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#### Chitosan-elicited defense responses in Cucumber mosaic virus (CMV)-infected tomato plants

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18	
19	Abbreviations: A, photosynthetic activity; ANOVA, analysis of variance; AR156, Bacillus cereus AR156;
20	CERK1, chitin elicitor receptor kinase 1; CHT, chitosan; CMV, Cucumber mosaic virus; DAS-ELISA,
21	double-antibody sandwich enzyme-linked immunosorbent assay; ET, ethylene; $F_v/F_m$ , maximum quantum
22	yield of PSII; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; $g_s$ , stomatal conductance to water
23	vapor; ISR, induced systemic resistance; JA, jasmonic acid; NPR1, non-expressor of pathogenesis-related
24	genes 1;  \$\phiPSII\$, quantum yield of PSII; PAL, phenylalanine ammonia lyase; PAR, photosynthetically active
25	radiation; PR, pathogenesis-related; PSY, phytoene synthase; PVX, Potato virus X; qPCR, Real-Time
26	quantitative PCR; ROS, reactive oxygen species; SA, salicylic acid; SAR, systemic acquired resistance;
27	TMV, Tobacco mosaic virus; TP, treated plants; TUB, tubulin; UK, uridylate kinase
28	
29	ABSTRACT
30	
21	The control of plant diseases by inducing plant resistance responses represents an interesting
1	The control of plant diseases by inducing plant resistance responses represents an interesting
32	solution to avoid yield losses and protect the natural environment. Hence, the intertwined

relationships between host, pathogen and inducer are increasingly subject of investigations. Here, 

34 we report the efficacy of chitosan-elicited defense responses in Solanum lycopersicum var. cerasiforme plants against Cucumber mosaic virus (CMV). Chitosan was applied via foliar spray 35 before the CMV inoculation to verify its effectiveness as a preventive treatment against the viral 36 infection. Virus accumulation, photosynthetic performance, as well as genes encoding for proteins 37 affecting resistance responses and biosynthetic pathways, were investigated. It was observed a 38 significant reduction of CMV accumulation in chitosan-treated plants that were successively 39 infected with CMV, compared to only CMV-infected ones (up to 100%). Similarly, a positive effect 40 of chitosan on gas exchange dynamics was revealed. The analysis of gene expression (CEVI-1, 41 NPR1, PSY2 and PAL5) suggested the occurrence of chitosan-induced, systemic acquired 42 resistance-related responses associated with a readjustment of the plant's oxidative status. In 43 44 addition, the absence of deleterious symptoms in chitosan-treated successively CMV-infected 45 plants, confirmed that chitosan can be used as a powerful control agent. Our data indicate that chitosan, when preventively applied, is able to elicit defense responses in tomato to control CMV 46 infection. Such finding may be recommended to protect the tomato fruit yields as well as other 47 48 crops.

49

Keywords: Chitosan; *Cucumber mosaic virus*; Defense/antioxidant-related protein; Disease
control; Photosynthetic performance; *Solanum lycopersicum* var. *cerasiforme*.

52

### 53 **1 Introduction**

54

55 Plants are susceptible to numerous pathogens responsible for diseases that can reduce crop yield

56 (e.g., (Vitti et al., 2015)), causing severe economic losses. Among the most dangerous

57 phytopathogens, viruses cannot be faced by using specific agrochemicals (Iriti and Varoni, 2015).

Hence, control of viral infections, protecting the natural environment, is the best strategy to ensure
an, at least, satisfactory harvest. In such a scenario, elicitors triggering plant defense responses and
inducing a systemic resistance state can represent a solution.

Chitosan (CHT) is a polycationic heteropolysaccharide composed by N-acetyl-D-glucosamine and 61 D-glucosamine linked by  $\beta$ -(1 $\rightarrow$ 4) glycosidic bonds (Iriti and Varoni, 2015). Although CHT is a 62 natural compound (present in Zygomycetes), it is mainly obtained by the deacetylation of chitin, a 63 component of the fungal cell wall and the arthropod exoskeleton (Iriti and Varoni, 2015). Chitosan 64 polymers may vary in molecular weight, viscosity, pKa value, polymerization and deacetylation 65 degree, affecting their physicochemical and biological properties (Iriti and Varoni, 2015). Chitosan 66 is a biodegradable and nontoxic compound inducing systemic acquired resistance (SAR) to 67 pathogens in plants (Xing et al., 2015). This compound also exhibits a direct antimicrobial activity, 68 69 likely mainly through electrostatic interactions (Xing et al., 2015). Chitosan bioactivities can be explained because the polycationic nature of CHT leads to affinity with the anionic contents of 70 target organism (Kumaraswamy et al., 2018). More specifically, as an antimicrobial, CHT 71 interaction with cell wall and cell membranes can destabilize them. In addition, its interaction with 72 DNA and proteins can interfere with the transcription and translation mechanisms (Kumaraswamy 73 et al., 2018). Furthermore, CHT can chelate essential nutrients, trace elements and metal ions that 74 are necessary for the microbial growth, as well as it can form a polymer film that compromises 75 metabolite excretion and nutrient uptake (Xing et al., 2015). However, the detailed mechanism of 76 77 action of CHT in reducing plant diseases has not been completely revealed (Hassan and Chang, 2017). Chitosan perception by plant is shortly followed by a vatiation in the ion fluxes and 78 membrane depolarization (Iriti and Varoni, 2015). Therefore, CHT can be recognized by plant as a 79 pathogen-mimicking stimulus, but the identification of a CHT receptor is still doubtful (Malerba 80 and Cerana, 2016; Povero et al., 2011). Although Petutschnig et al. (2010) found that the chitin 81 elicitor receptor kinase 1 (CERK1) also bound more weakly to CHT, later Povero et al. (2011) 82

