1 2	Occurrence of <i>Salmonella</i> Typhimurium resistance under sublethal/repeated exposure to cauliflower infusion and Infection effects on <i>Caernohabditis elegans</i> host test organism
3 4	Maria Sanz-Puig ^ª , Alejandra Arana-Lozano ^ª , M. Consuelo Pina-Pérez ^b , Pablo Fernandez ^c , Antonio Martinez ^ª , Dolores Rodrigo ^{ª*}
5 6	^a Instituto de Agroquímica y Tecnología de Alimentos – Consejo Superior de Investigaciones Científicas (IATA-CSIC). Unidad Asociada UPCT. Paterna, València, Spain.
7	^b Institute of Life Technologies, HES.SO VALAIS-WALLIS, Route du Rawil 64, Sion, Switzerland
8	^c Universidad Politécnica de Cartagena. Unidad Asociada al CSIC. Cartagena, Murcia, Spain.
9	
10	*corresponding author: lolesra@iata.csic.es
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12 Abstract

Resistant bacteria to antimicrobials are increasingly emerging in medical, food industry and livestock environments. The present research work assesses the capability of *Salmonella enterica* var Typhimurium to become adapted under the exposure to a natural cauliflower antimicrobial by-product infusion in consecutive repeated exposure cycles. *Caernohabditis elegans* was proposed as *in vivo* host-test organism to compare possible changes in the virulent pattern of the different rounds treated *S. enterica* var Typhimurium and untreated bacterial cells.

20 According to the obtained results, S. enterica var Typhimurium was able to generate resistance 21 against a repeated exposure to cauliflower by-product infusion 5 % (w/v), increasing the 22 resistance with the number of exposed repetitions. Meanwhile at the first exposure, 23 cauliflower by-product infusion was effective reducing S. enterica var Typhimurium ($\approx 1 \log_{10}$ 24 cycle), S. enterica var Typhimurium become resistant to this natural antimicrobial after the 25 second and third treatment-round, and was able to grow ($\approx 1 \log_{10}$ cycle). In spite of the 26 increased resistance observed for repeatedly treated bacteria, the present study reveals no 27 changes on C. elegans infection effects between resistant and untreated S. enterica var 28 Typhimurium, according to phenotypic parameters evaluation (lifespan duration and egg-29 laying).

31 Keywords: Natural antimicrobials, *S. enterica* var Typhimurium, resistance, *Caernohabditis* 32 *elegans*, infection.

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37 1.Introduction

Hurdle technology is a processing concept industrially well established to achieve efficient 38 39 microorganisms reduction by the combination of different preservation techniques applied at 40 slight intensive conditions maintaining natural valuable properties of foods treated products 41 from a nutritional and sensory point of view (Pasha et al., 2014; Pina-Pérez et al., 2013; Sanz-42 Puig et al., 2017a). The combination of natural or chemical antimicrobials with other physical 43 methods is one of the most commonly used practices to synergistically enhance microbial and 44 enzymes reduction improving the safety and shelf-life of treated products (Guerrero et al., 45 2017; Sudhaus et al., 2012). However, nowadays, consumers are really concerned about 46 chemical additives, many of them close related with allergies and other possible health 47 problems (Asioli et al., 2017). Natural antimicrobials (e.g. polyphenols, glucosinolates, vegetable pigments, antimicrobial peptides) and phytochemical extracts (from soya, Stevia 48 49 Rebaudiana Bertoni, cauliflower infusion, citrus fruits essential oils extracts) could be 50 introduce in food formulation as an alternative to synthetic antimicrobials (Gutierrez et al., 51 2009; Sansano et al., 2017; Sanz-Puig et al., 2015a; Tiwari et al., 2009), additionally with other 52 technological (thickening agents, gelling agents) and functional effects in novel food (enhanced

53 antioxidant potential, enhanced fiber content in food). Many bioactive compounds can be 54 obtained from agro-industrial by-products (Fernandez-Lopez et al., 2005; Sanz-Puig et al., 2015 a,b; Viuda-Martos et al., 2007). Sanz-Puig et al. (2015b) studied different concentrations of 55 56 vegetable by-product infusions (cauliflower, mandarin, orange, lemon, and okara) against 57 Salmonella enterica var Typhimurium, and according to the obtained results 5 % (w/v) of 58 cauliflower resulted the most effective bactericidal natural aqueous extract reducing close to 6 59 log₁₀ cycles the initial S. enterica var Typhimurium load. Valorization studies carried out to 60 make profit from valuable vegetable wastes from food industry is in the priority research axes 61 of the European Union H2020 in order to support sustainable development by means of 62 circular economy fundamentals (EC, 2017).

