

Accepted Manuscript

Microbial food safety in the 21st century: emerging challenges and foodborne pathogenic bacteria

Charles M.A.P. Franz, Heidi M.W. den Besten, Christina Böhnlein, Manfred Gareis, Marcel H. Zwietering, Vincenzina Fusco



PII: S0924-2244(17)30380-1

DOI: [10.1016/j.tifs.2018.09.019](https://doi.org/10.1016/j.tifs.2018.09.019)

Reference: TIFS 2326

To appear in: *Trends in Food Science & Technology*

Received Date: 16 June 2017

Revised Date: 9 May 2018

Accepted Date: 11 September 2018

Please cite this article as: Franz, C.M.A.P., den Besten, H.M.W., Böhnlein, C., Gareis, M., Zwietering, M.H., Fusco, V., Microbial food safety in the 21st century: emerging challenges and foodborne pathogenic bacteria, *Trends in Food Science & Technology* (2018), doi: <https://doi.org/10.1016/j.tifs.2018.09.019>.

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

1 **Microbial food safety in the 21st century: emerging challenges and foodborne pathogenic**
2 **bacteria**

3

4 Charles M.A.P. Franz^{1*}, Heidy M.W. den Besten², Christina Böhnlein¹, Manfred Gareis³,
5 Marcel H. Zwietering², and Vincenzina Fusco⁴

6

7 ¹*Max-Rubner-Institut, Department of Microbiology and Biotechnology, Hermann-Weigmann-*
8 *Straße 1, 24103 Kiel, Germany. Email: charles.franz@mri.bund.de*

9 ²*Laboratory of Food Microbiology, Wageningen University and Research, Bornse Weilanden*
10 *9, 6708 WG Wageningen, The Netherlands; Email: heidy.denbesten@wur.nl*

11 ³*Chair of Food Safety, Veterinary Faculty, Ludwig-Maximilians-University Munich,*
12 *Schoenleutnerstr. 8, 85764 Oberschleissheim, Germany; Email: manfred.gareis@lmu.de*

13 ⁴*Institute of Sciences of Food Productions, National Research Council of Italy, (ISPA-CNR),*
14 *Via G. Amendola 122/O, 70126, Bari, Italy. Email: vincenzina.fusco@ispa.cnr.it*

15

16 *Corresponding author

17

18

19 Keywords: food safety, pathogens, microbial risk assessment

20

21 **The challenge of foodborne disease**

22 With billions to feed worldwide, the need to produce adequate amounts of safe food,
23 unadulterated by bacterial, viral and protozoan pathogens, as well as harmful residues,
24 pesticides and allergens, remains one of the major challenges in modern times.

25 According to the World Health Organisation, unsafe food containing harmful bacteria,
26 viruses, parasites or chemical substances, causes more than 200 diseases – ranging from

27 diarrhoea to cancers. An estimated 600 million, i.e. almost 1 of 10 people in the world, fall ill
28 after eating contaminated food and 420,000 die every year, resulting in the loss of 33 million
29 healthy life years (DALY's = disability adjusted life years) (WHO, 2015a, b). Diarrhoeal
30 diseases are the most common illnesses resulting from the consumption of contaminated food
31 (WHO, 2015a, b). A data synthesis (Kirk et al., 2015) on the global and regional disease
32 burden of 22 foodborne diseases in 2010 estimated that these caused 580 million foodborne
33 illnesses in 2010. Norovirus alone was responsible for 125 million foodborne illnesses, the
34 largest number for any pathogen. Other pathogens resulting in high numbers of foodborne
35 cases were *Campylobacter* spp., non-typhoidal *Salmonella* spp., Enterotoxinogenic *E. coli*,
36 Enteropathogenic *E. coli*, STEC and *Shigella* spp. (Table 1) (Kirk et al., 2015).

