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

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1	FDP-titre in grapevines with different susceptibility to GYs
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3	Flavescence dorée phytoplasma titre in field-infected Barbera and Nebbiolo grapevines
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- 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

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

41 sequence into more than 30 groups and within the 'Candidatus Phytoplasma' genus (Firrao et al., 42 2004), (Zhao et al., 2010). 43 Flavescence dorée (FD) is an epidemic, economically important, quarantine disease of grapevine in 44 France, Italy and Spain (Boudon-Padieu, 2003). The disease is associated with a phytoplasma 45 belonging to the 16SrV (Elm yellows) taxonomic group (Lee et al., 2000). FD epidemics are caused 46 by phytoplasma strains belonging to three phylogenetic strain clusters, according to multilocus 47 sequence analysis of map, uvrB-degV and secY loci (Arnaud et al., 2007). In Italy the disease is 48 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), 49 50 although isolates of the less variable cluster 2 (FD-D) are also present. The leafhopper Scaphoideus 51 titanus Ball is the specific vector of the different FDP strains to grapevine under natural conditions 52 (Mori et al., 2002, Papura et al., 2009, Schvester et al., 1963). Recently, a role as potential vector of 53 FDP from alternative hosts to grapevine has been suggested for the planthopper Dictyophara 54 europea (L.) (Filippin et al., 2009). 55 FD-infected grapevines usually show symptoms the year after inoculation (Morone et al., 2007), 56 although longer latencies have been reported (Osler et al., 2002). Yellowing, downward curling of 57 leaves, fruit abortion, stunting, and lack of lignification of new shoots are among the most important 58 symptoms (Caudwell, 1983, Caudwell, 1990). Infected grapes dramatically reduce their production, 59 but following the first year of symptom expression, a spontaneous and cultivar-dependent remission 60 of symptoms may occur, and asymptomatic plants usually do not contain detectable FD 61 phytoplasma (Morone et al., 2007). 62 Control of the disease relies mainly on compulsory insecticide treatments to reduce vector 63 population and roguing of infected plants. No resistance is available to FD, but cultivars with 64 different susceptibility to the disease have been reported (Kuzmanovic et al., 2008). Barbera and 65 Nebbiolo are two traditional and economically important grapevine cultivars of Piemonte for the 66 production of red wines. Barbera is highly susceptible to FD and shows severe symptoms starting

67	with growth reduction at the beginning of the vegetative season, while symptoms on Nebbiolo vines
68	are milder, especially at the beginning of the vegetative season (Morone et al., 2001).
69	In this study we describe a new approach to real time quantification of FD phytoplasma in
70	grapevine which was used to determine a) relationships between cultivar susceptibility and
71	pathogen concentration in the plant, b) relationships between pathogen presence/concentration and
72	symptom severity, c) FD phytoplasma titre over the vegetative season in order to identify the best
73	timing for sampling and detection.
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76	Materials and methods
77	
78	Vineyard description
79	Two vineyards located in Piemonte (north-western Italy) were chosen: Cocconato (Asti province)
80	and Monteu Roero (Cuneo province). Barbera and Nebbiolo cvs are planted at the Cocconato
81	vineyard in 1999 on SO4 rootstock. Nebbiolo cv was planted at Monteu Roero vineyard in 1998
82	also on SO4 rootstock. Both vineyards were subjected to conventional agronomic and phytosanitary
83	practices. An electronic meteorological station was present in Cocconato between the neighbouring
84	Barbera and Nebbiolo vineyards. For the Monteu Roero vineyard, climatic data were collected at a
85	station located in a neighbouring municipality (Canale) in the same exposure and altitude
86	conditions. Mean month temperature (°C) and number of days in each month with minimum
87	temperature below 0°C (freezing days) were calculated from data collected at both stations.
88	
89	Establishment of a correct sampling procedure
90	To establish a correct sampling procedure, five FD-infected Barbera and Nebbiolo plants were
91	selected in the Cocconato vineyard at the growth stage $17 - 19$ (12 to 16 leaves separated; Coombe,
92	1995). Symptomatic and asymptomatic shoots were separately labeled on each plant and samples
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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 %.	
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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 $\mu$ l 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 $1 \times TBE$ 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  $\mu$ L 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, corresponding to 1.17·10<sup>8</sup>, 1.17·10<sup>6</sup>, 1.17·10<sup>3</sup> cells of FDP. For the absolute quantification of 168 grapevine DNA, three quantities of total healthy grapevine DNA (5 ng, 0.05 ng, and 0.005 ng) were 169 7

