This is a "Post-Print" accepted manuscript, which has been published in "International Journal of Food Microbiology".

Please cite this publication as follows:

Havelaar, A.H., Brul, S., De Jong, A., De Jonge, R., Zwietering, M.H., Ter Kuile, B.H. (2010) Future challenges to microbial food safety. International Journal of Food Microbiology 139 (Suppl 1), S79-S94.

You can download the published version at: <u>http://dx.doi.org/10.1016/j.ijfoodmicro.2009.10.015</u>

Confidential manuscript submitted to International Journal of Food Microbiology

1 Future Challenges to Microbial Food Safety

- 2 Arie H. Havelaar^{*1,2}, Stanley Brul³, Aarieke de Jong⁴, Rob de Jonge¹, Marcel H.
- 3 Zwietering⁵ and Benno H. ter Kuile^{3,6}
- 4 ¹ Laboratory for Zoonoses and Environmental Microbiology, Centre for Infectious Diseases Control
- 5 Netherlands, National Institute for Public Health and the Environment, PO Box 1, 3720 BA Bilthoven, the

6 Netherlands

- 7 ² Division of Veterinary Public Health, Institute for Risk Assessment Sciences, Utrecht University, PO Box
- 8 80175, 3508 TD Utrecht, the Netherlands
- 9 ³ Laboratory for Molecular Biology and Microbial Food Safety, Netherlands Institute for Systems Biology,
- 10 Swammerdam Institute for Life Sciences, University of Amsterdam, Nieuwe Achtergracht 166, 1018 WV
- 11 Amsterdam, the Netherlands
- ⁴ Dutch Food and Consumer Product Safety Authority, Hoogte Kadijk 401, 1018 BK Amsterdam, the
 Netherlands
- ⁵ Laboratory of Food Microbiology, Wageningen University, PO Box 8129, 6700 EV, Wageningen, the
 Netherlands
- ⁶ Dutch Food and Consumer Product Safety Authority, Office for Risk Assessment, PO Box 19506, 2500 CM
- 17 The Hague, the Netherlands
- 18 * Corresponding author. Tel.: +31 30 2742826, fax: +31 30 2744434.
- 19 E-mail address: arie.havelaar@rivm.nl

20

Confidential manuscript submitted to International Journal of Food Microbiology

22 Abstract

23 Despite significant efforts by all parties involved, there is still a considerable burden of 24 foodborne illness, in which microorganisms play a prominent role. Microbes can enter the food chain at different steps, are highly versatile and can adapt to the environment allowing 25 survival, growth and production of toxic compounds. This sets them apart from chemical 26 agents and thus their study from food toxicology. We summarize the discussions of a 27 28 conference organized by the Dutch Food and Consumer Products Safety Authority and the 29 European Food Safety Authority. The goal of the conference was to discuss new challenges to 30 food safety that are caused by microorganisms as well as strategies and methodologies to 31 counter these. Management of food safety is based on generally accepted principles of Hazard 32 Analysis Critical Control Points and of Good Manufacturing Practices. However, a more pro-33 active, science-based approach is required, starting with the ability to predict where problems 34 might arise by applying the risk analysis framework.

Developments that may influence food safety in future occur on different scales (from global 35 to molecular) and in different time frames (from decades to less than a minute). This 36 necessitates development of new risk assessment approaches, taking the impact of different 37 drivers of change into account. We provide an overview of drivers that may affect food safety 38 39 and their potential impact on foodborne pathogens and human disease risks. We conclude that 40 many drivers may result in increased food safety risks, requiring active governmental policy 41 setting and anticipation by food industries whereas other drivers may decrease food safety 42 risks.

43 Monitoring of contamination in the food chain, combined with surveillance of human illness
44 and epidemiological investigations of outbreaks and sporadic cases continue to be important
45 sources of information. New approaches in human illness surveillance include the use of

Confidential manuscript submitted to International Journal of Food Microbiology

- 46 molecular markers for improved outbreak detection and source attribution, sero-epidemiology
- 47 and disease burden estimation.

48 Current developments in molecular techniques make it possible to rapidly assemble 49 information on the genome of various isolates of microbial species of concern. Such 50 information can be used to develop new tracking and tracing methods, and to investigate the 51 behavior of microorganisms under environmentally relevant stress conditions. These novel tools and insight need to be applied to objectives for food safety strategies, as well as to 52 53 models that predict microbial behavior. In addition, the increasing complexity of the global 54 food systems necessitates improved communication between all parties involved: scientists, risk assessors and risk managers, as well as consumers. 55

56

57 Keywords: food safety, microbial hazards, future trends, monitoring, surveillance, risk

58 assessment, tracking and tracing, communication

Confidential manuscript submitted to International Journal of Food Microbiology

60 **1. Introduction**

61 The microbiological aspects of food safety have been studied intensively for many decades. In 62 the Netherlands, the standard of food safety has increased in the last decades (Van Kreijl et al., 2006), and political attention is shifting to other food-related problems such as obesity and 63 64 unhealthy diets. However, even in industrialized countries, there is still a considerable burden of foodborne illness. For example, in the Netherlands there are an estimated 700.000 cases of 65 illness and 80 deaths per year. The burden of foodborne disease for this country is at least 66 3800 Disability Adjusted Life Years and 65 million Euro per year (Havelaar et al., 2008). 67 Also other industrialized countries report a continuing burden of foodborne illness (Flint et 68 69 al., 2005, Hall et al., 2005, Adak et al., 2005, Anonymous 2007a, Jones et al., 2007). 70 Foodborne outbreaks appear to be on the rise again in some industrialized countries, with a 71 shift from traditional problems with foods from animal origin to fresh foods such as produce 72 (Anonymous, 2008a), shellfish (Pontrelli et al., 2008) and dry products and ingredients (e.g. peanuts, Anonymous, 2009). Furthermore, new threats continue to be identified. The attention 73 for viruses is more recent, but no less relevant. New risks are being encountered because of 74 changing characteristics of the relevant micro-organisms, changing production 75 76 methodologies, changes in the environment and the ecology, and an increase of the global 77 trade of foodstuffs. In addition, demands on food safety increase steadily. Due to the nature of 78 microbes and our food chain, measures to ensure food safety have to be implemented on a 79 global scale, necessitating a global approach. 80 To discuss the challenges that microbes pose to food safety on the longer term, the Dutch 81 Food and Consumer Products Safety Authority (VWA) and the European Food Safety

82 Authority (EFSA) organized a conference on "Future Challenges to Food Safety" (Wolfheze,

- the Netherlands, 9-12 June 2008). The goal of the conference was to discuss new challenges
- to food safety that are caused by microbes and strategies and methodologies to counter these.