37 Looking at the European situation, zoonoses monitoring activities carried out in 2016 in
38 37 European countries found campylobacteriosis the most commonly reported zoonosis,
39 followed by salmonellosis, yersiniosis, Shiga Toxin-producing *Escherichia coli* (STEC)
40 infections and listeriosis (Table 1). However, while the increasing EU trend for human
41 campylobacteriosis cases since 2008 stabilised during 2012-2016, within the same period the
42 decreasing EU trend for confirmed human salmonellosis cases ended, due to the recent
43 *Salmonella* Enteritidis outbreaks, accounting for 59% of all salmonellosis cases in EU (EFSA
44 and ECDC, 2017). On the other hand, the number of confirmed STEC infections in humans
45 remained stable whereas the decreasing EU trend of confirmed cases of yersiniosis since 2008
46 stabilised during 2012-2016. Moreover, a further increased number of confirmed human
47 listeriosis cases was registered in 2016 (EFSA and ECDC, 2017). Of the 4,786 weak- and
48 strong-evidence foodborne and waterborne outbreaks reported in 2016 by 27 member states,
49 bacteria were the most commonly detected causative agents of zoonoses (33.9%), followed by
50 bacterial toxins (17.7%), viruses (9.8%), other causative agents (2.2%) and parasites (0.4%).
51 Hereby *Salmonella* was accounting for 65% of the outbreaks caused by bacterial agents. The

52 main foods involved in the strong-evidence outbreaks were from foods of animal origin and
53 these were from ‘eggs’ (23.0%), ‘poultry meat’ (18.5%), ‘fish and fisheries’ including
54 ‘crustaceans, shellfish, molluscs and its products’ (22.4%), ‘meat and meat products other
55 than poultry’ (21.7%), and ‘milk and milk products’ (14.4%), while one-third of all strong-
56 evidence outbreaks involved ‘buffet meals’, ‘mixed food’ and ‘other foods’ including
57 ‘unspecified foods’.

58

59 **Rising to the challenge**

60 Acknowledging that we have made considerable progress in taking action for increasing
61 food safety in the last 15 years, we still have considerably high numbers of illnesses and
62 hence risks associated with the consumption of food, and the disease burden still is high (Kirk
63 et al., 2015). Moreover, because about 20% of the population of the United States and the
64 United Kingdom belong to the so-called “vulnerable people” (Lund and O’Brien, 2011; Lund,
65 2015) (especially the very young, the elderly and immunocompromised), we are more pressed
66 for finding solutions for increasing food safety. A “from farm-to-fork” approach of food
67 safety along the whole food chain has been adopted by many countries already a number of
68 years ago (EU, 2014). Recognizing that the farm-to-fork approach may not be sufficient, in
69 the last years the “one-health-initiative” emerged, stating that we have to start at the farm
70 level, with pathogen-controlled feed and with healthy livestock to assure food safety, as well
71 as with a healthy environment (Kahn, 2017). This came with the realisation that the health of
72 livestock affects human health, especially in connection with antibiotic resistant bacteria
73 (including foodborne pathogens). Use and misuse of antibiotics in both humans and animals
74 are responsible for the development of resistant bacteria (WHO, 2018) and antibiotic
75 resistance is therefore an important topic within the One-Health initiatives. Especially misuse

76 or overuse of antibiotics in animal husbandry could finally result in resistant bacteria
77 occurring in the food chain. Thus, although progress towards safe food production has been
78 made, new emerging challenges arise at the consumer, microorganisms or food processor
79 levels (Fig. 1), which require us to re-think food safety and keep a constant vigil for emerging
80 threats.

81

82 **Emerging challenges**

83 Having considered food safety from the ‘one health’ and the ‘farm-to-fork’ approaches,
84 one challenge remains at the level of the consumer, particularly the *vulnerable consumer*.
85 The very young may be particularly at risk, because of the immaturity of their immune and
86 physiologic systems (IUFoST, 2015). For the elderly (25% of the European population in
87 2017) and whose number is projected to further increase worldwide from the estimated 962
88 million in 2017, to 1.4 billion in 2030 and 2.1 billion in 2050 (United Nations, 2017),
89 weakness of the immune system also increases vulnerability. The vulnerable are also those
90 having poor nutritional status, existing health problems, and drug therapies which suppress
91 the immune system (IUFoST, 2015; Lund and O’Brien, 2011; Newman et al., 2015). Such
92 persons are more likely to acquire foodborne illness and are prone to more severe disease
93 outcomes, including higher mortality rates (IUFoST, 2015). The challenge will be to produce
94 foods with low microbial risks, to define and exclude high risk foods and to disseminate clear
95 advice about food safety.