170	used. Grapevine sample DNAs were diluted in SDW to a final concentration of 1 $ng/\mu l$ , and used (5
171	$\mu$ l) as template in real time assays together with Platinum SYBR Green qPCR Supermix UDG (Life
172	Technologies, Carlsbad, CA, USA) and specific primer pairs at a final concentration of 300 nM, in
173	a volume of 25 $\mu$ L. The PCR was performed in 96-well plates in a BioRad iCycler (BioRad).
174	Cycling conditions were as follows: one cycle at 95°C for 5 min, then 40 cycles at 95°C for 15 s,
175	and 61°C for 1 min. For each sample, plant- and FDP-DNAs were amplified in the same plate with
176	their corresponding reaction mixture, together with the three standard dilutions of healthy host DNA
177	(Hg) and p26SecYFD. Each sample was run in triplicate in the same plate. For each PCR, water
178	instead of DNA was used as negative control. Threshold levels, threshold cycles, and standard
179	curves were automatically calculated by the BioRad iCycler software, version 3.06070. Per-well
180	baseline cycles were determined automatically. Specificity of the reaction was tested by running a
181	melting curve analysis of the amplicons following each quantification reaction.
182	
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182	Data analyses
	Data analyses For the analyses of phytoplasma concentration, raw data (expressed as log values) were used and
183	
183 184	For the analyses of phytoplasma concentration, raw data (expressed as log values) were used and
183 184 185	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
183 184 185 186	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).
183 184 185 186 187	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 11 <sup>th</sup> edition rel. 11.1
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183 184 185 186 187 188 189	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 11 <sup>th</sup> 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

- 193
- 194 **Results**
- 195

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, Iin 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 Cocconato, respectively), and 13 in 2009
214	(9 from Monteu Roero and 4 from Cocconato). For early summer surveys, 4in 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 Cocconato), and 10 in 2009 (7
218	from Monteu Roero and 3 from Cocconato). For late summer surveys, 4in 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	2009, respectively. 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' TAATAATGATGGGGGATTCAA
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
	11

273	bp) and gel electrophoresis confirmed the absence of nonspecific products (not shown). Melting	
274	curve analysis of the amplicon showed a unique melting peak at 85.5°C (Fig. 1, Panel D). Serial	
275	dilutions of FDP plasmid DNA ranging from $21.13 \cdot 17 \times 10^7 \cdot 10^8$ to $21.13 \cdot 17 \times 10^3 \cdot 10^3 \cdot$	Formatted: Supersc
276	were tested in triplicate and the CT values were plotted against the copy number. Serial dilution of	
277	healthy grapevine DNA ranging from 50 ng to 0.05 ng were also tested in triplicate and the CT	
278	values were plotted against the DNA concentration. The linear correlation coefficient $(r)$ between	
279	the CT and the logarithm of the DNA copy number was repeatedly greater than 0.995 for the	
280	amplification of both targets (Fig. 1, panels A and C). PCR efficiencies of both curves were	
281	constantly around 108% and 103% (plant and phytoplasma, respectively).	
282		
283	Establishment of a correct sampling procedure	
284	Results of this preliminary experiment are presented in Table 4. FD phytoplasma was detected by	
285	nested PCR in all samples from Barbera and Nebbiolo symptomatic shoots, confirming the infected	
286	status of the analyzed plants. FD phytoplasma was also detected in three samples from	
287	asymptomatic shoots of two Barbera plants, while it was never detected in asymptomatic shoots of	
288	infected Nebbiolo plants. FD phytoplasma titre was measured in all samples from symptomatic	
289	shoots of Barbera and in two nested PCR positive sample from the two symptomatic shoots of one	
290	Nebbiolo plant. FD phytoplasma titre in symptomatic shoots of the five Barbera plants ranged from	
291	about 39 to 1180 cells/ng grapevine DNA, and it was above the quantification threshold in one of	
292	the three nested PCR positive samples from asymptomatic Barbera shoots. In Barbera, the variance	
293	among FDP titre within shoots of each plant was lower than that among plants (Table 5), therefore,	
294	collection of leaves from one symptomatic shoot per plant was set as sampling protocol for	
295	successive quantification experiments.	
296		