Confidential manuscript submitted to International Journal of Food Microbiology

85 It was aimed at achieving conceptual breakthroughs through an imaginative combination of 86 recent developments in microbiology, epidemiology, mathematical modeling and expert 87 knowledge; using these to propose new approaches to analyze and control food safety issues in the future. Such tools should enable risk assessors to pro-actively address imminent 88 89 problems before they cause harm. This paper provides an overview of the major themes 90 identified during the conference. The paper starts with a discussion of food safety from the risk management perspective. A systems approach to identify and structure future 91 92 developments that may help to develop a pro-active approach to food safety is introduced, 93 followed by a more detailed discussion of relevant developments with special attention for the 94 interaction between micro-organisms and their environment, and for microbial evolution. 95 Despite the need for pro-active approaches, surveillance and monitoring are discussed as important cornerstones of food safety policy and new developments in the available 96 methodology are discussed. Finally, needs related to communication between all actors in the 97 98 food chain are outlined.

99 2. Risk management

100 Food safety demands

101 Management of microbial food safety is a balancing act involving disparate factors. A high 102 level of safety can be achieved by rigorously heat-sterilizing the food, thereby destroying taste 103 and nutritious value. Irradiation would be another method for virtually absolute control of 104 microbial risks, but in addition to being expensive, it is not acceptable to the public at large in 105 many countries. Furthermore, some bacterial and fungal toxins are not inactivated by 106 currently used irradiation doses. The consumer demands fresh, tasty, healthy and wholesome 107 food products. Nevertheless, safety is in this framework considered an absolute requirement; 108 placing unsafe food on the market is not an option in the consumer's mind. Food laws

Confidential manuscript submitted to International Journal of Food Microbiology

109 everywhere are very clear on this point. For example, the EU General Food Law 110 (Anonymous, 2002) states that: "a high level of protection of human life and health should be 111 assured in the pursuit of Community policies". Still, placing chicken contaminated with 112 Salmonella or Campylobacter on the market is tolerated, because the consumer can 113 circumvent this risk by cooking the meat properly and taking adequate precautions against 114 cross-contamination, illustrating that responsibility for food safety is distributed over the entire chain. Nevertheless, there is an increasing pressure on producers to reduce 115 contamination levels of fresh meat as far as possible and economically feasible. It has been 116 117 demonstrated that it is very difficult to modify consumer behavior by education campaigns 118 (Nauta et al., 2008).

119 Microbial food safety differs fundamentally from chemical food safety. While chemical

120 residues and additives typically enter the food chain at more or less predictable steps,

microbes can enter at any step. They grow and die and interact with the food in ways that are 121 at best empirically described, but less understood in detail. The effects are also of a different 122 nature. Chemical contaminants, such as dioxins, can accumulate in the human body over the 123 years and still exert influence long after ingestion. Microbial pathogens can in some cases be 124 125 dormant for a certain time, but usually cause disease in a matter of days or weeks. The public 126 perception of microbial and chemical risks is also different. Residues of pesticides cause 127 public outcries if they exceed the norms, but usually will not have any noticeable detrimental 128 effect, while foodborne microbial and viral diseases are generally more accepted as facts of 129 life, as long as death or permanent harm do not occur (Hansen et al., 2003)

130 One of the challenges for managers of microbial food safety risks is to put in place effective

131 controls, without unnecessarily increasing costs or reducing taste and nutritional value.

132 Microbial hazards can be introduced at any step in the production chain and the most effective

133 opportunity for controlling those hazards can very well be a different step. Microbial risk

Confidential manuscript submitted to International Journal of Food Microbiology

134 management therefore requires a thorough understanding of the entire food production chain. 135 Monitoring the presence of pathogens in the end product usually is an inefficient approach to 136 hazard control, because it is impossible to test sufficient samples to obtain the necessary 137 degree of statistical power to detect contaminants at levels that may create unacceptable 138 health risks. Furthermore, by the time the potential presence of pathogens has been confirmed, 139 the optimal moment to take measures may have passed. Therefore, a pro-active approach is required, starting with the producer ensuring a safe product and process design, and predicting 140 141 where problems might arise, rather than detecting them after they have occurred.

142 At present HACCP (Hazard Analysis Critical Control Point) programs and GMP (good 143 manufacturing practice) are mainly used to manage microbial hazards in foods. While these 144 systems have proven very effective for the control of food safety (Van Der Spiegel et al., 2004, Arvanitoyannis and Traikou, 2005), it must be realized that they are designed on the 145 basis of known hazards, and do not necessarily take potential future developments in 146 consideration. For innovations, new validations and verifications are necessary. Furthermore, 147 148 it should be realized that the implications of microbial adaptability are not sufficiently taken 149 into account (McMeekin and Ross, 2002, McMeekin et al., 2006). Although documentation is 150 a very important aspect of HACCP and GMP procedures, the real confidence in control 151 comes for the validity of the effectiveness of the written guidelines and the adherence to them. 152 Producers and handlers of foodstuffs are more likely to adhere to the prescribed HACCP 153 procedures if they recognize these as useful and implementable (Taylor and Taylor, 2004).

154 Science-based risk-management

Within the food safety discipline the terms "risk manager" and "risk management" are not unequivocal and are used to indicate several functions and persons. As a consequence there are several persons with responsibility as risk managers, each at a different step of the food chain. Formally "the" risk manager is the government's minister of public health and/or

Confidential manuscript submitted to International Journal of Food Microbiology

159 his/her colleague of agriculture, as they set the standards to which food producers must 160 adhere. Furthermore, within government supervisory agencies the persons in charge of the 161 enforcement branch are often called risk managers. Often, however, the person responsible for 162 compliance to procedures within a food manufacturing company is called a "risk manager" as 163 well, as this term in fact very well describes his/her day-to-day activities. Although these functions are less clearly defined in smaller operations and in primary production, the role 164 should be fulfilled in any food operation. Risk assessors, on the other hand, have a different 165 function. They provide the risk manager with science-based advice on the magnitude of risks 166 and cost-effective ways to reduce these. This advice enables the different risk managers to 167 take decisions on measures to control risks, by setting standards, by implementing in-plant 168 control measures and/or enforce existing regulations. 169

