to the total identified proteins (Margaria & Palmano, 2011b). Perhaps different responses of the two cvs to FDP may account for the different multiplication level of the pathogen, but this hypothesis remains untested.

The results indicate that the best time for sampling and detection of FDP infection is early summer, when phytoplasma titre is highest. [However, our work, together with the one of Prezelj and co-workers \(2012\) provide evidence that FDP can be detected as early as at flowering season, or even before in the case of Barbera in our work.](#) Quantitative real time PCR is a useful tool for the evaluation of phytoplasma multiplication and can be used when screening for resistance/tolerance.

We have demonstrated that Barbera supports a higher FDP multiplication compared to Nebbiolo, and further studies are required to understand whether this has an effect on vector acquisition and

transmission efficiency, as already suggested for Pinot blanc and Merlot cultivars (Bressan et al., 2005).

Acknowledgements

We thank the wine growers L. Negro and P. Bava for allowing us to collect samples in their vineyards, F. Spanna (Piemonte Region, Settore Fitosanitario) for providing us climatic values at both locations, and C. Lovisolo (University of Torino) for his critical reading of the manuscript. This work was supported by the Italian Regional Grant (Piemonte Region) “Studi sui fattori che favoriscono le epidemie di Flavescenza dorata in Piemonte e loro superamento.”, and by the Piemonte Region CIPE "Adoption of a multidisciplinary approach to study the grapevine agroecosystem: analysis of biotic and abiotic factors able to influence yield and quality".

Literature cited

Angelini E, Bianchi GL, Filippin L, Morassutti C, Borgo M, 2007. A new TaqMan method for the identification of phytoplasmas associated with grapevine yellows by real-time PCR assay. *Journal of Microbiological Methods* **68**, 613-22.

Angelini E, Clair D, Borgo M, Bertaccini A, Boudon-Padieu E, 2001. Flavescence dorée in France and Italy - Occurrence of closely related phytoplasma isolates and their near relationships to Palatinate grapevine yellows and an alder yellows phytoplasma. *Vitis* **40**, 79-86.

Angelini E, Filippin L, Michielini C, Bellotto D, Borgo M, 2006. High occurrence of Flavescence dorée phytoplasma early in the season on grapevines infected with grapevine yellows. *Vitis* **45**, 151-2.

479 Arnaud G, Malembic-Maher S, Salar P, Bonnet P, Maixner M, Marcone C, Boudon-Padieu E,
480 Foissac X, 2007. Multilocus sequence typing confirms the close genetic interrelatedness of three
481 distinct Flavescence dorée phytoplasma strain clusters and group 16SrV phytoplasmas infecting
482 grapevine and alder in Europe. *Applied and Environmental Microbiology* **73**, 4001-10.

483 Baric S, Berger J, Cainelli C, Kerschbamer C, Letschka T, Dalla Via J, 2011. Seasonal colonisation
484 of apple trees by '*Candidatus* Phytoplasma mali' revealed by a new quantitative TaqMan real-time
485 PCR approach. *European Journal of Plant Pathology* **129**, 455-67.

486 Bertaccini A, 2007. Phytoplasmas: diversity, taxonomy, and epidemiology. *Frontiers in Bioscience*
487 **12**, 673-89.

488 Bianco PA, Casati P, Marziliano N, 2004. Detection of phytoplasmas associated with grapevine
489 Flavescence dorée disease using real-time PCR. *Journal of Plant Pathology* **86**, 257-61.

490 Boudon-Padieu E. The situation of grapevine yellows and current research directions: distribution,
491 diversity, vectors, diffusion and control. *Proceedings of the Proceedings of the 14th Meeting of the*
492 *International Council for the Study of Virus and Virus-like Diseases of the Grapevine 2003*.
493 Locorotondo, Bari, Italy, 47-53.