297 Flavescence dorée phytoplasma concentration in Barbera and Nebbiolo grapevines

298	Despite the low number of FD-D phytoplasma infected plants over the two years, tests indicated
299	that no significant differences were recorded in the concentration of FD-C and -D phytoplasmas
300	within each sampling of the different cvs (Table 6). Further tests also showed that the presence of
301	BNP had no effect on FDP concentration (Table 7). FDP concentration in Barbera grapevines
302	sampled in spring, early and late summer of 2008 and 2009 are illustrated in Fig. 2. Mean FDP
303	concentration in Barbera samples was 242 cells/ng grapevine DNA in spring 2008, and 2 cells/ng
304	grapevine DNA in spring 2009. In early summer samplings, FDP titre was 8,856 and 8,735 cells/ng
305	grapevine DNA in 2008 and in 2009, respectively; in late summer, FDP concentration was 6,420
306	cells/ng grapevine DNA in 2008 and 671 in 2009. Due to significant differences in the variance,
307	FDP titres in spring and early summer samplings were analyzed separately from those of late
308	summer. Significant differences were recorded in FDP titre between sampling seasons and years.
309	FDP concentration was consistently higher in 2008 than in 2009 (3.299 vs 2.456 DLog; $N1 = 31$ ;
310	N2 = 10; S.E.D. = 0.2799; least significant difference at 0.05 level = 0.5672), and the difference
311	was mainly due to phytoplasma titre in spring, since differences between FDP titre in 2008 and
312	2009 early summers were not significant (least significant difference at 0.05 level between two
313	years: spring = $0.0618$ ; early summer = $0.6108$ ). FDP concentration was also significantly higher in
314	2008 late summer sampling compared to 2009 ( $P$ = 0.072). A seasonal trend in FDP concentration
315	(low in spring, high in early summer and intermediate in late summer) was conserved in Barbera
316	vines sampled in both years (Fig. 2).
317	FDP titre in Nebbiolo vines (Fig. 2) was significantly different according to year (2008 and 2009)
318	and sampling season (spring, early and late summer), while it was similar in both vineyards (Table
319	8). Mean FDP titre in Nebbiolo vines was 134 cells/ng grapevine DNA in spring samples, 1,417 in
320	early summer samples and 489 in late summer samples of 2008. In 2009, mean FDP titre in spring
321	samples was 2 cells/ng grapevine DNA, in early summer 545 cells/ng grapevine DNA and in late
322	summer samples 334. All data were then pooled to analyze differences between years and sampling
323	seasons. Overall, in 2008, FDP in Nebbiolo was more abundant than in 2009 (2.837 vs 2.310 DLog; 13