170 A challenge to food microbiologists in their role of risk assessors is to translate complex 171 scientific problems in such a way as to help a risk manager to make a simple yes/no decision. The risk manager wants to know what standards to set, when to interfere in the production 172 process, prevent a batch from reaching the market or take another measure. Under the WTO 173 174 Agreements, and in particular the Agreement on the Application of Sanitary and 175 Phytosanitary Measures, considerations of food safety and animal and plant health are the 176 only legitimate reason for trade restrictions. The type of decisions in these cases is also of the 177 yes/no kind. The underlying science will have to provide the decision maker with a solid 178 rationale that will stand up in the international courts. This implies that data on the 179 microbiological status of the foods concerned must be communicated in terms of public 180 health risk with a limited margin of error, requiring a predictive power from food 181 microbiology. As zero-risk is unattainable, this approach also implies that the risk manager 182 defines a level of acceptable (tolerable) risk. In international trade, this is called the 183 Appropriate Level of Protection, equivalent to the currently realized risk level under the food

Confidential manuscript submitted to International Journal of Food Microbiology

184 safety system of the importing country. Food safety managers may wish to achieve a higher
185 level of protection in future by stating additional public health targets (Anonymous 2006,
186 Anonymous 2007b).

187 Food safety risk-management in the EU

Public concern about food safety increased sharply as a result of the food scandals in the last 188 decade of the twentieth century and confidence decreased in parallel. To counter these 189 190 sentiments national governments and the EU established food laws and regulations that 191 strictly separate risk management from risk assessment. The idea behind this was to create 192 transparency by having the risk assessor provide advice to the risk manager completely in the 193 open. The risk assessor operates independently, based on the best available science and free 194 from influence by politics, industry or any other stakeholder. The risk assessor should give 195 objective advice, based on science while taking account of other considerations that the risk 196 manager has indicated. The risk manager can then incorporate other factors such as public concern or political preferences into the decision making process. 197

198 The exact procedures for risk assessment differ considerably in the different member states. 199 The EU itself has mandated the European Food Safety Authority (EFSA) to carry out risk 200 assessments, either at the request of the Commission, a member state or the European 201 Parliament, or on its own initiative. The EFSA in turn has handed this task to 10 scientific 202 panels and the Scientific Committee, which it finances and supports scientifically and 203 otherwise. These panels write opinions which are passed on unchanged to the EU 204 Commission and published on the EFSA website. When strong public interest is expected a 205 press release is issued as well. The panels are comprised of scientists who operate 206 independently from risk management, and almost all come from EU-member states, though 207 this is not a formal requirement. The scientists are chosen in an EU-wide application

Confidential manuscript submitted to International Journal of Food Microbiology

- 208 procedure on the basis of their expertise and prominence in their scientific fields, and are
- appointed for a period of three years at a time.

210 The advice of EFSA's panels is presented as a scientific opinion to assist in the formulation of

- 211 risk management measures by the European Commission. Measures may be decided upon by
- the members states in a consensus procedure, allowing political, economic and other
- 213 considerations to influence the decision making process.

214 Interaction between scientists and regulators

215 Researchers, in their role as risk assessors, need to provide the regulators, in their role as risk

- 216 managers, with advice that can be implemented in a practical manner. Regulators may or may
- 217 not have a scientific background and thus results of risk assessments have to be

218 communicated in a way that can be understood by fully informed laypersons. The process of

- 219 risk analysis, as defined by Codex Alimentarius, consists of risk assessment, risk management
- and risk communication. The very first step in the process, hazard identification, is a
- 221 component of risk assessment. The hazard identification can come from any source. In
- 222 practice it is often the risk manager, who "identifies" the risk by asking for a risk assessment.
- For a good risk assessment, the problem at stake needs to be well understood by the assessor.
- 224 Therefore, correct phrasing of the request is essential for a successful risk assessment
- 225 procedure. The regulator needs to articulate his question so that it will not be misunderstood
- by the scientist. In short, they need to "speak each others language". This seems a trivial
- 227 point, but experience proves that it is not.

The roles of the risk assessor and the risk manager need not only to be formally separated, but also with respect to substance. A risk manager has to consider more issues than science only, such as stakeholder interests, public concerns and political pressure. The risk assessor needs

Confidential manuscript submitted to International Journal of Food Microbiology

- 231 to present a science-based advice, but may anticipate the risk manager to take other than
- 232 scientific factors into account when selecting the risk management options.

233 **3.** A systems approach to analyzing future challenges to food safety

234 In addressing the present or future state of food safety one should investigate effects both on large scales of space and time as well as on small scales (Table 1). Certain aspects slowly 235 change at a global scale, like climate change while others such as point mutations or the 236 237 acquisition of a plasmid by a micro-organism, occur on a molecular scale and on very short 238 time scales. In addition, certain aspects occur over longer time scales, but on a molecular 239 level, like subsequent adaptation of micro-organisms, or on a small time scale but on a larger 240 spatial scale, like the spread of a virulent micro-organism due to the large traffic of people or 241 goods. Furthermore these changes occur in micro-organisms and humans, in habitats and in 242 the environment. To accurately describe and predict processes in all these different organisms 243 and locations, on these very different spatial scales and time scales is virtually impossible, the more because all these aspects interact. The risk framework of the UK Foresight project on 244 245 infectious diseases (Tait et al., 2006) offers a useful starting point for the development of 246 scenario analysis in relation to food safety. The project has developed the following definitions, which are cited literally here: 247

Disease sources/emerging hazards: phenomena or biological events that: give rise to
 potential new diseases; enable existing diseases to become more harmful; enable
 existing diseases to infect new hosts; or enable existing diseases to spread to new
 areas.

Pathways: mechanisms or routes by which a disease organism can transfer from one
 host to another, within or between species.

Confidential manuscript submitted to International Journal of Food Microbiology

- Drivers: social, economic or physical factors that affect disease outcomes by changing
 the behavior of disease sources or pathways.
- Outcomes: diseases of plants and animals at the individual, community and ecosystem
 or farming system level, and diseases of humans at individual and societal levels.
- 258 A basic risk framework (Figure 1) shows the links between these different factors.

259 The PERIAPT project on emerging risks (Noteborn *et al.*, 2005) identified 8 major categories

260 of drivers, based on expert surveys. Later, we will present a tabular representation of these

261 drivers of change in the food system, and a qualitative analysis how they affect sources,

262 pathways and outcomes. To better explore these complex interrelationships among different

263 drivers, mathematical models may be helpful.