494 Bressan A, Spiazzi S, Girolami V, Boudon-Padieu E, 2005. Acquisition efficiency of Flavescence
495 dorée phytoplasma by *Scaphoideus titanus* Ball from infected tolerant or susceptible grapevine
496 cultivars or experimental host plants. *Vitis* **44**, 143-6.

497 Bustin SA, Benes V, Garson JA, Hellemans J, Huggett J, Kubista M, Mueller R, Nolan T, Pfaffl
498 MW, Shipley GL, Vandesompele J, Wittwer CT. (2009) The MIQE guidelines: minimum
499 information for publication of quantitative real-time PCR experiments. *Clinical Chemistry* **55**, 611-
500 622.

501 Caudwell A, 1983. L'origine des jaunisses à mycoplasmes (MLO) des plantes et l'exemple des
502 jaunisses de la vigne. *Agronomie* **3**, 103-11.

503 Caudwell A, 1990. Epidemiology and characterization of Flavescence dorée (FD) and other
504 grapevine yellows. *Agronomie* **10**, 655-63.

- 505 Coombe G, 1995. Growth stages of the grapevine. *Australian Journal of Grape and Wine Research*
506 **1**, 100-110.
- 507 Deng S, Hiruki C, 1991. Amplification of 16S rRNA genes from culturable and non culturable
508 mollicutes. *Journal of Microbiological Method* **14**, 53-61.
- 509 Ebadi A, Coombe BG, May P, 1995. Fruit-set on small Chardonnay and Shiraz vines grown under
510 varying temperature regimes between budburst and flowering. *Australian Journal of Grape and*
511 *Wine Research* **1**, 3-10.
- 512 Filippin L, Jovic J, Cvrkovic T, Forte V, Clair D, Tosevski I, Boudon-Padieu E, Borgo M, Angelini
513 E, 2009. Molecular characteristics of phytoplasmas associated with Flavescence dorée in clematis
514 and grapevine and preliminary results on the role of *Dictyophara europaea* as a vector. *Plant*
515 *Pathology* **58**, 826-37.
- 516 Firrao G, Andersen M, Bertaccini A, Boudon-Padieu E, Bove J M, Daire X, Davis R E, Fletcher J,
517 Garnier M, Gibb K S, Gundersen-Rindal D E, Harrison N, Hiruki C, Kirkpatrick B C, Jones P,
518 Kuske C R, Lee I M, Liefting L, Marcone C, Namba S, Schneider B, Sears B B, Seemuller E, Smart
519 C D, Streten C, Wang K, 2004. '*Candidatus* Phytoplasma', a taxon for the wall-less, non-helical
520 prokaryotes that colonize plant phloem and insects. *International Journal of Systematic and*
521 *Evolutionary Microbiology* **54**, 1243-55.
- 522 Foyer C H, Bloom A J, Queval G, Noctor G, 2009. Photorespiratory Metabolism: Genes, Mutants,
523 Energetics, and Redox Signaling *Annual Review Plant Biology* **60**, 455-84.
- 524 Franck N, Morales JP, Arancibia-Avendano D, Garcia de Cortazar V, Perez-Quezada J F, Zurita-
525 Silva A, Pastenes C, 2011. Seasonal fluctuations in *Vitis vinifera* root respiration in the field. *New*
526 *Phytologist* **192** 939-51.
- 527 Galetto L, Bosco D, Marzachi C, 2005. Universal and group-specific real-time PCR diagnosis of
528 Flavescence dorée (16Sr-V), Bois noir (16Sr-XII) and apple proliferation (16Sr-X) phytoplasmas
529 from field-collected plant hosts and insect vectors. *Annals of Applied Biology* **147**, 191-201.

530 Hren M, Boben J, Rotter A, Kralj P, Gruden K, Ravnikar M, 2007. Real-time PCR detection
531 systems for Flavescence dorée and Bois noir phytoplasmas in grapevine: comparison with
532 conventional PCR detection and application in diagnostics. *Plant Pathology* **56**, 785-96.