83 demonstrated that the perception of CHT was independent of CERK1. Recently, Liu et al. (2018) suggested wheat W5G2U8, W5HY42, and W5I0R4 as potential chitosan oligosaccharides receptors. 84 Interestingly, the application of CHT to promote the plant growth has also been studied 85 (Kumaraswamy et al., 2018; Sharif et al., 2018) and CHT nanoparticles can also be used to deliver 86 pesticides, herbicides, fertilizers, micronutrients as well as genetic material (Malerba and Cerana, 87 2016). The antimicrobial activity of CHT has been studied for bacteria, yeasts and moulds (Liu et 88 al., 2004). Furthermore, CHT is effective as an alternative treatment to conventional fungicides 89 aimed to control the postharvest decay, both after preharvest (Feliziani et al., 2015) and postharvest 90 applications. The latter is associated with CHT coating of fruits (Sivakumar et al., 2016). On cherry 91 92 tomato fruit, CHT exerted an inhibitive action on the gray mold (*Botrytis cinerea*), presumably 93 involving the mitogen-activated protein kinase signaling pathway and determining an increase of hydrogen peroxide, peroxidase activity, as well as PR1a1 and PR5 transcripts (Zhang et al., 2015). 94 Chitosan-induced responses against viruses, such as *Potato virus X* (PVX) and *Tobacco mosaic* 95 virus (TMV), were also evaluated (Chirkov et al., 2001; Jia et al., 2016; Nagorskava et al., 2014). 96 Cucumber mosaic virus (CMV) (genus Cucumovirus, family Bromoviridae) presents polyhedral 97 virions composed of 180 subunits (T=3 icosahedral symmetry) (Gallitelli, 1998). Virus particles are 98 isometric: separate particles separately contain RNA1 and RNA2, a third particle contains RNA3 99 100 and subgenomic RNA4 (Agrios, 1997) and possibly RNA3 and subgenomic RNA4A (Gallitelli, 2000). RNA1 and RNA2 code for two different proteins involved in RNA replication (Agrios, 101 1997). The 2b protein is translated from the RNA4A of RNA2 (Gallitelli, 2000). RNA3 encodes a 102 protein involved in virus movement and contains the open reading frame for the coat protein. The 103 coat protein cistron is translated via the RNA4 (Gallitelli, 2000). In order to infect plants, the co-104 infection with the three particles together is required (Gallitelli, 1998). The numerous strains 105 belonging to CMV differ in properties and characteristics such as host plants, symptoms produced, 106 ways of transmission (Agrios, 1997). Infecting at least 100 plant families and 1,200 species 107

108 (Edwardson and Christie, 1991), CMV has a wide range of hosts, such as ornamentals and many species of vegetables (Agrios, 1997). Furthermore, CMV causes distortion and discoloration of 109 leaves, fruits and flowers, reduction in quantity and quality of crop yield, up to reduced growth and 110 plant death (Agrios, 1997). It is known that about 80 species of aphids can represent vectors of 111 CMV. For this reason, control approaches can include the removal of aphids and the destruction of 112 CMV reservoirs weeds (Gallitelli, 1998). Interestingly, CMV, Alfalfa mosaic alfamovirus (AMV), 113 Potato M carlavirus (PVM), Potato Y potyvirus (PVY) and Tomato spotted wilt tospovirus (TSWV) 114 often compose mixed infections in tomato cultivated in the Mediterranean basin (Gallitelli, 1998). 115 116 Limited information has been reported on the tools and mechanisms controlling the virus diseases, especially regarding the economically relevant CMV. Specifically, a gap exists in the complete 117 understanding of the tomato-CMV-CHT interaction. For this reason, the viral titer, the plant 118 photosynthetic performance and the expression of genes related to antioxidant compounds and plant 119 resistance responses to pathogens, were investigated in tomato plants infected with CMV, with and 120 without CHT treatment. Therefore, the aim of this work was to investigate the efficacy of CHT as 121 an innovative and eco-friendly strategy to elicit defense responses in tomato plants against CMV, so 122 avoiding the negative consequences of the viral infection. 123

124

### 125 **2 Materials and methods**

### 126 2.1 Chitosan (CHT) and CMV sources and preparations

127

128 Low molecular weight CHT (50-190 kDa, 75-85% deacetylated) was purchased from Sigma-

129 Aldrich (448869; St. Louis, MO, USA). Chitosan (1 g) was dissolved in distilled water (40 mL)

130 containing 1 M acetic acid (9 mL) under overnight continuous stirring. The pH was 5.4. Eliciting

131 CHT solution was prepared by dissolving 1 g of this stock in 1 L of distilled water and foliar

sprayed (10 mL plant<sup>-1</sup>), while water was sprayed on untreated plants. Obtained as reported by Vitti
et al. (2015), *Cucumber mosaic virus* strain Fny inducing necrosis (CMV-Fny) was propagated in *Nicotiana tabacum* cv Xanthi plants. Subsequently, tobacco leaves exhibiting CMV-Fny symptoms
were macerated in 0.05 M sodium citrate buffer (pH 6.5) and the suspension was mechanically
rubbed on celite pre-dusted tomato leaves.