264 Mathematical models are a representation of the essential aspects of an existing system (or a system to be constructed), presenting knowledge of that system in a usable form. A 265 266 mathematical model usually describes a system by a set of numerical variables and a set of 267 equations that establish relationships between the variables. The variables represent some properties of the system, obtained by measurements or by expert opinion. The actual model is 268 269 the set of functions that describe the relations between the different variables. The purpose of 270 modeling is to increase our understanding of the world. The usefulness of a model rests not 271 only on its fit to empirical observations, but also on its ability to extrapolate to situations or 272 data beyond those originally described in the model (from Wikipedia, March 21, 2008).

Some examples of models that are currently used to analyze and to support decision makingon food safety:

Microbial risk assessment (MRA) models (hazard identification, exposure assessment,
 hazard characterization (including dose-response), risk characterization).

277 Used to understand the relationships of pathogen occurrence (both prevalence and

Confidential manuscript submitted to International Journal of Food Microbiology

278		concentration) in different steps of the food chain, to predict health risks associated
279		with pathogens in food, and the expected public health effects of interventions and the
280		setting of risk-based standards for food production.
281	•	Predictive microbiology (growth/death/survival).
282		Used to understand the growth or death of micro-organisms in relation to their
283		implicit properties and interactions, and the intrinsic properties of the food and the
284		extrinsic factors of the (processing) environment. It is an important component of
285		MRA models, and also used for prediction of shelf-life and intrinsic safety of foods.
286	•	Dynamic infectious disease models.
287		Used to understand the spread of diseases in human or animal populations, depending
288		on contact patterns and mode of spread of the pathogens, in relation to the
289		development of protective immunity.
290	•	Risk factor models (analytical epidemiology).
291		Used to relate the observed occurrence of responses (e.g. illness) to the occurrence of
292		potential predictive factors.
293	•	Attribution models.
294		Used to estimate the contribution of putative sources to the observed occurrence of
295		responses (e.g. illness).
296	•	Multi-criteria analysis models.
297		Used to support decision makers in making evaluations of different options, based on
298		a combination of variables of a different nature (health, economic, societal,) with
299		value-based weights.
300	All m	odels are based on a set of (simplifying) assumptions and have specific data needs. They

301 also differ in their form (linear or non-linear, deterministic vs. stochastic, static vs. dynamic)

Confidential manuscript submitted to International Journal of Food Microbiology

which impacts on their use for specific purposes. It is noted that all models described above
work at relatively low levels of aggregation, looking at specific (parts of) food chains in
relation to public health effects. The challenge is to develop models that are able to capture
the impact of different drivers on foodborne risks at higher levels of aggregation.

? The overall aim of this coupling of various aggregation levels is to better understand how 306 307 drivers affect the evolution of foodborne illness and to determine the most important drivers. 308 This can help to react with a timely and adequate response, and support a pro-active approach, 309 targeted at future events. Generally one needs to decrease complexity in the more detailed 310 parts and move upwards to more global aspects, but also vice-versa, since after identifying 311 certain important aspects on a global scale one might need to change focus to the more 312 detailed level. For this, intensified linking to models in other domains is necessary, like 313 biosphere models, geospatial modeling, catastrophe modeling, climate models, remote 314 sensing, network science, statistical physics, and data mining.

For these models, a huge amount of data is needed such as information of food production sites and global product flows (volume, origin) for high risk products (e.g. fresh meat, fresh produce, shellfish), risk maps, global atlases of food consumption and production, food categories with different levels of risk. Connecting them in dynamic systems can help to identify rates of change. However, it is necessary to begin with low granularity, with more detail based on sensitivity analysis.

In current risk evaluations considerable uncertainties already prevail, so, when modeling future risks even larger uncertainties will exist. However, important insights can still be gained using the available information, making the best informed decisions at a given point in time, that later can be detailed if more and better information becomes available. Scenario analysis will be particularly important to better understand the impact of different factors, their interrelatedness and their uncertainties. It is a process of analyzing possible future events

Confidential manuscript submitted to International Journal of Food Microbiology

by considering alternative possible future developments and outcomes (scenarios), and their
likelihood. These insights can then be used to develop a range of contingency plans to address
the most likely or most serious scenarios. The analysis is designed to allow improved
decision-making by providing more complete consideration of outcomes and their
implications. Typically, scenario analysis starts with the identification of possible important
drivers of change, and subsequently assigning a preliminary ranking of their importance.

333 4 Trends and future developments

334 In this section, we present an attempt to collect and structure available information on the current and future aspects of microbial food safety. Table 2 gives an overview of factors 335 identified so far, according to the model presented by Tait et al., (2006). We interpret the 336 337 sources category as referring to the pathogens; the pathways category is split into the three 338 major stages of the farm-to-fork pathway (farm, processing and consumption), and outcomes 339 are defined at the public health level. Note that there are complex interrelationships between 340 different drivers, sources, pathways and outcomes that are difficult to visualize in a two-341 dimensional table, and hence the information must be interpreted with care. Nevertheless, 342 important insights can be gleaned from this analysis, which provides a general background 343 against which specific situations can be analyzed. Currently, it is only possible to discuss the 344 impact of drivers of change on sources and outcomes in general, qualitative terms and so an 345 attempt has been made to indicate the anticipated direction of change for specific sources and 346 overall for drivers. It is clear from Table 2 that many drivers are expected to result in an 347 increased risk to food safety, although there are also favorable exceptions, such as the lesser 348 consumption of meat due to higher food prices. Controlling such threats is a challenge to 349 governments as the only drivers assumed to result in reduced threats relate to government and 350 policies. A second important challenge is to food industries that can modulate the effects of

Confidential manuscript submitted to International Journal of Food Microbiology

- 351 many drivers in such a way that a neutral or even positive effect on food safety can be
- 352 expected (see drivers under science, technology and industry).

353 Further elaboration of the crude framework sketched in this chapter requires considerable

- inputs. It is suggested that to further develop the framework, historical examples be analyzed.
- 355 This will provide more detailed insight regarding relevant drivers and sources, and their
- interaction as well as data to validate the approach. Such examples include the BSE epidemic,
- different meatborne zoonoses (e.g. VTEC O157), shellfish poisoning and more recent
- 358 outbreaks in fresh produce.

359 In order to improve future responsiveness, signals about changes or breakdowns in the food

360 safety system must be received and processed in time. This implies a pro-active approach.