533 Kuzmanovic S, Martini M, Ermacora P, Ferrini F, Starovic M, Tosic M, Carraro L, Osler R, 2008.
534 Incidence and molecular characterization of Flavescence dorée and stolbur phytoplasmas in
535 grapevine cultivars from different viticultural areas of Serbia. *Vitis* **47**, 105-11.

536 Lee I M, Davis R E, Gundersen-Rindal D E, 2000. Phytoplasma: Phytopathogenic mollicutes.
537 *Annual Review of Microbiology* **54**, 221-55.

538 Lee I M, Gundersen D E, Hammond R W, Davis R E, 1994. Use of mycoplasmalike organism
539 (MLO) group-specific oligonucleotide primers for nested-PCR assays to detect mixed-MLO
540 infections in a single host-plant. *Phytopathology* **84**, 559-66.

541 Margaria P, Palmano S, 2011a. Response of the *Vitis vinifera* L. cv. 'Nebbiolo' proteome to
542 Flavescence dorée phytoplasma infection. *Proteomics* **11**, 212-24.

543 Margaria P, Palmano S, 2011b. Proteome responses of *Vitis vinifera* L. to 'Flavescence dorée'
544 phytoplasma infection. *Bulletin of Insectology* **64 (Supplement)**, S-167-S8.

545 Margaria P, Turina M, Palmano S, 2009. Detection of Flavescence dorée and Bois noir
546 phytoplasmas, Grapevine leafroll associated virus-1 and -3 and Grapevine virus A from the same
547 crude extract by reverse transcription-RealTime Taqman assays. *Plant Pathology* **58**, 838-45.

548 Martini M, Botti S, Marcone C, Marzachi C, Casati P, Bianco P A, Benedetti R, Bertaccini A, 2002.
549 Genetic variability among Flavescence dorée phytoplasmas from different origins in Italy and
550 France. *Molecular and Cellular Probes* **16**, 197-208.

551 Martini M, Murari E, Mori N, Bertaccini A, 1999. Identification and epidemic distribution of two
552 flavescence doree-related phytoplasmas in Veneto (Italy). *Plant Disease* **83**, 925-930.

553 Marzachi C, Alma A, D'aquilio M, Minuto G, Boccardo G, 1999. Detection and identification of
554 phytoplasmas infecting cultivated and wild plants in Liguria (Italian Riviera). *Journal Plant*
555 *Pathology* **81**, 127-36.

- 556 Marzachi C, Bosco D, 2005. Relative quantification of chrysanthemum yellows (16Sr I)
557 phytoplasma in its plant and insect host using real-time polymerase chain reaction. *Molecular*
558 *Biotechnology* **30**, 117-27.
- 559 Marzachi C, Milne R J, Bosco D, 2004. Phytoplasma-plant-vector relationships. In: Pandalai S G,
560 Gayathri A, eds. *Recent Research Development in Plant Pathology*. Kerala, India: Research
561 Signpost, 211-41.
- 562 Marzachi C, Palermo S, Boarino A, Veratti F, d'Aquilio M, Loria A, Boccoardo G, 2001.
563 Optimisation of a one-step PCR assay for the diagnosis of Flavescence dorée related phytoplasmas
564 in field-grown grapevines and vector populations. *Vitis* **40**, 213-7.
- 565 [Mehle N, Prezelj N, Hren M, Boben J, Gruden K, Ravnikar M, Dermastia M, 2013. A real-time](#)
566 [PCR detection system for the Bois noir and Flavescence dorée phytoplasmas and quantification of](#)
567 [the target DNA. In: Dickinson M, Hodgetts J, eds. Phytoplasma: Methods and Protocols. New](#)
568 [York, USA: Humana Press, Springer Protocols, doi: 10.1007/978-1-62703-089-2_22 \(Methods in](#)
569 [Molecular Biology; vol. 938 pp. 253–68\).](#)
- 570 Minucci C, Boccoardo G, 1997. Genetic diversity in the Stolbur phytoplasma group.
571 *Phytopathologia Mediterranea* **36**, 45-9
- 572 Mori N, Bressan A, Martini M, Guadagnini M, Girolami V, Bertaccini A, 2002. Experimental
573 transmission by Scaphoideus titanus Ball of two Flavescence doree-type phytoplasmas. *Vitis* **41**, 99-
574 102.
- 575 Morone C, Boveri M, Giosue S, Gotta P, Rossi V, Scapin I, Marzachi C, 2007. Epidemiology of
576 Flavescence dorée in vineyards in northwestern Italy. *Phytopathology* **97**, 1422-7.
- 577 Morone C, Gotta P, Marzachi C, 2001. Riconoscimento dei sintomi di inizio stagione della
578 flavescenza dorata. *L'Informatore Agrario* **57**, 83-6.
- 579 Musetti R, Marabottini R, Badiani M, Martini M, di Toppi L S, Borselli S, Borgo M, Osler R, 2007.
580 On the role of H₂O₂ in the recovery of grapevine (*Vitis vinifera* cv. Prosecco) from Flavescence
581 dorée disease. *Functional Plant Biology* **34**, 750-8.