324	N1 = 32; $N2 = 26$ ; 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
337 338	<i>Flavescence dorée phytoplasma concentration and symptom severity</i> Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples
338	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples
338 339	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples showed a significant, positive correlation between FDP concentration and symptoms severity
<ul><li>338</li><li>339</li><li>340</li></ul>	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples showed a significant, positive correlation between FDP concentration and symptoms severity ( $r=0.412$ ; S.E.=0.367; N=25; DF=23; t=1.89; P= 0.071Spearman's correlation coefficient r=0.412;
<ul><li>338</li><li>339</li><li>340</li><li>341</li></ul>	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples showed a significant, positive correlation between FDP concentration and symptoms severity ( $r=0.412$ ; S.E.=0.367; N=25; DF=23; t=1.89; P= 0.071Spearman's correlation coefficient r=0.412; adjusted for ties = 0.367; N=25; DF=23; t=1.89; P= 0.071), and so did the Barbera samples of 2008
<ul> <li>338</li> <li>339</li> <li>340</li> <li>341</li> <li>342</li> </ul>	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples showed a significant, positive correlation between FDP concentration and symptoms severity ( $r=0.412$ ; S.E.=0.367; N=25; DF=23; t=1.89; P= 0.071Spearman's correlation coefficient r=0.412; adjusted for ties = 0.367; N=25; DF=23; t=1.89; P= 0.071), and so did the Barbera samples of 2008 (Spearman's correlation coefficient r=0.389; N=25; DF=23; t approximation 1.72; P= 0.099), but
<ul> <li>338</li> <li>339</li> <li>340</li> <li>341</li> <li>342</li> <li>343</li> </ul>	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples showed a significant, positive correlation between FDP concentration and symptoms severity ( $r=0.412$ ; S.E.=0.367; N=25; DF=23; t=1.89; P= 0.071Spearman's correlation coefficient r=0.412; adjusted for ties = 0.367; N=25; DF=23; t=1.89; P= 0.071), and so did the Barbera samples of 2008 (Spearman's correlation coefficient r=0.389; N=25; DF=23; t approximation 1.72; P= 0.099), but not in 2009, pooling both early and late summer samplings (r=0.389; S.E.=0.377; N=25; DF=23;
<ul> <li>338</li> <li>339</li> <li>340</li> <li>341</li> <li>342</li> <li>343</li> <li>344</li> </ul>	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples showed a significant, positive correlation between FDP concentration and symptoms severity $(r=0.412; S.E.=0.367; N=25; DF=23; t=1.89; P=0.071Spearman's correlation coefficient r=0.412; adjusted for ties = 0.367; N=25; DF=23; t=1.89; P=0.071), and so did the Barbera samples of 2008 (Spearman's correlation coefficient r=0.389; N=25; DF=23; t approximation 1.72; P=0.099), but not in 2009, pooling both early and late summer samplings (r=0.389; S.E.=0.377; N=25; DF=23; t=1.72; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; SPE-23; P=0.091), P=0.091), P=0.091, P=0.091,$
<ul> <li>338</li> <li>339</li> <li>340</li> <li>341</li> <li>342</li> <li>343</li> <li>344</li> <li>345</li> </ul>	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples showed a significant, positive correlation between FDP concentration and symptoms severity ( $r=0.412$ ; S.E.=0.367; N=25; DF=23; t=1.89; <i>P</i> = 0.071Spearman's correlation coefficient r=0.412; adjusted for ties = 0.367; N=25; DF=23; t=1.89; P= 0.071), and so did the Barbera samples of 2008 (Spearman's correlation coefficient r=0.389; N=25; DF=23; t approximation 1.72; P= 0.099), but not in 2009, pooling both early and late summer samplings ( $r=0.389$ ; S.E.=0.377; N=25; DF=23; t=1.72; <i>P</i> = 0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; adjusted for ties r = 0.373; N=22; DF=20; t=1.98; P=0.087), but not in 2008, pooling both early and
<ul> <li>338</li> <li>339</li> <li>340</li> <li>341</li> <li>342</li> <li>343</li> <li>344</li> <li>345</li> <li>346</li> </ul>	Considering both Barbera and Nebbiolo and both years from both vineyards, the spring samples showed a significant, positive correlation between FDP concentration and symptoms severity ( $r=0.412$ ; S.E.=0.367; N=25; DF=23; t=1.89; P= 0.071Spearman's correlation coefficient r=0.412; adjusted for ties = 0.367; N=25; DF=23; t=1.89; P= 0.071), and so did the Barbera samples of 2008 (Spearman's correlation coefficient r=0.389; N=25; DF=23; t approximation 1.72; P= 0.099), but not in 2009, pooling both early and late summer samplings (r=0.389; S.E.=0.377; N=25; DF=23; t=1.72; P= 0.099), and the Nebbiolo samples of 2009 (Spearman's correlation coefficient r=0.430; adjusted for ties r = 0.373; N=22; DF=20; t=1.98; P=0.087), but not in 2008, pooling both early and late summer samplings (r=0.389; S.E.=0.307; N=24; DF=20; t=1.98; P=0.087), but not in 2008, pooling both early and late summer samplings (r=0.389; DF=21; t=1.98; P=0.061). Early and late