361 Process or production failures, including fraud and terrorist action need specific attention.

362 Risk mapping (on a global scale) can be a helpful tool. This includes observation and

363 systematically analyzing consumption patterns, processing / production changes, knowledge

364 of international production chains (trade), and data & knowledge sharing. Where necessary,

365 available data can be supplemented with expert opinions (global and multidisciplinary).

366 Trends in food processing

The trend for mildly preserved foods comes with a range of approaches being investigated, 367 mostly using minimal heating, natural preservatives and non-thermal treatment as 368 369 technologies and often combined preservation/hurdle technology as the principle in designing 370 the overall treatment. Such processes need to be well controlled through adequate product and 371 process design and proper implementation and monitoring through HACCP. This places a 372 responsibility on industry, including small enterprises. There are more weak links, and overall 373 the processes are less robust and more accident prone. As the scale of operation of food 374 businesses continues to increase, errors may have a bigger impact.

Confidential manuscript submitted to International Journal of Food Microbiology

375 Consumers need to be aware of the criticality of the formulations and of the need to treat 376 manufactured foods either as perishable products needing proper refrigeration or requiring 377 specific conditions of preparation (for instance non-Ready-To-Eat products that need to be 378 cooked properly before consumption even though they may appear to be cooked). The 379 concern is that many consumers do not habitually read labels and are unaware of shelf-lives 380 and preparation requirements.

Can these new product types offer niches for concurrent, old, emerging or new microbial 381 382 hazards? Classical examples are refrigeration and the niche created for Yersinia and Listeria, 383 and for sporeformers by non-thermal treatments at the pasteurization level. As in many of 384 these foods spoilage organisms have been removed or suppressed, there is increased 385 opportunity for the growth of pathogens that recontaminate treated products – in the absence 386 of the "normal" spoilage signal. Noroviruses show prolonged survival during cold storage and 387 even freezing. The probability and extent of survival and of post-process contamination, rather than pathogen growth opportunity, may then determine the level of consumer risk. 388 389 While there would be a benefit from (and indeed a need for) for irradiation technology for certain applications (e.g. it is "safe", "invisible", no microbial issues of resistance, or of 390 391 recontamination when done in-pack), consumer concerns towards the technology itself and 392 about misuse to make spoiled food marketable, prohibit its wider use in practice. This implies 393 considerable communication challenges, should the technology prove to be the treatment most 394 relevant for certain applications.

While reducing packaging is a laudable initiative where the packaging is for "cosmetic" or bulk-transport purposes, industry and consumers should be aware of those situations where the packaging has a preservative function by minimizing growth and/or recontamination of microorganisms.

Confidential manuscript submitted to International Journal of Food Microbiology

399 Consumer behavior

400 Several changes in food composition related to consumer health are foreseen. These may also 401 have an impact on bacterial growth. For instance, components like salt and sugar are often 402 used to inhibit the growth of organisms, both by their water activity lowering effect, and 403 additionally solute specific effects. Their concentration cannot be safely reduced for health or 404 other non-safety reasons without adapting the product design. On the other hand, fat can be 405 seen as a vehicle for better stomach survival of pathogens, so less fat might reduce risks. 406 Lower fat may also increase the water activity of a product by having more diluted solutes in 407 the aqueous phase (Senhaji, 1977).

408 Considering the present increased level of general health of the population and better medical 409 treatments, proportionally more elderly will be present in society. These elderly often are 410 more vulnerable to foodborne diseases partly, as a result of a weakened immune system 411 increasing the risks of complications and even death. Other defence systems, such as stomach 412 acid may be impaired (achlorhydria), augmented by medication.

413 Due to changes in eating habits, certain risks might change in magnitude. For example an 414 increased consumption of fish for health benefits may result in increased microbial risks. The 415 increasing trend for fresh, pre-packaged produce or other foods that are consumed without 416 additional heating by consumers also increases consumer risk. Exotic and ethnic foods are 417 now trendy in the market, but do we understand the underlying preservation system? When 418 we change these products (adapted for new markets; altered ingredients), are we clear on how 419 this may affect safety? More and more animal species are used for food production and there 420 is little knowledge about zoonotic risks of such foods (e.g. reptile meat, Magnino et al., 421 2009). New culinary techniques, such as molecular gastronomy involve more and more 422 technical creativity and exotic ingredients to improve quality and consumer acceptance, which may result in unexpected risks. In the case of small restaurants, catering establishments and 423

Confidential manuscript submitted to International Journal of Food Microbiology

street vendors the scale of production is small and often, knowledge and sufficient technology
may be lacking, resulting in avoidable errors which may lead to serious consequences for
consumers.

427 Improving animal welfare (e.g. by increased outdoor access) and organic food production may lead to the re-introduction of pathogens with wildlife reservoirs such as Trichinella 428 429 spiralis and Toxoplasma gondii and increase the prevalence of other hazards such as 430 *Campylobacter* spp. but may reduce the prevalence of others such as *Salmonella* spp. 431 (Gebreyes et al., 2008). Reduced usage of antimicrobial agents may have a positive impact on 432 resistance development (Van der Giessen et al., 2007; Hoogenboom et al., 2008). Trends 433 towards continuously increasing herd size in intensive bio-husbandry may also lead to 434 increased public health risks by increased numbers of contacts between food animals, 435 including purchasing of animals from more suppliers. The potential for infections to spread 436 increases with herd size, and if such farms are concentrated in particular regions, there is also 437 an increased risk of spread between farms. On the other hand, establishment of larger, newly designed farms may improve conditions for biosecurity and farm management, potentially 438 439 reducing zoonotic risks (Kornalijnslijper et al., 2008).

440 Price

441 It is well known that cost is a very important consideration for the consumer when selecting 442 foods, and that profits on food products are generally quite low. Both aspects make it difficult 443 to be critical towards food safety and furthermore might occasionally result in fraud. Also, 444 costs and conservatism may lead to resistance against implementing reasonable interventions 445 or to interpret existing regulations liberally (e.g. use of approval of sick animals for 446 consumption). Such non-compliance may increase consumer risk. The Law Enforcement 447 Department of the Netherlands Food and Consumer Safety Authority experienced during the 448 2008-9 economic downturn that food producers and retailers more frequently violate

Confidential manuscript submitted to International Journal of Food Microbiology

regulations relating to cleaning and maintenance due to cost cutting (J. van der Kooij,
personal communication). Increasing food prices are likely to compromise food security (in
terms of food availability) on a global scale. For industrialized countries, food security is less
at risk, but consumers may choose less costly alternatives. This may lead to less consumption
of animal proteins (also driven by animal welfare and environmental considerations), leading
to other health-related issues. Higher food prices may cause consumers to use food more
frequently past its shelf life, and may increase recycling of food.