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Microorganisms are becoming adapted and resistant to (i) veterinary antibiotics, used in animals farming, (ii) clinical antibiotics used in human medicine and (iii) antimicrobial additives used in food preservation, representing a great concern for the scientific community (WHO, 2017).

68 Cross-resistances along the complete food chain, and changes in virulence of pathogenic 69 bacteria, may even emerge (Kalily et al., 2016, 2017; Kisluk et al., 2013). Zanini et al. (2014) 70 studied the bacterial adaptive responses to antibiotics induced by sublethal concentration of 71 citral on first- and second-generation cells of Listeria monocytogenes serovar 4b and 72 Salmonella enterica var Typhimurium. They concluded that the presence of citral in the culture 73 medium of L. monocytogenes 4b and S. enterica var Typhimurium increased the antibiotic 74 susceptibility of the first generations, while an increase in antibiotic resistance was observed in 75 the second generation of *S. enterica* var Typhimurium.

Recent *in vivo* models are being proposed to evaluate mechanisms of pathogenic infection,
and bacterial genes responsible for changes in virulence. Among those, Additionally, *C. elegans*has been defined as a valid model for human pathogen infection, disease and health

promotion (Balla and Troemel, 2013; Gammon, 2017; Kurz and Ewbank, 2000). The evaluation
of the *C. elegans* lifespan has been previously used as an indicator of the impact that certain
molecules, e.g. antioxidants, metabolites and synthetic compounds, could have on human
health and their influence in the anti-aging process (Liu et al., 2016; Rollins et al., 2017; Wang
and Wink, 2016).

Silva et al. (2015) evaluated the effect of two antimicrobial substances, carvacrol and citral, on *L. monocytogenes* and *L. innocua* cells, as well as possible virulence changes in sub-lethally damaged cells, using *C. elegans* as *in vivo* test organism. According to Silva et al. (2015) results, the lifespan of *C. elegans* population fed on a lawn of *L. monocytogenes* previously treated with carvacrol was reduced in comparison with lifespan exerted by *C. elegans* population fed on a lawn of *E. coli* OP50 (as negative control), *L. monocytogenes* treated with citral, and, treated/not treated *L. innocua*.

91 Understanding the behaviour of foodborne pathogens exposed to sublethal concentrations of 92 natural antimicrobials, and the possible implications in terms of bacterial resistance generation 93 and changes in virulence are key questions to search for alternatives to conventional 94 preservatives in food and to assess the safety of the hurdle technology processing application 95 (Pina-Perez et al., 2013; Sanz-Puig et al., 2017a). Evaluation of some C. elegans phenotypic 96 factors (such as lifespan, reproduction patterns, and mobility) under the exposure to 97 pathogenic human bacteria could be a preliminary indicator of risks for consumers when these 98 natural antimicrobials are used below the bactericidal dosage for food formulation with 99 preservative objectives (Kastbjerg et al., 2010; Schroeder et al., 2017).

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101 The main aim of the present study is the assessment of *S. enterica* var Typhimurium behaviour 102 under the repeatedly exposure to a cauliflower natural antimicrobial extract, subsequently 103 evaluating the effects of infection by repeatedly treated/untreated *S. enterica* var 104 Typhimurium on the *C. elegans in vivo* host-model organism.

106 **2. Materials and methods**

107 **2.1 Cauliflower by-product extract**

108 Cauliflower by-product (mainly external leaves of cauliflower plants) was provided as 109 dehydrated residues from primary production of TRASA S.L., and was washed in sterile water, 110 dried and homogenized using a laboratory grinder (Janke & Kunkel IKA Labortechnik) (Brandi 111 et al., 2006).

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Five per cent (w/v) of dried cauliflower by-product was selected in the previous study according to previous results of Sanz-Puig et al. (2015a). The infusion was prepared as follows: buffered peptone water (0.1% (w/v)) was boiled and then the dry by-product was added and allowed to infuse for 30 min (Sanz-Puig et al., 2016). The extracts were then centrifuged at 4,000 rpm for 15 min at 4 °C and filtered three times, using filters with a pore size of 11 and 2.5 μ m to eliminate smaller particles (Whatman), and 0.45 μ m (PVDF syringe filter) to sterilize.