96 The ‘one health’ initiative quite rightly connects environmental and animal health with
97 human health. Changes in the agri-food chain, social changes and advances in the detection
98 and reporting systems, coupled with bacterial adaptation and evolution, may lead to certain
99 microorganisms becoming new or *emerging zoonotic pathogens*. Examples of such include

100 shigatoxigenic/ enterohaemorrhagic *E. coli* (STEC/EHEC) and *Campylobacter* spp. in the
101 meat chain, *Listeria monocytogenes* in vegetable, meat or milk products, *Cronobacter* spp. in
102 infant milk formula, *Arcobacter* spp., *Yersinia enterocolitica* serobiotype O3/4, parasites such
103 as *Cyclospora* on fruit and *Cryptosporidium* and *Giardia* in water, as well as hepatitis E virus
104 in pork and boar meat (Duffy et al., 2008; Batzilla et al., 2011; Park et al., 2016; Ramees et
105 al., 2017). Recognising these current zoonotic pathogens and their potential for foodborne
106 transmission will be essential for identifying emerging foodborne pathogens.

107 Viruses (adeno-, calici- and enterioviruses) are important pathogens which in many
108 countries are the most numerous causes (norovirus) for foodborne infection. For adenoviruses
109 or caliciviruses no standardized methods for cultivation or detection exist. While standardized
110 procedures for cultivation of some enteroviruses exist, these methods are not capable of
111 distinguishing between virus types and are not applicable for all enteroviruses (Hartmann and
112 Halden, 2012). Detection is also challenging because viruses have a high mutation rate and
113 many have a high probability of infection even at 10 virions (Hartmann and Halden 2012).
114 This has obvious implications regarding the difficulty for the ***detection and monitoring of***
115 ***foodborne viruses***. Here, methods for virion concentration, as well as sensitive molecular
116 biological or serological methods, or even mass spectrometry, need to be developed for an
117 accurate and specific detection at low contamination levels.

118 Decreasing the excess use of antibiotics in animal husbandry and in human medicine is
119 especially important to decrease the occurrence and spread of ***antibiotic resistant bacteria***.
120 Yet antibiotic use cannot be decreased to zero in the interest of human and animal health. It
121 will be important, therefore, to define points of pathogen entry, trace transmission routes
122 along the food chain, to determine the evolution of transferable antibiotic genes and more
123 importantly to find control measures which prevent or diminish the entry and spread of
124 resistant microorganisms or resistance genes. Here, not only true foodborne pathogens are of

125 importance, but also opportunistic pathogens such as *Klebsiella* spp., *Enterobacter* spp.,
126 *Citrobacter* spp. and *Serratia* spp. These are well known to occur in various foods (e.g. meats,
127 vegetables, milk) and to cause hospital infections (Nordmann et al., 2012; Fusco et al., 2018).
128 Additionally, even non-pathogenic bacteria may become antibiotic resistant and can be
129 relevant in spread. The challenge here is to monitor the spread and evolution of such bacteria
130 to prove an animal/environment/human connection. One approach may be a syst-OMICS
131 approach, as was recently reported to be adopted for salmonellosis to ensure food safety and
132 reduce the economic burden. The study by Emold-Rheault et al. (2017) sets out to sequence
133 the genomes of 4500 *Salmonella* genomes and to build an analysis pipeline for the study of
134 *Salmonella* genome evolution, antibiotic resistance and virulence genes. This way, the study
135 aims to draw potential links between strains found in fresh produce, humans, animals and the
136 environment (Emold-Rheault et al., 2017). A similar approach would be worthwhile for
137 adopting for other bacterial pathogens such as *Campylobacter*, *Listeria monocytogenes*,
138 pathogenic *E. coli* strains, or the opportunistic pathogens mentioned above.

139 ***Climate change*** may well be important for microbial food safety in the 21st century.
140 There is reasonable evidence that the environment and weather play a role in the transmission
141 of e.g. *Salmonella* and *Campylobacter* spp. to humans, even though there is uncertainty about
142 the mechanisms behind this (Justus et al., 2017; Lake, 2017; Nichols et al., 2018). Possibly
143 global warming may have such an effect on increased transmission also with other pathogens,
144 or may even become a key factor in selecting for other emerging pathogens. Food will also be
145 produced in altered climatic conditions in modified surrounding ecosystems, and the
146 interactions between these changes and the food production systems are complex and
147 uncertain (Lake and Barker, 2018). For example, increased indoor animal husbandry to
148 counteract heat stress may elevate the potential for animal to animal transmission of zoonotic
149 pathogens. Increased growing seasons may lead to greater use or outdoor pastures and

150 increase the probability of transmission of pathogens from the environment. Flooding or
151 drought may favour the spread of pathogens to produce, or have consequences on water
152 quality and pathogen transmission (Lake and Barker, 2018). Another important aspect
153 concerns the increasing water shortage and the worldwide demand for fresh water. As a result,
154 an increase in the use of waste water for irrigation and sewage sludge could be expected,
155 accompanied by increasing risks of contamination of agricultural land and plants with
156 pathogens.