350	Overall, there was no highly significant correlation between FDP concentration and symptoms
351	severity.
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354	Discussion
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356	Flavescence dorée infection in 2008 and 2009 in the two vineyards of Cocconato and Monteu Roero
357	was caused mainly by Flavescence dorée type C phytoplasma (FD-C). This isolate is stably present
358	in the area since the early outbreak of the epidemics in 1998, and it is the most represented one in
359	FD-infected vines of Piemonte region (Marzachi et al., 2001). FD-D type phytoplasmas were
360	detected sporadically in infected Barbera and Nebbiolo grapevines collected in spring, early and late
361	summer of both years. Grapevines showing early symptoms were mostly infected with FD-C
362	phytoplasma, as reported in other grapevine-growing areas (Angelini et al., 2006), and this isolate
363	was detected throughout the vegetative season. Bois noir disease is also present in Piemonte
364	vineyards (Marzachi et al., 2001), and BNP was found only in mixed infections with FDP in
365	samples from the two vineyards. As we only sampled symptomatic grapevines in the two highly
366	FD-infected locations, the presence of BNP in asymptomatic plants cannot be ruled out.
367	The FDP type ( C or D) had no effect on symptom severity of the infected vines of both cultivars.
368	Also, the presence of BNP in mixed infection, a rare occurrence, had no effect on symptom
369	expression. Under these experimental conditions, a competition effect between BNP and FDP
370	eannot be ruled out .
371	For two consecutive years, infected Barbera vines showed more severe spring symptoms than
372	Nebbiolo ones. This observation coincides with the fact that Barbera is known as a highly FD-
373	sensitive cv. Both cvs showed more severe symptoms in 2009 than in 2008, and in 2009 lower
374	winter and spring mean temperatures and more freezing days were recorded at both localities than
375	in 2008. Low spring temperatures have an immediate effect on grapevine in delaying bud burst
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376	(Tomasi et al., 2007), but they also have a long term effect in reducing fruit set (%) and berry
377	number per bunch in a cv-specific manner (Ebadi et al., 1995). Temperature stresses also affect
378	Ca2 <sup>+</sup> homeostasis in grapevine (Wang et al., 2004). Also, low temperatures reduce respiration rate
379	in grapevine (Franck et al., 2011), thus reducing the production of H <sub>2</sub> O <sub>2</sub> (Foyer et al., 2009), a
380	molecule probably involved in defense response pathways to phytoplasmas (Musetti et al., 2007).
381	We can speculate that more severe FD symptoms in both cvs in early and late summer assessment
382	of 2009 may be an effect of a long-lasting stress of the plant caused by the occurrence of low
383	winter/spring temperatures in that year. Moreover, in 2008, temperature probes recorded lower
384	winter/spring mean temperatures and more freezing days in the Monteu Roero vineyard compared
385	to Cocconato, and significantly more severe early summer symptoms were recorded in Nebbiolo
386	infected vines in Monteu Roero than in Cocconato.
387	Quantification of FDP-DNA was obtained as the ratio of phytoplasma DNA per ng of grapevine
388	DNA as previously described for "Ca. P. asteris" (Marzachi & Bosco, 2005). Real time based PCR
389	assays for diagnosis and characterization of several phytoplasmas, including FDP, have been
390	proposed (Angelini et al., 2007, Bianco et al., 2004, Galetto et al., 2005, Hren et al., 2007, Mehle et
391	al., 2013, Pelletier et al., 2009), but this is one of the first attempts, together with the one of Prezelj
392	and co-workers (2012), to quantify FDP in field-infected vines of different cvs. A double absolute
393	quantification assay was developed and only samples with a threshold cycle value within that of the
394	lowest standard concentration were considered. The quantification of less than 3 phytoplasma cells
395	per ng plant DNA in three spring samples can still be considered reliable according to the MIQE
396	guidelines (Bustin et al., 2009) as it exceeds by three times the most sensitive limit of detection.
397	It is known that phytoplasmas have uneven distribution in the grapevine canopy, but a direct
398	correlation between the presence of phytoplasma specific symptoms and FDP detection was clearly
399	shown as all samples collected from symptomatic shoots of both cvs were positive for the presence
400	of FDP, while all samples from asymptomatic Nebbiolo and most from asymptomatic Barbera were
401	nested PCR negative. Similar results are reported for other grapevine varieties (Prezelj et al., 2012)
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402	where FDP was mainly associated with symptomatic tissues, although it was also detected in some
403	symptomless tissues of highly infected (and symptomatic) plants. Also, the differences between
404	FDP titre in different symptomatic shoots from the same Barbera plant were lower than the
405	differences among plants, therefore confirming that sampling from one symptomatic shoot can
406	provide a reliable and representative evaluation of FDP titre in the infected parts of the plant.
407	We can suggest that FD-C and D phytoplasmas multiply at the same level in grapevine, and that
408	the presence of BNP has no influence on the accumulation of FDP in doubly infected plants. Very
409	few quantitative information on plants co-infected by two phytoplasmas are available, however,
410	mild and severe isolates of "Ca. P. asteris" show different multiplication rate and movement in the
411	herbaceous host (Sinha, 1983) and, in the case of Ash yellows phytoplasma, strains that differ in
412	aggressiveness have a different multiplication and/or movement following co-inoculation to the
413	laboratory host periwinkle (Sinclair & Griffiths, 2000). In Apple proliferation (AP) infected apple
414	trees, the ratio of mild to severe strains in the phytoplasma population determines the virulence of
415	the infection and shifts in the population composition may alter virulence of multiple strain
416	accession (Seemueller et al., 2011).
417	In both cvs and in both years, the lowest FDP titre was observed in spring, the highest in early
418	summer and an intermediate one at the end of the vegetative season. A continuous increase of FDP
419	titre up to the end of the vegetative season was reported for other grapevine cvs in Slovenia (Prezelj
420	et al., 2012). Seasonal fluctuations of phytoplasma populations in AP-infected apple trees have been
421	measured with highest levels occurring from December to May (Baric et al., 2011). Despite the
422	occurrence of more severe symptoms throughout the vegetative season of 2009 compared to 2008,
423	at each sampling date, FDP titre was significantly higher in 2008 than in 2009, irrespective of the
424	cv. This difference was most evident in the spring samplings, when as many as hundreds FDP cells
425	were recorded in 2008 in contrast to only as few as 2 FDP cell per ng plant DNA in 2009. Since the
426	lowest FDP titre was consistently measured in both cvs, both years and at both sites in spring,
427	winter temperature together with plant dormancy and pruning, that eliminates most of the infected 17