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120 2.2 Bacterial strain

121 A pure culture of S. enterica var Typhimurium (CECT 443) was provided freeze-dried by the 122 Spanish Type Culture Collection. It was rehydrated with 10 mL of tryptic soy broth (TSB) 123 (Scharlab Chemie, Barcelona, Spain). After 20 min, it was transferred to 500 mL of TSB and 124 incubated at 37 °C with continuous shaking (Selecta Unitronic) at 200 rpm for 14 h to obtain 125 cells in a stationary growth stage. The cells were centrifuged (Beckman Avanti J-25) twice at 126 4.000 rpm and at 4 °C for 15 min and then resuspended in TSB. The cells were finally 127 resuspended in 20 mL of TSB, then dispensed in 2 mL vials with glycerol at 20% to a final concentration of 10⁸ CFU/mL obtained by plate count, and finally frozen and stored at –80 °C. 128

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130 **2.3 Evaluation of microbial resistance**

The initial population of *S. enterica* var Typhimurium (10⁷ CFU/mL) was exposed to three 131 132 consecutive antimicrobial treatments. Each treatment consisted in exposure to 5 % (w/v) cauliflower by-product infusion at 37 °C for 4 h with continuous shaking (200 rpm) (sub-lethal 133 134 treatment) (Sanz-Puig et al., 2015a). Afterwards, the sample was centrifuged to recover the 135 microbial cells. Recovered cells were grown in TSB culture overnight to achieve stationary phase (10⁹ CFU/mL), inoculated in cauliflower by-product extract (10⁷ CFU/mL) and treated 136 again using the same conditions as described above. Before and after each treatment, the 137 138 antimicrobial effect against the S. enterica var Typhimurium population was evaluated by plate 139 count in tryptic soy agar (TSA) (Scharlab Chemie, Barcelona, Spain). The entire experiment was 140 carried out in triplicate with three different infusion batches.

S. Typhimurium treated once and *S. enterica* var Typhimurium treated three times were
included in the *C. elegans* infection study, because they were the most different populations
with regard to their resistance to the cauliflower antimicrobial exposure.

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145 2.4 C. elegans studies

C. elegans strain N2, obtained from the College of Biological Sciences, Minnesota University,
 USA, was used to evaluate the possible virulence changes in *S. enterica* var Typhimurium
 populations as a consequence of repeated exposure to antimicrobial from cauliflower by product infusion. *C. elegans* was maintained in plates with Nematode Growth Medium (NGM)
 agar and a bacterial lawn of *E. coli* OP50 (Silva et al., 2015).

151 In order to evaluate the effect of different *S. enterica* var Typhimurium subpopulations on the 152 survival of *C. elegans*, 5 repetitions of 50 synchronized young adult nematodes (250 in total), 153 distributed in 5 plates of 10 worms each, were transferred to NGM agar with a lawn of 154 untreated *S. enterica* var Typhimurium (control) (10⁷ CFU/per plate). This was repeated for treated *S. enterica* var Typhimurium populations (*Salmonella* treated once and treated three times with cauliflower by-product extract) (10⁷ CFU/per plate). The worms were maintained at 20 °C during their life cycle (approximately three weeks) and were examined at 48 h intervals. Worms were considered dead when they did not move and did not respond to stimulation (contact with a platinum worm picker).

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The effect of selected *S. enterica* var Typhimurium populations on egg laying of *C. elegans* was also assessed. For this purpose, 10 worms were transferred to NGM plates with a lawn of untreated *S.* Typhimurium or *S.* Typhimurium treated with antimicrobial by-product extract, and the number of eggs laid was counted at 48 h intervals.

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166 **2.5 Statistical analysis of data**

An ANOVA analysis was carried out to detect significant differences (p-value ≤ 0.05) between microbial resistance under the exposure to cauliflower by-product infusion, and phenotypic differences in *C. elegans* infection due to different bacterial populations. Additionally, survival data for *C. elegans* were analysed by using the Kaplan–Meier method, and the hazard function and percentiles of estimated survival distribution were obtained. All statistical analyses were performed using Statgraphics Centurion XII software (Statpoint Technologies, Inc., Warrenton, VA, USA).

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175 **3.Results**

Figure 1 shows the microbial inactivation and growth (\log_{10} cycles) of the initial *S. enterica* var Typhimurium population (10^7 CFU/mL) caused by each cauliflower by-product infusion treatment round. As can be seen graphically, 1.04 \log_{10} cycles of inactivation were achieved after the first cauliflower by-product exposure. However, subsequently exposed cells were

resistant to the antimicrobial cauliflower effect, and even able to grow (≈1 log₁₀ cycle after the
 third exposure).