137

# 138 2.2 Experimental setup

139

140	Solanum lycopersicum var. cerasiforme seeds were sterilized (1 min in 1% Na-hypochlorite
141	solution) and then put to germinate on moist filter paper imbibed with sterile distilled water in Petri
142	dishes. After incubation for 24 h at 4 °C in the dark and for 2-3 days at 26 °C, seedlings were
143	transferred to pots filled with sterilized soil. At the four-leaf stage, plants were transplanted and
144	grown in a greenhouse at a temperature regime of 26/23 °C (day/night) and with a 16-h
145	photoperiod.
146	The experimental design included four experimental conditions (15 plants for each condition):
147	untreated plants; plants inoculated with CMV (CMV-TP); plants treated with CHT (CHT-TP);
148	plants treated with CHT and then inoculated with CMV 24 h after CHT treatment (CHT-CMV-TP).
149	Treatment/inoculation was performed starting from the tenth day after transplantation.
150	
151	2.3 CMV determination
152	
153	Twenty and ninety days after CMV inoculation, leaves from three plants randomly chosen, were
154	collected and used for a double-antibody sandwich enzyme-linked immunosorbent assay (DAS-

155 ELISA) according to Vitti et al. (2016). Measurements were performed spectrophotometrically

(model Multiskan GO: Thermo Fisher Scientific, Waltham, MA, USA) and the mean absorbance
value (OD<sub>405 nm</sub>) of six replicates for each experimental condition was taken.

158

### 159 2.4 Gas exchange and chlorophyll fluorescence

160

Photosynthetic activity (A), stomatal conductance to water vapor  $(g_s)$ , maximum quantum yield of 161 PSII ( $F_v/F_m$ ) (dark-adapted state) and quantum yield of PSII ( $\phi$ PSII) (light-adapted state) were 162 163 measured on clear days. Two apical fully developed leaves, belonging to one of three (for A and  $g_s$ determinations) or of two (for  $F_v/F_m$  and  $\phi$ PSII determinations) 2- and 4-month-old plants randomly 164 chosen for each experimental condition, were tested. Measurements were carried out using the LI-165 6400 portable photosynthesis system (LI-COR, Lincoln, NE, USA), operating between 10:00 and 166 12:00 a.m., at 398 ppm external CO<sub>2</sub> concentration, flow rate at 500  $\mu$ mol s<sup>-1</sup> and 1500  $\mu$ mol 167 photons m<sup>-2</sup> s<sup>-1</sup> photosynthetically active radiation (PAR). Temperature inside the leaf chamber was 168 maintained equal to environmental air temperature (28 °C) by instrument automatic temperature 169 regulation. 170

The same plants used for gas exchange measurements were chosen to measure chlorophyll 171 fluorescence at 10:00-12:00 a.m. using a leaf chamber fluorometer (LI-6400-40; Li-Cor, Inc.). On 172 each plant, both sun-adapted and dark-adapted leaves were chosen to measure fluorescence 173 parameters. On dark-adapted leaves (covered by silver film for 18 h before the measurements by 174 homemade clip holders), the maximum quantum yield of PSII photochemistry was calculated as 175 176  $F_{\rm v}/F_{\rm m} = (F_{\rm m} - F_{\rm o})/F_{\rm m}$  (Murchie and Lawson, 2013), where  $F_{\rm m}$  is the maximum fluorescence in the dark and  $F_0$  is the minimum level of fluorescence. On sun-adapted leaves, the quantum yield of PSII 177 ( $\phi$ PSII) was calculated as  $(F_{\rm m}' - F')/F_{\rm m}'$  (Murchie and Lawson, 2013), where  $F_{\rm m}'$  is the maximum 178 fluorescence in the light and F' is the steady-state fluorescence yield measured under actinic light. 179 The value of PAR inside the leaf chamber (light with a 90% red fraction at a wavelength of 630 nm 180

181	and a 10% blue fraction at 470 nm) during fluorescence measurements was 950 $\mu mol\ m^{-2}\ s^{-1}.$ This
182	value was chosen keeping into account (1) the measured average light saturation point (900 to 1000
183	$\mu$ mol m <sup>-2</sup> s <sup>-1</sup> ) and (2) the mean environmental irradiance monitored by the LI–6400 external
184	quantum light sensor every 3 seconds.
185	
186	2.5 SPAD measurements
187	
188	Leaf chlorophyll content was measured with a portable meter (SPAD-502, Minolta Camera Co.
189	Ltd., Osaka, Japan) between 10:00 and 11:00 a.m. Three apical fully developed leaves, belonging to
190	one of two 2- and 4-month-old plants randomly chosen for each condition were tested. The mean
191	value of the six measurements was recorded.
192	
193	2.6 Extraction and determination of total phenolic content
194	
195	Extractions and determinations were carried out in two randomly chosen plants for each condition,
196	analyzing leaves collected 60 h after the only or last treatment/inoculation. The total phenolic
197	content was analyzed spectrophotometrically using the Folin-Ciocalteu reagent and catechol as
198	standard, as reported by Sofo et al. (2017). All values were expressed as mg catechol equivalents
199	100 g <sup>-1</sup> of leaves fresh weight. The mean absorbance value ( $OD_{650 nm}$ ) of four replicates for each
200	condition was taken.
201	
202	2.7 Gene expression analysis
203	
200	

205 month-old plants for each experimental condition, were ground in liquid nitrogen in a