157 Research into *novel food preservation methods* (or technologies) remains a challenge,
158 particularly when considering the production of foods with low microbial diets for vulnerable
159 people. Against this background we need to discuss whether all food needs to be made
160 suitable and available to the vulnerable, or whether specific safe diets need to be formulated
161 or especially produced? Specific preservation technologies that have been researched and to
162 some extent applied in the last years include high hydrostatic pressure, pulsed electric fields,
163 high voltage arc discharge and cold plasma (Stoica et al., 2013), as well as pulsed light or
164 UV-C treatments. One promising biocontrol tool would also be the use of lytic bacteriophages
165 to specifically control pathogens or antibiotic resistant opportunistic pathogens. Due to their
166 host specificity, lytic bacteriophages would act very target specific (Jordan et al., 2014). This
167 would be of obvious advantage also for use in specific foods suitable for this technology, in
168 which a pathogen of concern needs to be inactivated to improve its safety for the vulnerable
169 people group.

170 Interestingly, the Executive Summary of Food Safety by the EFSA (2009), reporting on a
171 survey of consumer risk perception showed that the consumer is more likely to worry about
172 risks caused by external factors, over which they have no control, e.g. consumers expressed
173 concern regarding contamination of food by bacteria and unhygienic conditions outside home.
174 On the other hand, they seemed less concerned about factors linked to their own behaviour

175 (e.g. food preparation, food hygiene at home). Apart from such optimistic bias and illusion of
176 control, other reasons for unsafe food preparation by the consumer were shown to include
177 habits and lack of knowledge concerning food safety during domestic food preparation, as
178 well as disagreement with some recommendations for safe food handling (Al-Sakkaf, 2012;
179 Young and Waddell, 2016). Regarding the latter for example, a study by Kosa et al. (2015) on
180 consumer-reported handling of poultry products at home showed that there was low
181 adherence to current recommended food safety practices by the consumers regarding that they
182 should not wash raw poultry before cooking, proper refrigerator storage of raw poultry, use of
183 a food thermometer to determine doneness, and proper thawing of raw poultry in cold water.
184 Clearly, therefore, risk assessment agencies or communicators should in future spend more
185 effort in gathering and utilizing such information to develop and update science-based
186 education materials.

187 **Microbial Risk Assessment**

188 Risk assessment is a science-based process consisting of hazard identification, hazard
189 characterization, exposure assessment, and risk characterization (CAC, 2014). Microbial risk
190 assessment (MRA) can largely help to understand the behaviour of pathogens over a food
191 chain, to predict health risks and the expected public health effects of interventions and
192 standards (Havelaar et al. 2010). For risk assessment studies, many quantitative data are
193 needed, like prevalences of foodborne pathogens, characteristics of organisms, food products
194 and processes, virulence of organisms and susceptibility of humans, as well as public health
195 and epidemiological data. In the last decades, more and more of these data became available,
196 not always perfect, but the quantity and the availability of data has increased largely. This
197 information is even in certain cases overwhelming (big data), not only regarding
198 microorganisms characteristics (genomics) and behaviour (transcriptomics and
199 metabolomics), but also the tenacity (survival or dying) or growth with regard to products,