In a second step, *C. elegans* was used as a host-model organism for infection, fed with *S. enterica* var Typhimurium: untreated (control), treated once (sensitive to cauliflower effect), and treated three times (resistant to cauliflower bactericidal effect and able to grow). Lifespan, movement, and egg-laying were phenotypic factors under study to evidence possible changes in bacterial virulence between untreated and different populations of resistant treated *S. enterica* var Typhimurium bacteria.

188 Figure 2 shows the percentage of surviving nematodes during their life cycle, when they were 189 fed with (i) untreated S. enterica var Typhimurium, (ii) treated once, and (iii) treated three 190 times with cauliflower by-product infusion. The C. elegans survival curve indicated that during 191 their life cycle (until day 18) the number of living nematodes was higher (p-value < 0.05) for C. 192 elegans fed with treated S. enterica var Typhimurium (one or three times) than for C. elegans 193 fed with untreated S. enterica var Typhimurium. Owing to the life cycle of the worm, the 194 number of nematodes decreased with time; but, survival was up to 21 days for worms fed with 195 untreated S. enterica var Typhimurium, and 23 and 25 days for nematodes fed with S. enterica 196 var Typhimurium treated once and three times, respectively. Nevertheless, non-significant 197 differences were found between survival data of nematode populations fed with S. enterica 198 var Typhimurium treated once and three times (p-value > 0.05).

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Survival data were analysed by using a Kaplan–Meier method to obtain the hazard function for different *C. elegans* individuals fed with different *S*. Typhimurium subpopulations (see Figure 3) and percentiles (Table 1). As can be seen in the figure 3, the hazard was always higher when the nematodes were fed with untreated *S*. Typhimurium than when they were fed with treated *S. enterica* var Typhimurium. From day 14 onwards, the hazard increased more slowly

for *C. elegans* fed with *S. enterica* var Typhimurium treated once and three times than for *C. elegans* fed with untreated *S. enterica* var Typhimurium.

Also the percentiles of estimated survival distribution were obtained. Table 1 shows percentiles of estimated survival distribution for each of the *C. elegans* populations. If we focus on the 5 % percentile (percentage of worms surviving for a given time), it can be seen that there are significant differences (p-value < 0.05) between *C. elegans* fed with treated *S. enterica* var Typhimurium (treated once: 19.3 days, and three times: 19.9 days) and untreated *S.* Typhimurium (16.5 days).

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214 Egg lying was also studied as a complementary test. In optimal conditions, C. elegans lays 215 about 300–350 eggs during its life cycle (approximately 21 days) (Labrousse et al., 2000). 216 However, when the nematodes were fed and infected with any of the subpopulations of S. 217 enterica var Typhimurium considered in this study, egg-laying was only maintained until the 5 $^{
m th}$ 218 day. Figure 4 shows a comparison of the number of eggs laid by C. elegans fed with the three 219 different S. enterica var Typhimurium subpopulations for two periods of time (from 0 to 2 days 220 and from 2 to 4 days). In the first period of time (0 to 2 days) significant differences (p-value < 221 0.05) were found in the number of eggs laid by nematodes fed with the three S. Typhimurium 222 subpopulations. Nematodes fed with untreated S. enterica var Typhimurium laid fewer eggs 223 than nematodes fed with S. enterica var Typhimurium treated once. The highest number eggs 224 laid in the first period are corresponding with nematodes fed with S. enterica var Typhimurium 225 treated three times. For the second time period (2 to 4 days) the number of eggs laid by the 226 worms was almost the same in all cases, and lower than the number of eggs laid in the first 227 period of time. These results are important because, despite the effect of S. enterica var 228 Typhimurium on the egg-laying rate during the life cycle of the nematode, it appears that S. 229 enterica var Typhimurium treated with the cauliflower by-product extract was less virulent in

relation to egg laying than the untreated *Salmonella*, thus corroborating the findings observedin the survival data and the Kaplan–Meier analysis.