206 pestle and mortar. From the powder obtained, RNA was extracted using the RNAqueous Total RNA Isolation kit (AM1912, Ambion, Life Technologies, Thermo Fisher Scientific, Waltham, MA, 207 USA). On ice, RNA was purified adding 1/10 volume of 3M sodium acetate and 7/10 volume of 208 100% isopropanol. Samples were stored at -80 °C for 30 min, then centrifuged at 4 °C for 15 min 209 at maximum speed. To the pellet saved, 400 µL of 70% ethanol was added and two centrifugations 210 at 4 °C for 2 min at maximum speed taking off the supernatant were carried out. The open tubes 211 were put at 37 °C for 5 min in a heat block (ThermoStat Plus, Eppendorf, Hamburg, Germany) and 212 finally, RNase-free water was added to dissolve the RNA pellet. Purified RNA concentration and 213 purity were spectrophotometrically determined at 260 nm (NanoDrop ND-1000 UV/Vis 214 Spectrophotometer, NanoDrop Technologies, Wilmington, DE, USA). The Experion RNA StdSens 215 216 Analysis Kit (Bio-Rad Laboratories, Hercules, CA, USA) was used to assess the RNA quality. To the same concentration (1 µg) of all the RNA samples then utilized for the reverse transcription, 217 DNase (TURBO DNA-free Kit, AM1907, Invitrogen, Thermo Fisher Scientific, Waltham, MA, 218 USA) was added to degrade contaminating gDNA. cDNA synthesis was carried out starting from 219 oligo(dT)-primers and random hexamers (PrimeScript RT Reagent Kit, Perfect Real Time, 220 RR037A, Takara, Japan), then the cDNA obtained was 10-fold diluted using 1/10 TE buffer (1 mM 221 Tris-HCl, 0.1 mM EDTA, pH 8.0). Real-Time quantitative PCR (qPCR) was performed 222 coupled with SYBR Green fluorescent dye, using the 7500 Real-Time PCR System (Applied 223 Biosystems, Lennik, Belgium). Cycling conditions were 95 °C for 20 s, 50 cycles of 3 s at 95 °C 224 and 30 s at 60 °C. A final volume of 10 µL contained 2 µL of cDNA produced, 5 µL of Fast SYBR 225 Green Master Mix (4385612, Applied Biosystems, Lennik, Belgium), 0.3 µL of each forward and 226 reverse primer (300 nM) (Table 1) and 2.4 µL of RNase-free water. A dissociation curve followed 227 to evaluate the amplification specificity. Sequences for the reference genes, as well as for the genes 228 of interest, were searched in the NCBI (https://www.ncbi.nlm.nih.gov) and JGI Phytozome 229 (https://phytozome.jgi.doe.gov/pz/portal.html) databases, and related primers were designed using 230

231	Primer3 Program and nBLAST-NCBI (Table 1). Primers efficiency ( $E = 10^{(-1/slope)} - 1$ ) was
232	measured using serial dilutions from 1/2 to 1/64 of cDNAs collected in a pooled sample and
233	realizing C <sub>q</sub> versus log(dilution) calibration lines (Table 2). The GrayNorm algorithm
234	(https://github.com/gjbex/GrayNorm) was adopted to select the combination of reference genes able
235	to yield the highest possible accuracy. The expression of the genes encoding (EC 1.11.1.7)
236	peroxidase (CEVI-1), non-expressor of pathogenesis-related genes 1 (NPR1), (EC 2.5.1.32)
237	phytoene synthase 2 (PSY2) and (EC 4.3.1.5) phenylalanine ammonia lyase (PAL5) was considered
238	relative to five reference genes selected for normalization: SAND, TIP4, TUB (tubulin), UK
239	(uridylate kinase) and GAPDH (glyceraldehyde-3-phosphate dehydrogenase). In Tables 1 and 2 are
240	respectively reported sequences and efficiencies of primers for the reference genes and genes of
241	interest for qPCR analysis. Genes of interest relative expression was calculated as $2^{-\Delta Cq}$ , and the $2^{-}$
242	$\Delta Cq$ values geometric average was used for data normalization.
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253	

Name	Sequence	Tm (°C)
SAND	5'-CCAGCTAACTTTCTCCATGCTTAC-3'	58.1
	5'-ACCAACAAGACTGATAACCTTTTGT-3'	55.3
TIP4	5'-CTGTTAAAGTGAGAGTCATGCCTAG-3'	58.4
	5'-TGCAAACGAGTGTCTCTTAGTCT-3'	57.8
TUB	5'-AGAATGCCGATGAATGTATGGT-3'	55.0
	5'-CAGGGAATCTCAAACAGCAAG-3'	55.2
UK	5'-TGGTAAGGGCACCCAATGTGCTAA-3'	59.7
	5'-ATCATCGTCCCATTCTCGGAACCA-3'	59.9
GAPDH	5'-GATGTCTCCGTTGTCGATCTT-3'	55.1
	5'-CAAGATACCCTTCAATTTACCCTCT-3'	55.9
CEVI-1 (GenBank accession number Y19023)	5'-TCACCAACAAGGGAATGGAT-3'	52.0
	5'-TGGATCAGGGCTACCACTTC-3'	54.9
NPR1 (Phytozome Solyc07g040690.2)	5'-CGATGATTTGCGTATGAAGC-3'	54.3
	5'-CCAGGGGTAATTCAGACGTG-3'	54.4
PSY2 (Phytozome Solyc02g081330.2)	5'-CCGAATTCCGAGGTCTCATA-3'	54.5
	5'-CCTGTCTCCCACCTTTCTTG-3'	54.5
PAL5 (GenBank accession number M90692)	5'-CTCTGGCAATGGGTGCTAAT-3'	55.2
	5'-CAGGGGTCATCAGCATAGGT-3'	55.3

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256			
257			
258			
259			
260			

Primer	PCR efficiency	Equation	R <sup>2</sup>
	(80-120%)		Coefficient
SAND	93.03%	y = -3.5010x + 25.795	$R^2 = 0.9923$
TIP4	100.51%	y = -3.3097x + 24.447	$R^2 = 0.9964$
TUB	100.78%	y = -3.3035x + 23.606	$R^2 = 0.9984$
UK	87.58%	y = -3.6605x + 25.201	$R^2 = 0.9987$
GAPDH	96.92%	y = -3.3981x + 21.682	$R^2 = 0.9992$
CEVI-1	105.70%	y = -3.1926x + 25.276	$R^2 = 0.9944$
NPR1	103.53%	y = -3.2401x + 25.306	$R^2 = 0.9997$
PSY2	92.89%	y = -3.5049x + 24.470	$R^2 = 0.9939$
PAL5	100.73%	y = -3.3045x + 22.149	$R^2 = 0.9990$

263

## 264 **2.8 Statistical data analysis**

265

Normal distribution of data was tested performing the Shapiro-Wilk test ( $P \le 0.05$ ) and

homoscedasticity was tested performing the Bartlett's test ( $P \le 0.05$ ). Data were analyzed by one-

and two-way analysis of variance (ANOVA). Parametric and non-parametric as multiple

comparisons were performed using the Tukey's HSD test and the Kruskal-Wallis test, respectively.