582 Osler R, Zucchetto C, Carraro L, Frausin C, Pavan F, Vettorello G, Girolami V, 2002. Trasmissione
583 di flavescenza dorata e legno nero e comportamento delle viti infette. *L'Informatore Agrario* **19**, 61-
584 5.

585 Papura D, Delmotte F, Giresse X, Salar P, Danet J L, Van Helden M, Foissac X, Malembic-Maher
586 S, 2009. Comparing the spatial genetic structures of the Flavescence dorée phytoplasma and its
587 leafhopper vector *Scaphoideus titanus*. *Infection Genetics and Evolution* **9**, 867-76.

588 Pelletier C, Salar P, Gillet J, Cloquemin G, Very P, Foissac X, Malembic-Maher S, 2009. Triplex
589 real-time PCR assay for sensitive and simultaneous detection of grapevine phytoplasmas of the
590 16SrV and 16SrXII-A groups with an endogenous analytical control. *Vitis* **48**, 87-95.

591

592 [Prezelj N, Nikolic' P, Gruden K, Ravnikar M, Dermastia M, 2012. Spatiotemporal distribution of](#)
593 [flavescence dorée phytoplasma in grapevine. Plant Pathology. Doi: 10.1111/j.1365-](#)
594 [3059.2012.02693.x](#)

595 Rozen S, Skaletsky H J, 2000. Primer3 on the WWW for general users and for biologist
596 programmers. In: Krawetz S M S, eds. *Bioinformatics Methods and Protocols: Methods in*
597 *Molecular Biology*. Totowa, NJ: Humana Press, 365-86.

598 Schneider B, Cousins M T, Klinkong S, Seemuller E, 1995. Taxonomic relatedness and
599 phylogenetic positions of phytoplasmas associated with diseases of faba bean, sunnhemp, sesame,
600 soybean, and eggplant. *Zeitschrift Fur Pflanzenkrankheiten Und Pflanzenschutz-Journal of Plant*
601 *Diseases and Protection* **102**, 225-32.

602 Schvester D, Carle P, Montous G, 1963. Transmission de la Flavescence Dorée de la vigne par *S.*
603 *littoralis* Ball. *Annales des Epiphyties* **14**, 175-98

604 [Seemueller E, Kampmann M, Kiss E, Schneider B, 2011. HflB gene-based phytopathogenic](#)
605 [classification of 'Candidatus Phytoplasma mali' Strains and evidence that strain composition](#)
606 [determines virulence in multiply infected apple trees. Molecular Plant Microbe Interactions](#) **24**,
607 [1258-66.](#)

608 ~~Sinclair W A, Griffiths H M, 2000. Variation in aggressiveness of ash yellows phytoplasmas. *Plant*~~
609 ~~*Disease* **84**, 282-8.~~

610 ~~Sinha R, 1983. Relative concentration of mycoplasma like organisms in plants at various times after~~
611 ~~infection with aster yellows. *Canadian Journal of Plant Pathology* **5**, 7-10.~~

612 Smart C D, Schneider B, Blomquist C L, Guerra L J, Harrison N A, Ahrens U, Lorenz K H,
613 Seemuller E, Kirkpatrick B C, 1996. Phytoplasma-specific PCR primers based on sequences of the
614 16S-23S rRNA spacer region. *Applied and Environmental Microbiology* **62**, 2988-93.