200 processes, intrinsic and extrinsic factors of foods, and even human behaviour. The exact
201 meaning of data and defining its quality and applicability can then become problematic.
202 Difficulty apart from searching, collecting, defining, interpreting and valuing the data sources
203 is also how to make use of it, since large variability and uncertainty exist (Zwietering, 2015,
204 Koutsoumanis & Aspridou, 2016, Membré & Guillou, 2016). Adaptation and evolution of
205 microorganisms within a changing environment might affect the genotypes or lineages of
206 pathogens which become problematic. Thus genotype-specific risk assessment (Carlin et al.,
207 2013) and individual cell-based modelling (Koutsoumanis, 2008; Metselaar et al., 2016) are
208 becoming increasingly important. These approaches might contribute to fine-tune the hazard
209 identification, hazard characterization and exposure assessment elements of microbiological
210 risk assessments and thereby reducing the uncertainty in risk characterizations (see for a more
211 in-depth discussion e.g. Cocolin et al., 2018; Den Besten et al., 2018; Haddad et al., 2018;
212 Membré & Guillou, 2016; Pielaat et al., 2015; Rantsiou et al., 2018). Reports with data,
213 databases (e.g. Combase, <http://www.combase.cc>), and many tools are developed (e.g. PMM-
214 Lab, <https://foodrisklabs.bfr.bund.de/pmm-lab/>, Baseline, <http://www.baselineapp.com/>) that
215 can make implementation of risk assessments more available for more people. Ultimately the
216 integration of genotypic data that can be obtained with omics technologies and quantitative
217 phenotypic data (i.e. quantitative descriptors for growth, survival and inactivation for
218 genotypes and heterogeneity between individual cells) and simulation tools and experimental
219 challenge tests make it possible to get better grip on magnitudes and sources of risks. This is
220 needed to evaluate various ways to effectively control the microbial risks with technical
221 solutions, behavioural changes, changes in product formulations and in standards and
222 legislation, for a balanced control of hazards in our foods.

223

224

225 **References**

- 226 Al-Sakkaf, A. (2012). Evaluation of food handling practice among New Zealanders and other
227 developed countries as a main risk factor for campylobacteriosis rate. *Food Control*, 27,
228 330-337.
- 229 CAC, 2016. Codex Alimentarius Commission PROCEDURAL MANUAL. Twenty-fifth ed.
230 [http://www.fao.org/fao-who-codexalimentarius/sh-](http://www.fao.org/fao-who-codexalimentarius/sh-proxy/en/?lnk=1&url=https%253A%252F%252Fworkspace.fao.org%252Fsites%252Fcodex%252FShared%20Documents%252FPublications%252FProcedural%20Manual%252FManual_25%252FManual_25e.pdf)
231 [proxy/en/?lnk=1&url=https%253A%252F%252Fworkspace.fao.org%252Fsites%252Fcodex%252FShared%20Documents%252FPublications%252FProcedural%20Manual%252FManual_25%252FManual_25e.pdf](http://www.fao.org/fao-who-codexalimentarius/sh-proxy/en/?lnk=1&url=https%253A%252F%252Fworkspace.fao.org%252Fsites%252Fcodex%252FShared%20Documents%252FPublications%252FProcedural%20Manual%252FManual_25%252FManual_25e.pdf).
232
233
- 234 Carlin F., Albagnac C., Rida A., Guinebretière M.-H., Couvert O. & Nguyen-the C. (2013).
235 Variation of cardinal growth parameters and growth limits according to phylogenetic
236 affiliation in the *Bacillus cereus* Group. Consequences for risk assessment. *Food*
237 *Microbiology*, 33, 69-76
- 238 Cocolin L., Mataragas M., Bourdichon F., Doulgeraki A., Pilet M.-F., Jagadeesan B.,
239 Rantsiou K. & Phister, T. (2018). Next generation microbiological risk assessment meta-
240 omics: The next need for integration. *International Journal of Food Microbiology*. *In*
241 *press*, <https://doi.org/10.1016/j.ijfoodmicro.2017.11.008>
- 242 Den Besten H.M.W., Amézquita A., Bover-Cid, S., Dagnas S., Ellouze M., Guillou S.,
243 Nychas G., O'Mahony C., Pérez-Rodríguez F. & Membré J.M. (2018). Next generation of
244 microbiological risk assessment: Potential of omics data for exposure assessment.
245 *International Journal of Food Microbiology*, *In press*.
246 <https://doi.org/10.1016/j.ijfoodmicro.2017.10.006>