428	plant parts, have a cooperative effect in reducing phytoplasma titre in the plant. Multiplication of
429	pathogen resumed with the vegetative season and the differences in FDP titres between the two
430	years were reduced, although FDP-specific symptom expression was more severe in 2009 than in
431	2008. We can speculate that colder winter/early spring climatic conditions have a long-lasting stress
432	effect on the grapevines, increasing both their sensitivity to FD and symptom severity even in the
433	presence of a low pathogen load. Also, the hypothesis that the pathogen may multiply less
434	efficiently in the stressed plant, therefore reaching a low titre despite the severe symptoms, cannot
435	be ruled out.
436	FDP titre in Barbera grapevines was always higher than in Nebbiolo. This is in good correlation
437	with more severe symptoms expressed by FD-infected Barbera vines compared to Nebbiolo ones
438	and provides the most likely explanation for the milder symptoms shown by FDP-infected
439	Nebbiolo. Actually this latter cv sustains a low multiplication of the pathogen. Proteomic and
440	transcriptomic analyses of FDP-infected Nebbiolo plants show that most proteins modulated during
441	infection belong to the "cell rescue, defense and virulence" class (Margaria & Palmano, 2011a).
442	On the other hand, preliminary studies on Barbera cv, show a lower presence of proteins of the
443	'defense' category compared to the total identified proteins (Margaria & Palmano, 2011b). Perhaps
444	different responses of the two cvs to FDP may account for the different multiplication level of the
445	pathogen, but this hypothesis remains untested.
446	The results indicate that the best time for sampling and detection of FDP infection is early summer,
447	when phytoplasma titre is highest. However, our work, together with the one of Prezelj and co-
448	workers (2012) provide evidence that FDP can be detected as early as at flowering season, or even
449	before in the case of Barbera in our work. Quantitative real time PCR is a useful tool for the
450	evaluation of phytoplasma multiplication and can be used when screening for resistance/tolerance.
451	We have demonstrated that Barbera supports a higher FDP multiplication compared to Nebbiolo,
452	and further studies are required to understand whether this has an effect on vector acquisition and