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233 Discussion

234 Antimicrobial resistance and adaptation is mediated by specific genes encoded for bacterial 235 protection or generally due to spontaneous mutations in the bacterial chromosome. Once the 236 development of resistance has occurred, the mutated gene is directly transferred to the 237 bacteria's progeny during replication (Tiwari and Tiwari, 2011). The results of this study are 238 revealing the capability of S. Typhimurium to develop microbial resistance/adaptation to 239 cauliflower by-product infusion after being exposed to consecutive sub-lethal treatments that 240 inactivated only part of the initial microbial population. Various authors (Di Pasqua et al., 2006; 241 Ultee et al., 2000) have shown specific adaptation of E. coli, S. Senftenberg, S. Typhimurium 242 and Bacillus cereus to sublethal concentrations of essential oils. Kalily et al. (2016) found 243 linalool-resistant mutants of S. Senftenberg. Development of resistance to these natural 244 antimicrobial has been associated to structural and functional changes in microbial cells, 245 specifically in their cell membranes (Ali et al., 2018; Kalily et al., 2016; McMahon et al., 2007). 246 Recent scientific studies have also demonstrated development of microbial resistance of S. 247 enterica var Typhimurium to several antibiotics such as cephalosporins or fluoroquinolones or 248 cross-protection against antibiotics (Kalily et al., 2017; Kariuki et al., 2015; Zanini et al., 2014). 249 Additional studies are also demonstrated the development of resistance when S. enterica var 250 Typhimurium is treated with non-essential oil natural antimicrobials (polyphenols) (Gupta and 251 Birdi, 2017). Under other sub-lethal physical treatments, like Pulsed Electric Fields and High 252 Hydrostatic Pressure, S. enterica var Typhimurium has also demonstrated ability to be adapted 253 (Sanz-Puig et al., 2018)

In order to further understand the mechanism of resistant bacteria infection, and the possible
evident changes in virulence due to this adaptation step in relation to untreated bacteria, the

256 C. elegans in vivo model was used. Up to date, over 40 human pathogens have been analyzed 257 in the *C. elegans* infection model (Battisti et al., 2017). The rapid growth and short generation 258 time of *C. elegans* permit extensive screens for pathogens infective potential, some of the 259 factors identified in these screens have also been shown to play roles in mammalian infections. 260 In the present study, C. elegans was fed with different populations, untreated S. enterica var Typhimurium and cauliflower treated bacteria, once and three times, the last one highly 261 262 resistant to the natural cauliflower antimicrobial exposure. Firstly, the nematode intestinal 263 epithelium should recognize and respond to pathogens in a microbe-specific manner. According to several previous reported studies, the accumulation of the bacteria in the 264 265 intestine is traduced in several pathologies in C. elegans. The survival of C. elegans is 266 dependent on the type and availability of bacteria on which to feed, among other factors 267 (Allen et al., 2015). Although some studies have indicated that the lifespan of worms fed with 268 S. enterica var Typhimurium decreased significantly (Aballay et al., 2000; Labrousse et al., 269 2000), neither of those studies contemplated infecting the worms with naturally occurring resistant subpopulations of S. enterica var Typhimurium, repeatedly exposed to natural 270 271 cauliflower infusion. According to results, it seems that the lifespan of worms fed with treated 272 S. enterica var Typhimurium was longer than the lifespan of worms fed with untreated 273 Salmonella. This behaviour indicating that resistant S. enterica var Typhimurium are not more 274 virulent for C. elegans than untreated type. Ahmed et al. (2016) identify the antimicrobial-275 resistance genes in Salmonella Typhimurium to be mainly blaTEM and floR, detected at 276 frequencies 53 and 73 % respectively in Salmonella isolated from chicken meat. Wild types of 277 Salmonella are specifically capable to produce enterotoxins controlled by the expression of a 278 series of virulence-genes such as stn (Barilli et al., 2018). Antimicrobial exposure can modulate 279 the virulence of *S. enterica* var Typhimurium. In that sense, several reports are supporting the 280 capability of bacteria (Mycobacterium, Escherichia coli, Salmonella enterica, Pseudomonas 281 aeruginosa) to acquire de novo resistance under inadequate treatment regimes (Schroeder et al., 2017). Adaptative resistance (induced genetic changes, enzymatic driven antimicrobial inactivation, changes in the antimicrobial target, changes in cell permeability) is another of the possible explanations for these increased survival of *S. enterica* var *typhimurium* under the repetitive exposure to cauliflower infusion (Brooks and Brooks, 2014; Sanz-Puig et al., 2017b, 2018). In the present study, cauliflower resistant *S. enterica* var *Typhimurium* seems to be less virulent for the fed nematode than untreated one.