- 247 EFSA (European Food Safety Authority) and ECDC (European Centre for Disease Prevention
248 and Control) (2017). The European Union summary report on trends and sources of
249 zoonoses, zoonotic agents and food-borne outbreaks in 2016. *EFSA Journal*, 15(12), 5077.
- 250 Emond-Rheault, J.-G., Jeukens, J., Freschi, L., Kukavica-Ibrulj, I., Boyle, B., Dupont, M.-J.,
251 Colavecchio, A., Barrere, V., Cadieux, B., Arya, G., Bekal, S., Berry, C., Burnett, E.,
252 Cavestri, C., Chapin, T.K., Crouse, A., Daigle, F., Danyluk, M.D., Delaquis, P., Dewar, K.,
253 Doualla-Bell, F., Fliss, I., Fong, K., Fournier, E., Franz, E., Garduno, R., Gill, A.,
254 Gruenheid, S., Harris, L., Huang, C.B., Huang, H., Johnson, R., Joly, Y., Kerhoas, M.,
255 Kong, N., Lapointe, G., Lariviere, L., Loignon, S., Malo, D., Moineau, S., Mottawea, W.,
256 Mukhopadhyay, K., Nadon, C., Nash, J., Feze, I.N., Ogunremi, D., Perets, A., Pilar, A.V.,
257 Reimer, A.R., Robertson, J., Rohde, J., Sanderson, K.E., Song, L., Stephan, R., Tamber, S.,
258 Thomassin, P., Tremblay, D., Usongo, V., Vincent, C., Wang, S., Weadge, J.T.,
259 Wiedmann, M., Wijnands, L., Wilson, E.D., Wittum, T., Yoshida, C., Youfsi, K., Zhu, L.,
260 Weimer, B.C., Goodridge, L. & Levesque, R.C. (2017). A syst-OMICS approach to
261 ensuring food safety and reducing the economic burden of salmonellosis. *Frontiers in*
262 *Microbiology*, 8, 996.
- 263 EU (European Union) (2014). From farm to fork. Safe and healthy food for everyone.
264 [https://publications.europa.eu/en/publication-detail/-/publication/946e612c-6b31-4805-](https://publications.europa.eu/en/publication-detail/-/publication/946e612c-6b31-4805-89bc-b0c90fa881cf/language-en/format-PDF/source-69594759)
265 [89bc-b0c90fa881cf/language-en/format-PDF/source-69594759](https://publications.europa.eu/en/publication-detail/-/publication/946e612c-6b31-4805-89bc-b0c90fa881cf/language-en/format-PDF/source-69594759)
- 266 Fusco, V., Abriouel, H., Benomar, N., Kabisch, J., Chieffi, D., Cho, G.-S., & Franz, C.M.A.P.
267 (2018). Opportunistic foodborne pathogens. In: Food safety and preservation: modern
268 biological approaches to improving consumer health (chapter 10). 1st Edition. Editors:
269 Alexandru Grumezescu Alina Maria Holban. ISBN: 9780128149560. Academic Press.

- 270 <https://www.elsevier.com/books/food-safety-and-preservation/grumezescu/978-0-12->
271 814956-0
- 272 Haddad N., Johnson N.B., Kathariou S., Metris A., Phister T., Pielaat A., Tassou C., Wells-
273 Bennik M.H.J. & Zwietering M.H. (2018). Next generation microbiological risk
274 assessment – Potential of omics data for hazard characterisation. *International Journal of*
275 *Food Microbiology*. *In press*. <https://doi.org/10.1016/j.ijfoodmicro.2018.04.015>.
- 276 Havelaar, A.H., Brul, S., Jong, A., de Jonge, R., Zwietering, M.H., & ter Kuile, B.H. (2010).
277 Future Challenges to Microbial Food Safety. *International Journal of Food Microbiology*
278 139, S79-S94
- 279 IUFOST (International Union of Food Science and Technology) (2015). Foodborne disease
280 and vulnerable groups. IUFOST Scientific Information Bulletin March 2015.
281 [http://www.iufost.org/iufostftp/IUFOST%20SIB%20%20-](http://www.iufost.org/iufostftp/IUFOST%20SIB%20%20-%20Foodborne%20Disease%20and%20Vulnerable%20Groups%20March%202015.pdf)
282 [%20Foodborne%20Disease%20and%20Vulnerable%20Groups%20March%202015.pdf](http://www.iufost.org/iufostftp/IUFOST%20SIB%20%20-%20Foodborne%20Disease%20and%20Vulnerable%20Groups%20March%202015.pdf)
- 283 Jordan, K., Dalmaso, M., Zentek, J., Mader, A., Bruggeman, G., Wallace, J., De Medici, D.,
284 Fiore, A., Prukner-Radovnik, E., Lukac, M., Axelsson, L., Holck, A., Ingmer, H., &
285 Malakauskas, M. (2014). Microbes versus microbes: control of pathogens in the food
286 chain. *Journal of the Science of Food and Agriculture*, 94, 3079-3089.
- 287 Kahn, L.H. (2017). The one-health way. *Nature*, 543, S47.
- 288 Kirk, M.D., Pires, S.M., Black, R.E., Caipo, M., Crump, J.A., Devleeschauwer, B., Döpfer,
289 D., Fazil, A., Fischer-Walker, C.L., Hald, T., Hall, A.J., Keddy, K.H., Lake, R.J., Lanata,
290 C.F., Torgerson, P.R., Havelaar, A.H., & Angulo, F.J. (2015). World Health Organisation
291 estimates of the global and regional disease burden of 22 foodborne bacterial, protozoal,
292 and viral diseases, 2010: A data synthesis. *PLOS Medicine* 12, e1001921.