456 Global aspects

Due to the increase of international travel, organisms can be spread easily and quickly over 457 458 the globe, and people come in contact with organisms and specific strains to which they have 459 not been exposed earlier, increasing the risk of illness. Global trade will result in a longer 460 transit distances and durations in the food chain, possibly increasing risk. Furthermore, 461 complex food chains with stakeholders in many different countries will make the management of safety more difficult, especially at the initial stages of the food chain, the primary 462 production as it consists of many small farms and is increasingly global in nature. On the 463 464 other hand more powerful stakeholders (trade companies, supermarkets) will have the 465 intention to influence these complex chains in order to guarantee food safety. Several large 466 retailers that operate internationally have organized the GlobalG.A.P. quality control system 467 (www.globalgap.org) that aims to supervise the primary production process of all agricultural 468 products. Sourcing food from various climate areas means more variation in hazards. Border 469 controls are effective in regard to the control of only a small proportion of imported foods and 470 are less effective than hygiene controls imposed in the country of origin. Furthermore, global 471 food chains may be more vulnerable to terrorist attacks.

472 More and more harmonization will occur in international regulation of food safety by the473 activities of e.g. the Codex Alimentarius Commission and the European Union. As a positive

Confidential manuscript submitted to International Journal of Food Microbiology

effect, fairness in trade will increase but on the other hand the same level of contamination infood may result in very different health risks in different parts of the world, due to differences

- 476 in e.g. demography, immune status, food preparation and consumption habits, and relevance
- 477 of various routes of infection.

478 *Climate change*

479 Climate change is considered to be one of the greatest current challenges to mankind,

480 affecting all sectors of society, including nutrition, food security and food safety

481 (Anonymous, 2008b). Due to climate change various risks may change, as a result of changed

482 ecological conditions on various places on the earth. Changing ecology is expected to affect483 the distribution of plant and animal diseases. Water shortages may lead to limited quantities

484 or quality problems with irrigation water, process water or ingredient water. This may lead to

485 shifts in production areas and cultured crops, as well as an increased use of agrochemicals.

486 This trend may be increased by competition for land-use, e.g. for biofuels or for settlements.

487 Flooding may lead to increased contamination of crops in the field, or increased exposure of

488 food animals to zoonotic agents. Control of cold chains may be impeded by rising ambient

489 temperatures. Humidity may increase production of mycotoxins, whereas certain foodborne

change affect food safety are highly complex and interrelated with many other societal

490 pathogens may thrive better under warm conditions. The mechanisms by which climate

492 factors, The outcome of these changes strongly depends on the adequateness of societal

493 responses, both of a technical and a political nature.

494 Science

491

495 More and more public health risks, physiological and ecological traits of foodborne

496 pathogens, routes of contamination, effects of interventions, will be investigated and

497 quantified, making it possible to better balance risks, and evaluate optimal interventions to

Confidential manuscript submitted to International Journal of Food Microbiology

498 control risks to an appropriate level. That is, science will have a greater role in setting criteria499 both nationally and in international trade agreements.

The genomics revolution will facilitate easier and faster detection and identification methods, and can in particular lead to a better mechanistic understanding of the behavior of microorganisms, both their physiology as well as their ecology. Furthermore new pathogens can be uncovered, for example better detection of injured and thus less easily culturable organisms is possible and more advanced methods to investigate cases and outbreaks become available.

505 Antimicrobial resistance

Usage of antimicrobial agents, both in the agricultural sector and in human health care 506 settings, contributes to the emergence of resistant microbes. While resistance in the 507 508 agricultural sector might be considered an economic problem with limited other 509 consequences, it is developing into a cause of growing concern for human health care (see Newell et al., in an accompanying paper in this Special Issue). The overall use of 510 511 antimicrobial agents in food animals is high and in certain countries largely exceeds the 512 human use. The ban on usage of antimicrobials as growth promotors has, in some countries, 513 barely had an influence, as it has been replaced by increased use for therapeutic purposes 514 (Mevius and Van Pelt, 2006). Micro-organisms have an immense diversity and can easily 515 transfer genetic information, making the emergence of new hazards, and adaptation to 516 previously effective intervention methods possible.

At present, it is not clear what proportion of resistance encountered in human pathogens originates from selection in and transfer from animal reservoirs. It is not known if measures to reduce usage in the agricultural setting will lead to a rapid reduction of resistance, as this can also contribute in other ways to overall fitness of the micro-organism. The benefits to human health care of such measures are not quantified, as resistance may also develop due to other

Confidential manuscript submitted to International Journal of Food Microbiology

- factors. In spite of these uncertainties, the development of antimicrobial resistance in
 agricultural settings is a cause of growing concern to public health and prudent use is
 advocated.
- 525 **5. Epidemiology and surveillance**

526 Monitoring of contamination in the food chain, combined with surveillance of human illness and epidemiological investigations of outbreaks and sporadic cases continue to be important 527 sources of information to evaluate the success of current food safety management systems and 528 to identify new hazards. Surveillance is defined as "the ongoing and systematic collection, 529 analysis, and interpretation of data about a disease or health condition; used in planning, 530 implementing, and evaluating public health programs" (Anonymous, 2000a). Surveillance can 531 532 be aimed at outbreaks or sporadic cases of foodborne disease, and continues to be a 533 cornerstone of food safety management (see the accompanying paper by Tauxe *et al.*, in this 534 issue).

Outbreak surveillance primarily aims to stop the outbreak by identifying incriminated 535 536 products and taking them from the market. Furthermore, investigations may aim to prosecute 537 those responsible, or to learn from outbreaks so as to avoid future outbreaks by identifying unsafe practices that had led to the outbreak. Outbreak investigations have and will continue 538 539 to be an important instrument for identifying new pathogens (e.g. Cyclospora cayatenensis, 540 Herwaldt, 2000), new vehicles for known pathogens (e.g. Salmonella Tennessee in peanut 541 butter, Anonymous, 2007c), new disease syndromes associated with known pathogens (e.g. 542 febrile gastroenteritis associated with *Listeria monocytogenes*, Dalton et al., 1997), and the re-543 emergence of problems that were thought to be under control (e.g. botulinum toxins in canned 544 foods, Ginsberg et al., 2007). They are important sources of data for establishing the

Confidential manuscript submitted to International Journal of Food Microbiology

545	economic impact of foodborne illness on populations and may provide dose-response

546 information for microbial risk assessment (Teunis *et al.*, 2008).