270 Statistical analyses were performed using the software RStudio: Integrated Development for R,

version 1.0.136 (RStudio, Inc., Boston, MA, USA).

272

### 273 **3 Results**

274

Twenty days after CMV inoculation, only CMV-TP showed mosaic and chlorosis in leaves, as well
as their deformation (Figure 1A). On the contrary, the absence of symptoms induced by CMV was

282	3.1 CMV load
281	
280	symptoms was detected in tobacco CHT-CMV-TP (Figure 1B).
279	days after CMV inoculation, tobacco CMV-TP showed mosaic in leaves, but the absence of CMV
278	the same inoculations and treatments methods, were also applied to Nicotiana tabacum. Eleven
277	detected in CHT-CMV-TP (Figure 1A). Furthermore, the same four experimental conditions, with

CHT-CMV-TP showed the mean optical density value significantly lower than that determined in
CMV-TP, both in 20 days and 90 days after CMV inoculation determinations (-86% and -100%,
respectively). The absence of CMV was detected in untreated and CHT-TP, and no significant
difference was revealed between these two conditions (Figure 2).

288

#### 289 **3.2** Gas exchange and chlorophyll fluorescence

290

Considering the A value between 2- and 4-month-old plants, CMV-TP and CHT-TP showed the 291 lowest and the highest A value, respectively (Figure 3A). Although not significantly, CHT-CMV-292 TP had the A value of 44% higher than that determined in CMV-TP, and untreated plants showed 293 the A value lower (-25%) than that determined in CHT-TP (Figure 3A). The 2-month-old CHT-TP 294 and CMV-TP showed the highest and the lowest g<sub>s</sub> value, respectively (Figure 3B). Chitosan 295 treatment in CHT-CMV-TP caused a g<sub>s</sub> value of 146% significantly higher than that determined in 296 297 CMV-TP, and a g<sub>s</sub> value not significantly different than that determined in CHT-TP. Untreated plants showed a g<sub>s</sub> value not significantly different than that of CMV-TP, and a g<sub>s</sub> value 298 significantly lower (-59%) than that of CHT-TP (Figure 3B). Results were not significantly 299 different between the four conditions, in 4-month-old plants (Figure 3B). The 4-month-old CHT-TP 300 and CHT-CMV-TP showed a significant decrease in the g<sub>s</sub> value, compared to the g<sub>s</sub> value of 2-301

302	month-old plants (-58 and -50%, respectively) (Figure 3B). The 4-month-old CMV-TP showed the
303	lowest $F_v/F_m$ value, lower (-6%) than the average value of the $F_v/F_m$ values of all the other
304	conditions (Figure 3C). The 4-month-old CMV-TP also showed a significant decrease of the $F_v/F_m$
305	value, compared to that determined in 2-month-old plants (-7%) (Figure 3C). The 4-month-old
306	untreated, CHT-TP and CHT-CMV-TP showed a significant increase in the $\phi$ PSII value, compared
307	to that of 2-month-old plants (298, 217 and 218%, respectively) (Figure 3D).
308	
309	3.3 SPAD
310	
311	In 2-month-old plants, CHT-TP had a SPAD value 11% significantly higher than that found in
312	untreated plants. Finally, among 4-month-old plants, the CHT-CMV-TP showed the lowest SPAD
313	value (Figure 4).
314	
315	3.4 Total phenolic content
316	
317	Untreated plants and CHT-CMV-TP showed the highest and the lowest total phenolic content,
318	respectively. CHT-CMV-TP showed a total phenolic content significantly lower than that
319	determined in untreated and CMV-TP (-41 and -40%, respectively) (Figure 5).
320	
321	3.5 Gene expression analysis
322	
323	Reference genes were TUB, UK and GAPDH for the experimental conditions represented in Figure
324	6; SAND, TIP4 and GAPDH for those represented in Figure 7.
325	Untreated plants were assumed as a control, compared to CMV-TP, and no significant difference
326	was found in the relative expression of all the genes of interest assayed (Figure 6). Although not 14