615 Tomasi D, Pascarella G, Sivilotti P, Gardiman M, Pitacco A, 2007. Grape bud burst: thermal heat
616 requirement and bud antagonism. *Acta Horticulturae* **754**, 205-12.

617 Wang L-J, Huang W-D, Li J-Y, Liu Y-F, Shi Y-L, 2004. Peroxidation of membrane lipid and Ca²⁺
618 homeostasis in grape mesophyll cells during the process of cross-adaptation to temperature stresses.
619 *Plant Science* **167**, 71-7.

620 Weintraub P G, Beanland L, 2006. Insect vectors of phytoplasmas. *Annual Review of Entomology*
621 **51**, 91-111.

622 Zhao Y, Wei W, Davis R E, Lee I M, 2010. Recent advances in 16S rRNA gene-based phytoplasma
623 differentiation, classification and taxonomy. In: Weintraub P G, Jones P, eds. *Phytoplasmas:*
624 *genomes, plant hosts and vectors* Wallingford, UK: CAB International, 64-92.

625

626

627 **Captions for figures**

628

629 Figure 1. A: Standard curve of Flavescente dorée phytoplasma DNA obtained by plotting threshold
630 cycles (CT) values vs log of 2 x 10⁷, 2 x 10⁵, 2 x 10³, and 2 x 10 copy number of p26SecYFD
631 plasmid. **BC**: Melting curve analysis of the amplicon obtained with primers FdSecyFw/FdSecyRv.
632 **EB**: Standard curve of plant DNA (*Vitis vinifera*) obtained by plotting threshold cycles (CT) values

633 vs log of 50, 5, 0.5, and 0.05 ng of healthy grapevine DNA. D: Melting curve analysis of the
634 amplicon obtained with primers Vitis18SF1/Vitis18SR1.

635

636 Figure 2. Mean Flavescence dorée phytoplasma concentration (cells / ng grapevine DNA) and
637 standard error in Barbera and Nebbiolo plants sampled in spring, early and late summer surveys of
638 2008 and 2009 in Cocconato (C) and Monteu Roero (MR) vineyards.

639

640

641

642

643

Table 1. Number of samples from Barbera and Nebbiolo grapevines showing phytoplasma-specific symptoms of different categories collected during spring, early and late summer surveys of 2008 and 2009. In spring, symptom severity of infected grapes was rated on a scale of 1 to 3 according to the percentage of symptomatic shoots on the fruit bearing branch (1: maximum 30%; 2: more than 30 % and up to 65%; 3: more than 65% and up to 100%). In early and late summer surveys, symptom severity of infected grapes was rated as A, B, C or D, according to the proportion of the plant canopy showing phytoplasma-specific symptoms (A: plants lacking FD-specific symptoms, but not looking healthy; B: up to 30 % FD-specific symptoms; C: above 30 % and up to 60 %; D: above 60 %).

Year	Cultivar	Spring			Early summer				Late summer			
		1	2	3	A	B	C	D	A	B	C	D
2008	Barbera	1	9	5	2	9	5	0	2	4	4	0
	Total plants	15			16				10			
2009	Barbera	2	1	0	0	1	2	5	0	1	1	5
	Total plants	3			8				7			
2008	Nebbiolo	2	1	2	4	6	5	0	3	6	5	0
	Total plants	5			15				14			
2009	Nebbiolo	2	1	0	0	2	4	7	0	2	3	5
	Total plants	3			13				10			

Table 2. Summary of significant to highly significant comparisons between FD symptom severities in Barbera (B) and Nebbiolo (N) grapevines in 2008 and 2009 at Cocconato (C) and Monteu Roero: (MR).