547

548 reported at an aggregated level annually (Anonymous, 2007a) or over a number of years

Many countries have surveillance systems for outbreaks of foodborne illness and data are

549 (Wang et al., 2007, Cretikos et al., 2008). New tools are becoming available to detect

550 international outbreaks for foodborne viruses (Verhoef *et al.*, 2009). Supranational agencies

such as EFSA and ECDC in Europe present regular reports on foodborne outbreaks in a larger

region or globally (Anonymous, 2007a). Outbreak summary reports provide important

insights in current and emerging food safety problems but it is essential that such summaries

are based on systematic surveillance activities. Reports in the peer-reviewed literature may

555 suffer from publication bias and overestimate the impacts of milk/milk products,

556 miscellaneous foods (e.g. sandwiches) and desserts while underestimating those of poultry,

557 fish and shellfish, red meat/meat products and eggs/egg products (O'Brien *et al.*, 2006).

558 Molecular tools identifying causative agents in environmental and clinical samples, and

559 molecular typing techniques identifying nucleotide sequences of single genes (i.e. *fla*-typing),

560 techniques identifying sets of genetic elements (MLST, MLVA) and various restriction

techniques (i.e. PFGE) have proven to be very useful aids in the epidemiology of foodborne

562 illness. Developments in genotyping of pathogens and informatics have enabled the

563 recognition of diffuse or multinational outbreaks which were previously unnoted (Gerner-

564 Smidt *et al.*, 2006, Kirk *et al.*, 2004, Kroneman *et al.*, 2008).

Estimating the incidence of sporadic cases of foodborne illness is more complex. Most existing surveillance systems are based on either notifiable disease reporting or laboratory surveillance. Both systems are passive in nature, and record only a minor proportion of all cases in the population. To estimate the true incidence of diseases that can be transmitted by food, active surveillance is necessary and more accurate estimates are needed for under-

Confidential manuscript submitted to International Journal of Food Microbiology

570 reported illness. The UK (Wheeler et al., 1999, Tompkins et al., 1999) and the Netherlands 571 (De Wit et al., 2001a, De Wit et al., 2001b, De Wit et al., 2001c) have carried out population-572 based prospective studies of infectious gastro-enteritis, combined with laboratory diagnostics 573 to assess the proportion of cases due to specific pathogens. Currently, the UK has launched 574 the second IID study (http://www.iid2.org.uk). Even in these large-scale projects, in a large 575 proportion of cases (60%) it was not possible to identify a causal pathogen. However, it appears to be possible to reduce this diagnostic gap by the application of molecular methods 576 577 (Amar et al., 2007). Population-based studies are expensive and time consuming, and several 578 countries have attempted to develop less costly alternatives. These include FoodNet in the 579 USA (Jones et al., 2007), OzFoodnet in Australia (Kirk et al., 2008), and the International 580 Collaboration on Enteric Disease Burden of Illness Studies (Flint et al., 2005, Roy et al., 2006, Thomas et al., 2006). As laboratory-based surveillance only detects a fraction of all 581 582 illness occurring in the population, modeling approaches have been used to reconstruct the 583 surveillance pyramid (Michel et al., 2000, Voetsch et al., 2004, Majowicz et al., 2005). Serosurveillance is now being explored as a new tool to provide internationally comparable 584 585 estimates of the exposure of populations to foodborne pathogens (Simonsen et al., 2008). 586 Although most surveillance activities are focused on gastro-intestinal illness, other symptoms 587 are also commonly associated with foodborne illness. These may be more serious or of longer 588 duration than GI illness. Furthermore, most pathogens that can be transmitted by food may 589 also be transmitted by other pathways such as water, direct human and animal contact. 590 Therefore, there is a need for source attribution to quantify the proportion of all cases that is 591 foodborne, and the food vehicles that are most frequently associated with illness (Batz et al., 592 2005). Molecular typing has successfully been used for source attribution of salmonellosis 593 (Van Pelt et al., 1999, Hald et al., 2004) and more recently for campylobacteriosis (Wilson et 594 al., 2008). Other methods being explored include case-control studies, outbreak studies, risk

Confidential manuscript submitted to International Journal of Food Microbiology

595 assessment modeling, natural or deliberate intervention studies at population level and expert 596 elicitation. Each method is subject to specific biases, and may attribute illness to different 597 points in the food chain. Therefore, interpreting the results from attribution studies should be 598 done with care (Pires et al., 2009). 599 The World Health Organization has recently launched a new initiative to estimate the burden 600 of foodborne illness on a global scale (Stein et al., 2007). This initiative is advised by experts 601 of the Foodborne Epidemiology Reference group (FERG), which assembles and appraises 602 global evidence on foodborne disease epidemiology. This action is considered necessary in 603 view of globalization, and to contribute towards meeting the Millennium Development Goals¹. 604 Results will be a basis for action at the global scale. Virtually no data on morbidity and 605 mortality exist in large areas of the world, and even more data gaps are expected for attribution. Therefore, systematic reviews will be carried out and extrapolation will be 606 necessary. As an example approach, the estimates of death from diarrhea for children under 5 607 608 from the Childhood Epidemiology Reference group (CHERG) will be used (Boschi-Pinto et al., 2008); complemented with other methods, including expert opinion. 609 In addition to surveillance of human illness, systematic food chain surveillance is necessary to 610 611 inform food safety decision making. Recent EU-wide baseline studies on the prevalence of 612 zoonotic pathogens have illustrated the benefits of such standardized sampling and analytical 613 approaches. For example, in the baseline survey on the prevalence of *Salmonella* in slaughter 614 pigs, which took place between October 2006 and September 2007, it was demonstrated that 615 approximately one out of every ten slaughter pigs in the European Union was infected with Salmonella in the lymph nodes, while one out of twelve pig carcasses was contaminated with 616 617 Salmonella. The survey also indicated large differences between Member States (Anonymous,

618 2008c). These data will be the basis for risk assessment and cost-benefit analysis of

¹ http://www.un.org/millenniumgoals/

Confidential manuscript submitted to International Journal of Food Microbiology

619 *Salmonella* control in the slaughter pig chain to support decision making by European risk620 managers.

The previous sections have described a highly complex set of interrelated factors affecting future trends in food safety. Predicting the impact of these factors is highly complex and surrounded by uncertainties. Hence, to be able to respond timely and appropriately, active, real-time surveillance in both the human and food system and communication to professionals responsible for infection control is of utmost importance.