453	transmission efficiency, as already suggested for Pinot blanc and Merlot cultivars (Bressan et al.,
454	2005).
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## 627 Captions for figures

- 628
- 629 Figure 1. A: Standard curve of Flavescence dorée phytoplasma DNA obtained by plotting threshold
- 630 cycles (CT) values vs log of  $2 \times 10^7$ ,  $2 \times 10^5$ ,  $2 \times 10^3$ , and  $2 \times 10$  copy number of p26SecYFD
- 631 plasmid. <u>BC</u>: Melting curve analysis of the amplicon obtained with primers FdSecyFw/FdSecyRv.
- 632 CB: 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 dorèe phytoplasma concentration (cells / ng grapevine DNA) and

- , led i. .coero (MR) v. 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.
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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	С	D	A	B	С	D
2008	Barbera	1	9	5	2	9	5	0	2	4	4	0
2000	Total plants		15		16				10			
	Barbera	2	1	0	0	1	2	5	0	1	1	5
2009	Total plants	3 8						7				
2008	Nebbiolo	2	1	2	4	6	5	0	3	6	5	0
2008	Total plants	5			15			14				
2000	Nebbiolo	2	1	0	0	2	4	7	0	2	3	5
2009	Total plants		3			1	3			1	0	

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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 <sup>a</sup>	U <sup>b</sup>	$P^{c}$		
Sp <u>ring</u> ⁴	С		2008+2009	B > N	17, 3	3.0	0.991
		В		2009 > 2008	15, 8	14.5	0.999
	С	Ν		2009 > 2008	4, 4	0.0	0.986
E <u>arly</u> s <u>ummer</u>			2008	B > N	15, 4	2.0	0.999
e	MR	Ν		2009 > 2008	11, 9	23.5	0.975
	MK	Ν	2008	MR > C	4, 11	0.0	0.999
	С	В		2009 > 2008	9, 7	7.5	0.996
L <u>ate</u>	MR	Ν		2009 > 2008	8, 7	9.0	0.990
s <u>ummer</u> ‡	С	N	2008	Es <sup>e</sup> < Ls <sup>f</sup>	4, 4	0.0	0.986

<sup>a</sup> N: sample numbers; <sup>b</sup> U: value of U in the Mann-Whitney U test; <sup>c</sup> 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); <sup>d</sup>-Sp: spring; <sup>e</sup>-Es: early summer; <sup>f</sup>-Ls: late summer.

;<sup>d</sup>Sp: spring; <sup>e</sup>Es: early summer; <sup>f</sup>Ls: late summer.

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				
			(Positiv	/e/ total sam	pled)	(Positive/total sampled)				
		0	Sp	Es	Ls	Sp	Es	Ls		
		В	14/15	15/16	8/10	1/15	1/16	2/10		
	Cocconato	Ν	0	3/4	2/4	1/1	1/4	2/4		
2008			G							
	Monteu Roero	Ν	1/4	7/11	9/10	3/4	4/11	1/10		
					),					
		В	3/3	8/8	6/7	0	0/8	1/7		
2000	Cocconato	Ν	2/2	4/4	3/3	0	0	0		
2009		N	1 /1	<u>8/0</u>		0	1./0	1 /7		
	Monteu Roero	Ν	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 (<u>P1/P7</u> primers) and nested (<u>R16(V)F1/R1 primers</u>) 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.