327	significant, an increase in CEVI-1 transcripts was detected in the leaves of CMV-TP (Figure 6A).
328	Finally, <i>PAL5</i> expression was the most stable in the considered conditions (Figure 6).
329	CMV-TP were then assumed as a control, compared to CHT-TP and CHT-CMV-TP (Figure 7). A
330	significant increase in PAL5 expression was observed both in CHT-TP and CHT-CMV-TP,
331	compared to CMV-TP (Figure 7D). Conversely, no significant up-regulation of NPR1 and down-
332	regulation of CEVI-1 and PSY2 transcripts were observed (Figures 7B, A and C, respectively).
333	
334	4 Discussion
335	
336	The plant-virus interaction affects the chloroplast. More specifically, the virus replication and viral
337	movement involve chloroplast factors (Zhao et al., 2016). As a result, the chloroplast structure and
338	the expression of photosynthesis-related proteins are perturbed as well as the viral symptoms are
339	manifested (Zhao et al., 2016).
340	Studies have shown the efficacy against strains of CMV infection of treatment with Trichoderma
341	harzianum T-22 (Vitti et al., 2016, 2015), Paenibacillus lentimorbus B-30488 (Kumar et al., 2016)
342	and benzo-(1,2,3)-thiadiazole-7-carbothioic acid S-methyl ester (BTH) (Anfoka, 2000).
343	The present research investigated the ability of CHT to elicit defense response in tomato plants
344	inoculated with CMV.
345	
346	4.1 CMV symptoms and load monitoring
347	
348	A phenotypical observation showed the capacity of CHT to control CMV symptoms, testing tomato
349	and tobacco as host plants. Indeed, neither tomato nor tobacco CHT-treated then CMV-infected
350	plants displayed viral infection symptoms (Figures 1A and B, respectively). The results of DAS-

ELISA showed the efficacy of CHT to control foliar CMV load in tomato CHT-CMV-TP, both at 20 days and at 90 days after CMV inoculation measurements (Figure 2). In agreement with our results, determined by ELISA, a CHT-induced resistance in potato against PVX was suggested as probably mediated by the enhanced ribonuclease activity and callose deposition (Chirkov et al., 2001). Furthermore, in *Nicotiana tabacum* L. cv. Samsun leaves, Nagorskaya et al. (2014) showed that CHT limited TMV coat protein content and infectivity as well as increased the hydrolases (proteases and RNases) activity. They also detected a highest content of abnormal virions.

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#### **4.2 Plant physiological responses to CHT and CMV**

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Photosynthetic activity was considerably analogous, between the short and long term (when plants 361 were 2- and 4-month-old, respectively). Although not significantly, CHT treatment improved the 362 photosynthetic activity (Figure 3A). Van et al. (2013) reported the effects of CHT nanoparticles on 363 Robusta coffee and, according to our results, they found an enhanced photosynthesis net rate and 364 besides, they supposed increased stomatal cells opening degree and stomatal conductance because 365 of the polycation property of CHT that raise the osmosis pressure of stomatal cells. Salachna et al. 366 (2017) suggested that the positive effect of CHT on plant growth parameters may cause the 367 increased stomatal conductance with CHT foliar application. However, the reduction of stomatal 368 apertures width after foliar CHT treatment of bean has also been reported (Iriti et al., 2009). 369 370 Differently from A data, effect of treatment/inoculation on stomatal conductance was more evident when plants were 2-month-old (Figure 3B). As reported by Vitti et al. (2016), the 2-month-old 371 tomato plants inoculated with CMV showed reduced stomatal conductance to water vapor besides 372 decreased photosynthetic activity (Figures 3B and A, respectively). 373

374 In agreement with the results obtained by Marler et al. (1993) on papaya leaves inoculated with Papaya ringspot virus (PRV), a significantly lower maximal quantum yield of PSII value  $(F_v/F_m)$ 375 was detected in CMV-TP than untreated ones, in 4-month-old plants (Figure 3C), probably 376 indicating a damage to PSII or photoinactivation caused by a decrease of the opened reaction 377 centers. The quantum yield of PSII values detected in 4-month-old plants were higher (some 378 379 significantly) than ones detected in 2-month-old plants (Figure 3D). No significant influence of CHT treatment was recorded in  $F_v/F_m$  ratio and similarly occurred in  $\phi$ PSII value, compared to 380 untreated and only infected plants (CMV-TP) (Figures 3C and D, respectively). 381 SPAD meter measures the relative chlorophyll content by estimating the leaf greenness. Indeed, 382 chlorophyll reflects a green light, inducing this color in plants (Shi et al., 2018). Results indicated 383 no significant difference in SPAD readings of CMV-TP, compared to untreated plants, in the same 384 385 interval measurements. However, in 4-month-old plants, CMV-TP had a SPAD value lower than that of untreated plants, reflecting the changed pigmentation responsible for the symptoms 386 displayed. Furthermore, our data indicated that CHT-TP showed a significant increase in 387 chlorophyll content, compared to untreated ones, in 2-month-old plants (Figure 4). Van et al. (2013) 388 also reported that CHT nanoparticles improved the content of chlorophylls as well as the uptake of 389 nitrogen and magnesium that constitute the chlorophyll chemical structure. 390 Phenols produced by plants vary in the defense response against environmental stresses (Sofo et al., 391 2017). In our case, 60 h after CMV inoculation, foliar total phenolic content was not significantly 392 393 different between CMV-TP and untreated plants (Figure 5). The same result was obtained six days after the inoculation of CMV-Y in tobacco plants (Ipper et al., 2008). Furthermore, the treatment 394 with CHT, both alone and before CMV inoculation (CHT-TP and CHT-CMV-TP, respectively), 395 caused a decrease of total phenols, compared to ones determined in only infected and untreated 396 plants (Figure 5). In agreement with our result, Coqueiro et al. (2011) assessed the effect of low 397 molecular weight CHT in tomato plants, then inoculated with Xanthomonas gardneri three days 398

after CHT treatment. They observed that the total phenolic compounds of the CHT-treated plants
increased starting from the second day after the inoculation of *Xanthomonas gardneri*, hence neither
before nor within 24 h after inoculation. Therefore, in our case, the absence of a CHT-conditioned
accumulation of phenolics as a response to CMV in CHT-CMV-TP may be due to the chosen
timing between CHT application and CMV inoculation (24 h) and/or between inoculation and
analysis (60 h).