288 The increase in the hazard from day 14 is probably related with the age of the nematodes, 289 which are already considered old at that time, and with their immunity system, which is less 290 efficient than in younger nematodes and consequently more sensitive to infection under 291 virulent subpopulations of S. enterica var Typhimurium. Thomas et al. (2004) also found that 292 the rate at which wild-type C. elegans was killed by the bacterial pathogens tested (S. enterica 293 var Typhimurium, among others) increased as nematodes aged, irrespective of the precise 294 mechanism of killing. Similar results were also found by Kurz et al. (2003) in nematodes 295 infected with Serratia marcescens.

296 With regard to egg laying, egg-laying events tend to be clustered in short bursts, or active 297 phases, which are separated by longer inactive phases during which eggs are retained (Hart et 298 al., 2006). Reproductive function begins on about day 5 of adulthood, and reproduction ceases 299 after 10–14 days of adulthood (Christopher and Kerry, 2013). Results of the present study 300 indicated that S. enterica var Typhimurium treated once and three times and untreated S. 301 Typhimurium modified the laying pattern of the worms. For all studied scenarios for infection, egg laying stopped on the 5th day after contact with the pathogen. Previous research studies 302 303 have indicated that live cells of S. Typhimurium are accumulated in the lumen of the intestine of *C. elegans*, that becomes completely infected by the 5th day of contact with this bacterium 304 305 (Aballay et al., 2000; Gardner et al., 2013; Labrousse et al., 2000). Other previous studies have 306 also registered the C. elegans egg-laying pattern as indicator of microbial virulence, showing 307 that under the exposure to E. faecalis, C. elegans increases the egg-in-uterus retention, being

308 the number of eggs retained related with the virulence of *E. faecalis* strain used in each case 309 (Gardner et al., 2013). This retention of progeny observed in the present study is 310 corresponding in time with the progressive accumulation of *S. enterica* var *Typhimurium* in the 311 lumen of the nematode. Additionally, other research studies have demonstrated that S. 312 Typhimurium infection can affect C. elegans egg laying and the eggs hatch internally, which 313 contributes significantly to killing the worms in the first days of their lifespan (Garden et al., 2013; Labrousse et al., 2000). In spite of the fact that egg laying stopped on the 5th day in all 314 315 cases, higher number of eggs was detected for C. elegans fed with resistant bacteria in 316 comparison with egg laying pattern of *C. elegans* fed with untreated *S. Typhimurium*.

In the present study, the survival data, the Kaplan–Meier analysis and the egg-laying behaviour used to assess the effect of infection by resistant bacteria on *C. elegans* revealed that an increase in resistance does not mean that microorganisms become more virulent for *C. elegans*, although *S. enterica* var Typhimurium still maintains some level of virulence, as can be deduced from the egg-laying study. Complex relationship between antimicrobial resistance and virulence genes involved in each specific treatment process should be elucidated in order to develop safest processes and products.

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326 Conclusions

327 The present research work provides valuable information regarding the resistance 328 development of S. enterica var Typhimurium under the repeated exposure to a natural 329 antimicrobial obtained from the agri-food by-products. Valorization of natural compounds 330 from vegetables, and the development of novel ingredients is feasible and demanded by 331 industry and consumers. Potential functionalities can appear related to creation of more 332 convenient, healthy and minimally processed products. However, associated risks should be accurately predicted. No virulence increment by the exposure to cauliflower infusion was 333 334 detected in the present study using one of the most concerning foodborne pathogens using an 335 in vivo test-organism C. elegans.

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337 **Conflict of interest**

The authors declare that there is no conflict of interests regarding the publication of thispaper.

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TABLE 1. Percentiles for *C. elegans* when fed with the different *S.* Typhimurium sub 545 populations.

		Untreated	S. Typhimurium	S. Typhimurium
		S. Typhimurium	treated once	treated three times
	Percentile	Time (days)	Time (days)	Time (days)
	75.0	2.4	3.9	3.3
	50.0	5.3	8.5	8.2
	25.0	10.2	13.9	13.4
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FIGURE 1. Evolution of *S*. Typhimurium resistance when subjected to repeated antimicrobial

treatments with 5% cauliflower by-product infusion (Initial bacterial count 10⁷ CFU/mL).

563 Treated once, Treated two times, Treated three times







Box-and-Whisker Plot



614

- 615 **FIGURE 4.** Eggs laid during first two time intervals by *C. elegans* fed with different *S.*
- 616 Typhimurium populations.
- 617 S= Salmonella without treatment
- 618 SINFE 1= Salmonella treated once
- 619 SINFE 3= Salmonella treated three times
- 620 2 days= Eggs laid at day 2
- 621 4 days= Eggs laid at day 4
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