Season	Vineyard	Cv	Year/s	Comparison	N ^a	U ^b	P ^c
Spring ^d	C		2008+2009	B > N	17, 3	3.0	0.991
		B		2009 > 2008	15, 8	14.5	0.999
Early summer ^e	C	N		2009 > 2008	4, 4	0.0	0.986
			2008	B > N	15, 4	2.0	0.999
	MR	N		2009 > 2008	11, 9	23.5	0.975
		N	2008	MR > C	4, 11	0.0	0.999
Late summer ^f	C	B		2009 > 2008	9, 7	7.5	0.996
	MR	N		2009 > 2008	8, 7	9.0	0.990
	C	N	2008	Es ^e < Ls ^f	4, 4	0.0	0.986

^a N: sample numbers; ^b U: value of U in the Mann-Whitney U test; ^c P:

the exact probability (adjusted for ties) that one group is equal to, greater or smaller than the other

group (according to the sign =, >, < specified in the table); ^d Sp: spring; ^e Es: early summer; ^f Ls: late summer.

; ^d Sp: spring; ^e Es: early summer; ^f Ls: late summer.

For Peer Review

Table 3. Number of Barbera (B) and Nebbiolo (N) grapevines PCR positive for the presence of Flavescence dorée phytoplasma (FDP) as either single or mixed infection with Bois Noir phytoplasma (BNP+FDP) among the total samples collected from vineyards in Cocconato and Monteu Roero during spring (Sp), early (Es) and late summer (Ls) growing seasons of 2008 and 2009.

Year	Vineyard	Cv	FDP			BNP+FDP		
			(Positive/ total sampled)			(Positive/total sampled)		
			Sp	Es	Ls	Sp	Es	Ls
2008	Cocconato	B	14/15	15/16	8/10	1/15	1/16	2/10
		N	0	3/4	2/4	1/1	1/4	2/4
	Monteu Roero	N	1/4	7/11	9/10	3/4	4/11	1/10
2009	Cocconato	B	3/3	8/8	6/7	0	0/8	1/7
		N	2/2	4/4	3/3	0	0	0
	Monteu Roero	N	1/1	8/9	6/7	0	1/9	1/7

Table 4. Number of symptomatic and asymptomatic shoots from the five Barbera and five Nebbiolo FD-infected grapevines from which samples were collected for successive diagnostic direct (P1/P7 primers) and nested (R16(V)F1/R1 primers) PCRs and FDP quantification. Only samples with a threshold cycle value within that of the lowest standard concentration were considered for FDP quantification. Titre of FDP is expressed as phytoplasma cells/ng plant DNA. SE: standard error.

Formatted: Font: (Default) Times New Roman, 12 pt

Cultivar	Symptoms	Shoots (n)	Direct PCR (positive / tested)	Nested PCR (positive / tested)	qPCR (positive / tested)	FDP titer (mean)	SE
Barbera	Symptomatic	11	9/11	11/11	11/11	345.1	103.94
	Asymptomatic	9	1/9	3/9	1/9	9.6	na
Nebbiolo	Symptomatic	7	1/7	7/7	2/7	45.3	5.39
	Asymptomatic	13	0/13	0/13	0/13	na	na

Table 5. Mean FDP titer (expressed as the difference between the logarithm of the number of FDP cells and the logarithm of the ng of grapevine DNA, DLog) in symptomatic shoots of the five Barbera plants, variance (Var) among FDP titres within shoots of each plant and among plants (General, among plants).