- 293
- 294 Kosa, K.M., Cates, S.C., Bradley, S., Chambers, E., & Godwin, S. 2015. Consumer-reported
295 handling of raw poultry products at home: Results from a national survey. *Journal of Food*
296 *Protection*, 78, 180-186.
- 297 Koutsoumanis, K. (2008). A study on the variability in the growth limits of individual cells
298 and its effect on the behavior of microbial populations. *International Journal of Food*
299 *Microbiology*, 128 (1), 116-121.
- 300 Koutsoumanis, K.P. & Aspidou, Z. (2016). Moving towards a risk-based food safety
301 management. *Current Opinion in Food Science*, 12, 36-41.
- 302 Lake I.R. & Barker G.C. (2018). Climate change, foodborne pathogens and illness in higher-
303 income countries. *Current Environmental Health Reports* 5, 187-196.
- 304 Lund, B.M. & O'Brien, S.J. (2011). The occurrence and prevention of foodborne disease in
305 vulnerable people. *Foodborne Pathogens and Disease*, 8, 961-973.
- 306 Metselaar K.I., Abee, T., Zwietering M.H. & den Besten H.M.W. (2016). Modeling and
307 validation of the ecological behavior of wild-type *Listeria monocytogenes* and stress-
308 resistant variants. *Applied and Environmental Microbiology*, 82 (17), 5389-5401.
- 309 Membré, J.M. & Guillou, S. (2016). Latest developments in foodborne pathogen risk
310 assessment. *Current Opinion in Food Science*, 8, 120-126.
- 311 Newman K.L., Leon J.S., Rebolledo P.A., & Scallan E. (2015). The impact of socioeconomic
312 status on foodborne illness in high-income countries: a systematic review. *Epidemiology*
313 *and Infection*, 143: 2473-2485.

- 314 Nordmann, P., Dortet, L., and Poirel, L. (2012). Carbapenem resistance in Enterobacteriaceae:
315 here is the storm! *Trends in Molecular Medicine*, 18(5), 263-272.
- 316 Pielaat A., Boer M.P., Wijnands L.M., van Hoek A.H.A.M, Bouw E., Barker G.C., Teunis
317 P.F.M., Aarts H.J.M. & Franz E. (2015). First step in using molecular data for microbial
318 food safety risk assessment; hazard identification of *Escherichia coli* O157:H7 by coupling
319 genomic data with in vitro adherence to human epithelial cells. *International Journal of*
320 *Food Microbiology*, 213, 130-138.
- 321 Rantsiou K., Kathariou S., Winkler A., Skandamis P., Saint-Cyr M.J., Rouzeau-Szynalski K.
322 & Amézquita A. (2018). Next generation microbiological risk assessment: opportunities
323 of whole genome sequencing (WGS) for foodborne pathogen surveillance, source tracking
324 and risk assessment. *International Journal of Food Microbiology*. *In press*.
325 <https://doi.org/10.1016/j.ijfoodmicro.2017.11.007>.
- 326 Stoica, M., Mihalcea, L., Borda, D., & Alexe, P. (2013). Non-thermal novel food processing
327 technologies. An overview. *Journal of Agroalimentary Processes and Technologies*, 9,
328 212-217.
- 329 United Nations (2017). World Population Prospects: The 2017 Revision, Key Findings and
330 Advance Tables. United Nations, Department of Economic and Social Affairs, Population
331 Division, New York.
332 <http://www.un.org/en/development/desa/population/publications/index.shtml>
- 333 WHO (World Health Organisation) (2014). Advancing food safety initiatives. Strategic plan
334 for food safety including foodborne zoonoses 2013-2022.
335 http://apps.who.int/iris/bitstream/10665/101542/1/9789241506281_eng.pdf?ua=1