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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).

2 2.1135 0.07722 3 2.850333 0.09827 4 2.134833 0.09812 5 2.7865 0.00098 General,	Plant	Mean FDP titer in shoots (DLog)	Var
2       2.1135       0.07722         3       2.850333       0.09827         4       2.134833       0.09812         5       2.7865       0.00098         General, among       2.412667       0.16141	1	2 178167	0.07131606
3       2.850333       0.09827         4       2.134833       0.09812         5       2.7865       0.00098         General, among       2.412667       0.16141			0.0772245
4 2.134833 0.09812 5 2.7865 0.00098 General, among 2.412667 0.16141			0.09827222
General, among 2.412667 0.16141			0.0981245
among 2.412667 0.16141	5	2.7865	0.00098272
among 2.412667 0.16141			
6			
plants		2.412667	0.16141599
	plants		

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: Cocconato 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<sub>C</sub>, N<sub>D</sub>: 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 <sub>C</sub>	LOG[FD-D] (mean)	ND	F pr	LSD	Comment
			Es	4.0242	14	3.640	1	0.43 6	1.034	N.S.
В	С	2008	Ls	3.8911	7	3.785	1	0.71 4	0.6746	N. S.
		2008	Es	3.2713	4	2.7967	3	0.20 6	0.8409	N.S.
Ν	N MR	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. Log[FD]<sub>FD</sub>, Log[FD]<sub>FD+BN</sub>: Log of FD phytoplasma concentration. N<sub>FD</sub>, N<sub>FD+BN</sub>: 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	LOG[FD] <sub>FD</sub> (mean)	N <sub>FD</sub>	LOG[FD] <sub>FD +BN</sub> (mean)	N <sub>FD+BN</sub>	F pr	LSD	Comment
		• • • • •	Sp	2.384	14	2.370	1	0.989	2.204	N.S.
В	С	2008	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.
	C+MR <sup>a</sup>	2008+2009	Sp	0.820	4	2.366	2	0.165	2.531	N.S.
Ν	C+MR <sup>b</sup>	2008	Es	3.2644	7	3.3878	4	0.651	0.595 5	N.S.

2.04

<sup>a</sup>, <sup>b</sup>: 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]<sub>CC</sub>, LOG[FD-D]<sub>MR</sub>: 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] <sub>CC</sub> (mean)	N <sub>CC</sub>	LOG[FD] <sub>MR</sub> (mean)	N <sub>MR</sub>	F pr	LSD	Comment
FD-C	2008+2009 <sup>a</sup>	Sp	0.349	2	1.291	2	0.549	5.681	N.S.
гD-С	2008 <sup>b</sup>	Es	3.255	3	3.271	4	0.962	0.826	N.S.
FD-C+BN	2008 <sup>b</sup>	Es	2.942	1	3.536	3	0.386	2.324	N.S.

<sup>a</sup>, <sup>b</sup>: Results allowing groupings as reported in Table 7 at corresponding symbols

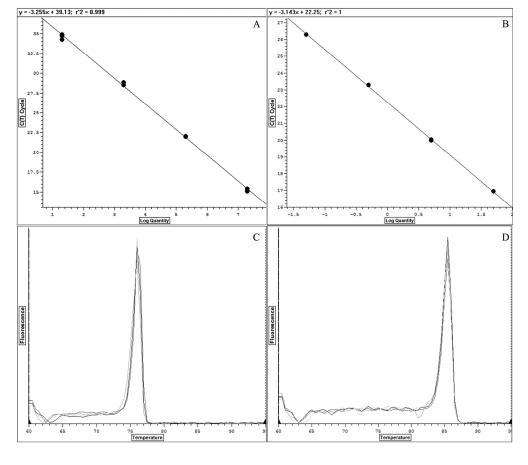


Figure 1 205x181mm (300 x 300 DPI)