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## 406 4.3 Expression analysis of defense-related genes

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The systemic acquired resistance (SAR) provides to the plant a long-lasting systemic resistance to 408 consecutive infections by many pathogens (Mou et al., 2003). The SAR involves plant responses, 409 such as the production of reactive oxygen species (ROS) and pathogenesis-related (PR) proteins, as 410 well as the lignification and cell wall reinforcement through the cell wall structural proteins cross-411 linking (Pandey et al., 2017). More specifically, the CHT-induced resistance can enhance the 412 activities of defense-related enzymes, such as peroxidase, PAL, polyphenol oxidase, superoxide 413 414 dismutase and catalase (Xing et al., 2015). The induced systemic resistance (ISR) is triggered by some bacteria and fungi and requires jasmonic acid (JA) and ethylene (ET). Differently, the SAR 415 requires salicylic acid (SA), exogenously applied (Mou et al., 2003) or endogenously produced. 416 To optimize the GrayNorm output for the molecular analysis, the four different experimental 417 conditions were divided into two groups (Figures 6 and 7). It was appropriate to compare the 418 419 expression of the assayed genes in CHT-TP and CHT-CMV-TP to untreated plants. Such comparison denoted that CHT treatment causes an up-regulation of CEVI-1, NPR1 and PAL5 420 expressions, whereas the increase of *PSY2* is only observed in even infected plants. 421 Particularly, peroxidases are enzymes that catalyze the hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) decomposition 422 oxidating many phenolic and non-phenolic substrates (Pandey et al., 2017). The implication of plant 423

peroxidases in processes, such as lignification, suberization, cell wall metabolism, defense against 424 pathogens and ROS metabolism is well-known (Pandey et al., 2017). Although not statistically 425 different, a strong increase in CEVI-1 expression occurred in CMV-TP, compared to untreated 426 427 plants (Figure 6A). Such finding is in accordance with the observation that at seven days post Tomato mosaic virus (ToMV) inoculation of tomato plants, CEVI-1 expression was induced in 428 leaves (Mayda et al., 2000). CEVI-1 expression strong up-regulation could be associated with the 429 cell wall reinforcement and the ROS content. Mayda et al. (2000) also reported that were able to 430 induce *CEVI-1* expression neither incompatible interactions nor some infiltrated signal molecules. 431 Compared to CMV-TP, the CHT treatment in CHT-CMV-TP seemed to limit the CEVI-1 432 transcripts amount (Figure 7A), suggesting that CHT plays a role in the regulation of the ROS 433 levels, thus controlling such CMV infection effect. 434 435 Cytosolic non-expressor of pathogenesis-related genes 1 (NPR1) regulates the salicylate- and jasmonate-dependent pathways cross-talk (Spoel et al., 2003). Moreover, in Arabidopsis, Bacillus 436 cereus AR156-induced ISR to Botrytis cinerea required NPR1 and JA/ET-signaling pathway (Nie et 437 al., 2017). Interestingly, another study demonstrated that SAR induction by AR156 required NPR1 438 and SA-signaling pathway (Niu et al., 2016). Wu et al. (2012) suggested that Arabidopsis NPR1 439 binds SA through cysteines<sup>521/529</sup> via copper. Just during SAR, NPR1 activates the PR gene 440 expression (Mou et al., 2003). In our study, the expression of NPR1 was analyzed as well, but no 441 significant differences in the transcripts amount were detected (Figures 6B and 7B). However, 442 NPR1 expression was slightly up-regulated in CHT-TP and CHT-CMV-TP, compared to CMV-TP 443 (Figure 7B) and untreated ones. This could indicate the CHT efficacy against CMV by triggering 444 SAR-related defense responses in tomato plants. Jia et al. (2016) reported that CHT oligosaccharide 445 pretreatment induced TMV resistance in Arabidopsis through the SA signalling pathway. An 446 optimal low concentration (50 mg  $L^{-1}$ ) applied one day before TMV inoculation was used. 447

Interestingly, Doares et al. (1995) observed an increase in the level of JA in leaves of excisedtomato plants after supplying CHT oligosaccharides.

Photosynthesis and photoprotection are processes involving plant carotenoids (Giorio et al., 2008). 450 451 The first biosynthetic step of carotenoids involves two molecules of geranylgeranyl pyrophosphate (GGPP) and is catalyzed by the enzyme phytoene synthase (PSY), that is encoded by PSY2 in 452 chloroplasts and PSY1 of chromoplasts in tomato. From phytoene, sequential reactions differently 453 form lycopene and then cyclic carotenoids such as lutein, zeaxanthin and violaxanthin (Fraser and 454 Bramley, 2004; Giorio et al., 2008; Meléndez-Martínez et al., 2010). However, such two genes 455 encode PSY2 and PSY1, respectively (Giorio et al., 2008). In our case, PSY2 expression was up-456 regulated in CMV-TP, but such a result was not significant; On the whole, no significant difference 457 was observed in the PSY2 transcript amount (Figures 6C and 7C). Interestingly, Ibdah et al. (2014) 458 459 found that CMV-Fny infection caused an increased phytoene content in Nicotiana tabacum L. cv. Samsun NN roots; however, the carotenoid production was reduced because the enzyme phytoene 460 desaturase was down-regulated by CMV. 461