- 336 WHO (World Health Organisation) (2015a). Food Safety. Fact sheet no. 399.
337 <http://www.who.int/mediacentre/factsheets/fs399/en/>
- 338 WHO (World Health Organisation) (2015b). WHO estimates of the global burden of
339 foodborne diseases. Foodborne diseases burden epidemiology reference group 2007-2015.
340 http://www.who.int/foodsafety/publications/foodborne_disease/fergreport/en/
- 341 WHO (World Health Organisation) (2018). Antibiotic resistance.
342 [http://www.euro.who.int/en/health-topics/disease-prevention/antimicrobial-](http://www.euro.who.int/en/health-topics/disease-prevention/antimicrobial-resistance/antibiotic-resistance)
343 [resistance/antibiotic-resistance](http://www.euro.who.int/en/health-topics/disease-prevention/antimicrobial-resistance/antibiotic-resistance)
- 344 Young I. & Waddell L. (2016). Barriers and facilitators to safe food handling among
345 consumers: A systematic review and thematic synthesis of qualitative research studies.
346 *PLoS One*, 11(12), e0167695.
- 347 Zwietering, M.H. (2015). Risk assessment and risk management for safe foods: Assessment
348 needs variability and uncertainty, management needs discrete decisions. *International*
349 *Journal of Food Microbiology* 213, 118-123.

350
351
352

353 Table 1: Estimated global cases and reported European cases of food borne illness

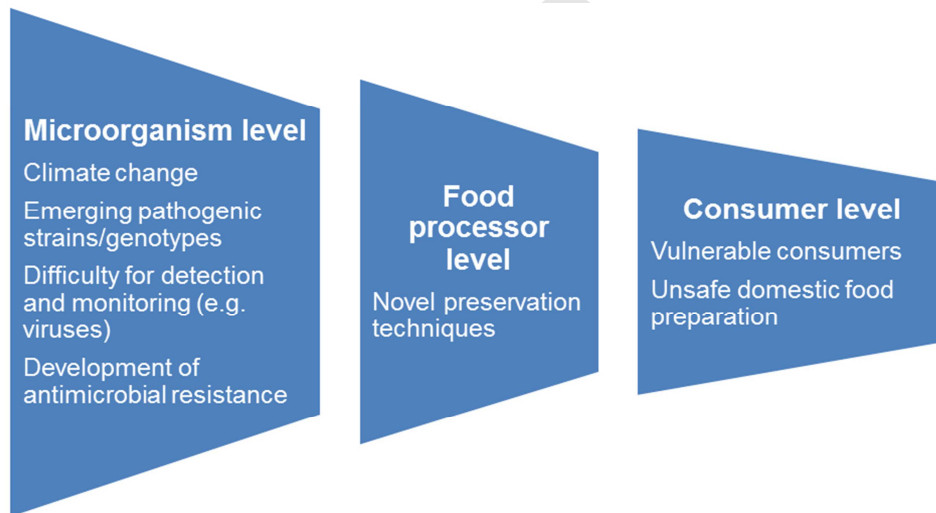
354

Bacterial zoonotic pathogen	Estimated global cases of foodborne illnesses in 2010 (Kirk et al., 2015)	Reported European cases of foodborne illnesses in 2016 (EFSA and ECDC, 2017)
<i>Campylobacter</i> spp.	95 613 970	246 307
Non-typhoidal <i>Salmonella</i> spp.	78 439 785	94 530
Enterotoxinogenic <i>E. coli</i>	86 502 735	n.r.*
Enteropathogenic <i>E. coli</i>	23 797 284	n.r.*
STEC	1 176 854	6 378
<i>Shigella</i> spp.	51 014 050	n.r.*
<i>Listeria monocytogenes</i>	14 169	2 536
<i>Yersinia</i>	n.r.*	6 861

355 *n.r.: not reported

356

357



358

359

360 **Figure Legend**

361

362 **Figure 1:** Challenges for food safety emerging at the consumer, microorganisms or food
 363 processor levels

364

ACCEPTED MANUSCRIPT

Highlights

High number of infection are still caused by foodborne microorganisms

Increasing number of vulnerable people needs safer food

emergence and spread of antibiotic resistant bacteria should be controlled

new methods for effective food preservation are needed

magnitudes and sources of risks and ways to effectively control these are needed