Plant	Mean FDP titer in shoots (DLog)	Var
1	2.178167	0.07131606
2	2.1135	0.0772245
3	2.850333	0.09827222
4	2.134833	0.0981245
5	2.7865	0.00098272
General, among plants	2.412667	0.16141599

Table 6. Analysis of variance (ANOVA) for grouping FD-C and FD-D phytoplasma titres of Barbera (B) and Nebbiolo (N) plants sampled at different times in the two vineyards. C, MR: Coconato and Monteu Roero vineyards, respectively. Es, Ls: early and late summer samplings. Log[FD-C], LOG[FD-D]: Log of FD-C or -D phytoplasma concentration. N_C, N_D: number of samples infected by FD-C or FD-D phytoplasmas. *F* pr: F probability. LSD: least significant difference. N.S.: not significant.

Cv	Vineyard	Year	Season	LOG[FD-C] (mean)	N _C	LOG[FD-D] (mean)	N _D	<i>F</i> pr	LSD	Comment
B	C	2008	Es	4.0242	14	3.640	1	0.43 6	1.034	N.S.
			Ls	3.8911	7	3.785	1	0.71 4	0.6746	N. S.
N	MR	2008	Es	3.2713	4	2.7967	3	0.20 6	0.8409	N.S.
		2009	Es	2.7845	6	3.5642	2	0.31 5	1.740	N.S.
			Ls	3.0948	5	2.551	1	0.39 7	1.594	N.S.

Table 7. Analysis of variance (ANOVA) for grouping FDP tires in FD singly infected and FD and BN doubly infected Barbera (B) and Nebbiolo (N) plants sampled at different times in the two vineyards. C, MR: Cocconato and Monteu Roero vineyards, respectively. $\text{Log}[\text{FD}]_{\text{FD}}$, $\text{Log}[\text{FD}]_{\text{FD+BN}}$: Log of FD phytoplasma concentration. N_{FD} , $N_{\text{FD+BN}}$: number of samples infected only by FD or by FD and BN in mixed infection. Sp, Es, Ls: spring, early and late summer samplings. F pr: *F* probability. LSD: least significant difference. N.S.: not significant.

Cv	Vineyard	Year	Season	$\text{LOG}[\text{FD}]_{\text{FD}}$ (mean)	N_{FD}	$\text{LOG}[\text{FD}]_{\text{FD+BN}}$ (mean)	$N_{\text{FD+BN}}$	F pr	LSD	Comment
B	C	2008	Sp	2.384	14	2.370	1	0.989	2.204	N.S.
			Es	4.0242	14	3.1773	1	0.100	1.034	N.S.
		2009	Ls	2.935	6	2.179	1	0.603	3.498	N.S.
N	C+MR ^a	2008+2009	Sp	0.820	4	2.366	2	0.165	2.531	N.S.
	C+MR ^b	2008	Es	3.2644	7	3.3878	4	0.651	0.595 5	N.S.

^a, ^b: according to corresponding symbol in Table 8.

Table 8. Analysis of variance (ANOVA) for grouping FDP titres from infected Nebbiolo plants collected in the two vineyards. Plants infected with FDP (FD-C) of FDP and BNP (FD-C+BN) in double infection were analysed separately. Sp, Es: spring and early summer samplings.

LOG[FD]_{CC}, LOG[FD-D]_{MR}: Log of FDP titre in plants collected at Cocconato or Monteu Roero vineyards, respectively. F pr: *F* probability. LSD: least significant difference. N.S.: not significant.

Infection	Year	Season	LOG[FD] _{CC} (mean)	N _{CC}	LOG[FD] _{MR} (mean)	N _{MR}	F pr	LSD	Comment
FD-C	2008+2009 ^a	Sp	0.349	2	1.291	2	0.549	5.681	N.S.
	2008 ^b	Es	3.255	3	3.271	4	0.962	0.826	N.S.
FD-C+BN	2008 ^b	Es	2.942	1	3.536	3	0.386	2.324	N.S.

^a, ^b: Results allowing groupings as reported in Table 7 at corresponding symbols

Figure 1
205x181mm (300 x 300 DPI)