Additionally, phenylalanine ammonia lyase (PAL) converts the L-phenylalanine to ammonia and 462 trans-cinnamic acid. PAL-catalyzed reaction is the first in phenylpropanoid metabolism. Lee et al. 463 (1994) demonstrated that multiple initiation sites in PAL5 allow the tomato plant to respond to 464 different environmental stimuli in a tissue-specific fashion. In our experiment, a significant increase 465 in PAL5 expression occurred in CHT-TP and CHT-CMV-TP, compared to CMV-TP (Figure 7D). 466 467 Such a result is in accordance with Mejía-Teniente et al. (2013), who observed that after the first CHT treatment in Capsicum annuum L., PAL activity as well as pal expression increased. No 468 significant difference was observed in PAL5 expression level in CMV-TP, compared to untreated 469 plants (Figure 6D). Ogawa et al. (2006) found increased PAL A and PAL B transcripts as well as 470 PAL activity in TMV-infected (hypersensitive reaction lesion-bearing) tobacco leaves, suggesting 471

the phenylalanine pathway as the main route of SA synthesis. Compared to CMV-TP, the

473	significantly higher PAL5 expression detected in CHT-CMV-TP (Figure 7D), suggests the
474	involvement of phenylpropanoid-derived products as lignin and SA.
475	The RNA silencing participates in the antiviral plant mechanism, though it is overcome by the
476	viruses encoding RNA silencing suppressors (Carr et al., 2010), such as the 2b protein of CMV.
477	Unfortunately, how virus infection is controlled in plants exhibiting SAR has not
478	been fully understood (Carr et al., 2010).
479	
480	5 Conclusions
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482	This paper reports the ability of chitosan as a preventive treatment able to elicit defense responses in
483	tomato plants against CMV-Fny infection. Chitosan was able to reduce the CMV titer and improved
484	the gas exchange of the infected plants. Furthermore, a SAR-related response induced by chitosan,
485	also by influencing the plant oxidative status, is hypotizable. Such findings represent a new and
486	additional piece of the puzzle depicting effective, sustainable and environmentally safe methods of
487	plant disease control. Further studies could clearly define the whole set of the resistance responses
488	triggered specifically in such host-pathogen-elicitor combination.
489	
490	Conflict of interest statement
491	Declarations of interest: none.
492	

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496

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- 692 Figure 1. Phenotypical observations of symptoms induced by *Cucumber mosaic virus* (CMV) 20 days and 11 days after
- 693 CMV inoculation of representative A) Solanum lycopersicum var. cerasiforme and B) Nicotiana tabacum cv Xanthi
- 694 plants, respectively. Four different experimental conditions: untreated plants; plants inoculated with CMV (CMV-TP);
- 695 plants treated with CHT (CHT-TP); plants treated with CHT and then inoculated with CMV 24 h after CHT treatment
- 696 (CHT-CMV-TP).
- 697



- Figure 2. *Cucumber mosaic virus* (CMV) load 20 days (light grey bars) and 90 days (dark grey bars) after CMV
- inoculation of tomato plants. Mean values (n = 6) are represented. Standard deviations are represented by bars.
- Significant differences ( $P \le 0.05$ ) among treatments and time are indicated by different letters, according to non-
- 703 parametric two-way ANOVA. Four different experimental conditions: untreated plants; plants inoculated with CMV
- 704 (CMV-TP); plants treated with CHT (CHT-TP); plants treated with CHT and then inoculated with CMV 24 h after CHT
- 705 treatment (CHT-CMV-TP).
- 706



- Figure 3. Gas exchange and chlorophyll fluorescence determinations. A) Photosynthetic activity (A), B) stomatal 709
- 710 conductance to water vapor ( $g_s$ ), C) maximal quantum yield of PSII ( $F_v/F_m$ ) and D) quantum yield of PSII ( $\phi$ PSII),
- measured in 2- (light grey bars) and 4-month-old (dark grey bars) tomato plants. Mean values ( $n \ge 6$  for gas exchange; n711
- 712 = 4 for chlorophyll fluorescence) are represented. Standard deviations are represented by bars. Significant differences
- 713  $(P \le 0.05)$  among treatments and time are indicated by different letters, according to parametric and non-parametric  $(g_s)$
- 714 two-way ANOVA. Four different experimental conditions: untreated plants; plants inoculated with CMV (CMV-TP);
- 715 plants treated with CHT (CHT-TP); plants treated with CHT and then inoculated with CMV 24 h after CHT treatment
- 716 (CHT-CMV-TP).





Figure 4. Chlorophyll content (SPAD) measured in 2- (light grey bars) and 4-month-old (dark grey bars) tomato plants. Mean values (n = 6) are represented. Standard deviations are represented by bars. Significant differences ( $P \le 0.05$ ) among treatments and time are indicated by different letters, according to non-parametric two-way ANOVA. Four different experimental conditions: untreated plants; plants inoculated with CMV (CMV-TP); plants treated with CHT (CHT-TP); plants treated with CHT and then inoculated with CMV 24 h after CHT treatment (CHT-CMV-TP).



- Figure 5. Total phenolic content of tomato leaves. Mean values (n = 4) are represented. Standard deviations are
- represented by bars. Significant differences ( $P \le 0.05$ ) among treatments are indicated by different letters, according to
- 731 non-parametric one-way ANOVA. Four different experimental conditions: untreated plants; plants inoculated with
- 732 CMV (CMV-TP); plants treated with CHT (CHT-TP); plants treated with CHT and then inoculated with CMV 24 h
- 733 after CHT treatment (CHT-CMV-TP).



- 736 Figure 6. Genes expression in CMV-TP relatively expressed to the Untreated tomato plants. A) peroxidase (CEVI-1), B)
- 737 non-expressor of pathogenesis-related genes 1 (NPR1), C) phytoene synthase 2 (PSY2) and D) phenylalanine
- 738 ammonia lyase (*PAL5*). Mean values (n = 4) are represented. Standard errors are represented by bars. Significant
- 739 differences ( $P \le 0.05$ ) among treatments are indicated by different letters, according to parametric one-way ANOVA.
- 740 Two different experimental conditions: untreated plants; plants inoculated with CMV (CMV-TP).