626 6. Methodology

627 Molecular Methods for complex food analysis

One of the key challenges in food microbiology that has always been around and can now be 628 629 addressed is to assess what molecular mechanistic processes underlie the observed physiological behavior of pathogens in food (see e.g. McMeekin et al., 2007). Much of this 630 631 work relies on a proper identification of (a) the microorganisms in the food at hand and (b) the food components that are relevant in determining the microbial stability of such foods. 632 633 The latter range from small molecules (flavor-like molecules, food preservatives and other 634 organic molecular) to the macro-ingredients i.e. proteins (peptides), sugar (polymers) and fats. 635 In many foods the microbes that are to be analyzed for are non-uniformly dispersed 636 throughout the product. This is the case for many ready-to-eat products from the chilled food 637 chain and is equally so for liquid products such as sauces and soups in which particles, as putative sources of microorganisms, may be non-uniformly mixed. 638 639 Analysis of microorganisms in foods may be done with two objectives in mind. On the one 640 hand it may be a direct assessment related to production processes or inspection, on the other 641 hand it may be research-oriented in which physiological inferences are made from molecular 642 data. Rapid analysis techniques for use in industrial practice have to be easy to perform, low

Confidential manuscript submitted to International Journal of Food Microbiology

643 cost, optimally selective and must demonstrate reproducible sensitivity and specificity. Such 644 methods need to be validated and written down in standardized protocols, preferably being 645 able to provide quantitative data of use in risk assessment and in food safety management. Currently they are generally based on DNA detection systems, either specific for ribosomal 646 647 genes, or in the more advanced systems for specific sequences that occur along the entire 648 genome (see e.g. Wattiau et al., 2008; Scaria et al., 2008). Comparative genome sequencing is 649 certainly at hand nowadays. Thus, in the case of relevant, closely related bacterial isolates it is 650 increasingly easy to identify unique sequences (see e.g. the discussions in Earl et al., 2008 651 and Medini et al., 2008). Many of these may then be used to derive sequences amenable to 652 use in DNA chip and / or PCR based detection platforms to the benefit of the safety 653 assessment of food processing.

Although these methods are fast, highly specific and relatively sensitive, the application of 654 655 molecular-based techniques in the control of food safety seems to be limited as they suffer from some serious drawbacks. The development of a horizontal method is seriously hampered 656 by the fact that food products may contain interfering components. The development of 657 658 horizontal methods becomes even more difficult due to a constant introduction of new 659 matrices. While they are very sensitive, low copy numbers are difficult to detect when the 660 sample size is very small. Introduction of an enrichment step preceding DNA-detection is a 661 solution, but this makes results qualitative, rather than quantitative unless cumbersome MPN 662 techniques are used. While this may not be problematic for quality assurance purposes, 663 quantitative results may be necessary for risk assessment studies. Sample preparation needs 664 close attention. Preferably, such sampling needs to be rapid and as homogeneous as possible. 665 Innovative strategies focus on the use of magnetic beads coated with cell-recognizing 666 molecules, on physical methods such as floatation, and on lysis of whole food matrices 667 (Wagner and Dahl, 2008). The latter was described originally by Hein and co-workers who

Confidential manuscript submitted to International Journal of Food Microbiology

obtained enough bacteria from a complex set of food matrices in one-step approach taking 668 669 only a few hours to be able to recover DNA for further study (Rossmanith et al., 2007). It has 670 yet to be established that such a procedure will also be effective for determining the concentration of bacterial spores. A limitation of currently available molecular techniques is 671 672 also that they fail to discriminate between viable and inactivated organisms. Recent research 673 may provide future practical solutions to this as transcriptional activity around bacterial cell survival / death reveals molecular markers for cell viability (Kort et al., 2008). 674 675 Finally, assays based on detection of multiple virulence genes can still give ambiguous results 676 if there is a mixed culture to begin with. This may be for instance the case for the detection of the Shiga like toxin (stx) and Intimin (eaeA) genes from Escherichia coli in direct molecular 677 analyses on food samples (e.g. Monday et al. 2007). The outcome will be positive with one 678

679 strain having both or two strains, each having one of the virulence genes.

680 The issues discussed above corroborate the notion that it is necessary to properly validate 681 such newly developed techniques. How do they compare to standard reference culture 682 techniques described in ISO-protocols, and which controls must be used? Information is 683 available on the efficacy of protocols using spiked samples, but little information is available 684 on the efficacy of developed protocols in case of naturally contaminated samples. Without 685 multi-laboratory validation, protocols for molecular techniques can be used for in-house 686 purposes, but to be used as a standard method for the detection of pathogens, molecular 687 techniques have to be validated, using a multi-laboratory approach, according to the ISO-688 16140 protocol.

In analyzing the composition of foods it is also more and more possible to detail

690 comprehensively the full chemical spectrum of the compounds observed. To this end tools

691 such as liquid chromatography (LC) or gas chromatography (GC) coupled to mass

692 spectrometry (MS) are increasingly successfully used (reviewed in Hounsome et al., 2008). A

Confidential manuscript submitted to International Journal of Food Microbiology

693 full analysis provides valuable information on product quality as well as the environmental 694 parameters that can be most relevant to microbial survival (Beckmann et al., 2007). The 695 analysis may also be used to detect the presence of microbial spoilage (Ellis et al., 2007). 696 Pattern analysis to identify relevant compounds is the area where developments are rapid. 697 While the costs of detection at the DNA level are increasingly reducing and interpretation of 698 the data can now be automated to a large extent, this is not always as straightforward with the 699 measurements of small to medium sized (mostly) organic molecules (see e.g. review by 700 Hounsome et al., 2008).

701 In the research area, molecular techniques are widely used. They can be used for the identification of organisms, for behavioral studies or for studying genes involved in (the 702 703 regulation of) virulence and stress in response to different environmental conditions, while 704 they can also be used in evolutionary studies. This approach marks the second objective, i.e. 705 the use of genomics data to underpin physiological observations and to mechanistically 706 'explain' them. Here the analysis platform used need not be restricted to the cheaper methods, 707 focused on biomarkers only; in fact the approach should be wider in nature while costing 708 marginally more (though cost-effectiveness remains an important parameter). To pinpoint 709 which type of compound has the most effect on the microorganisms at hand it is useful to 710 analyze the genome-wide expression pattern and use the data obtained as a bioassay to 711 identify the physiologically most sensitive environmental cues. Translation of molecular data 712 into a biological meaning remains an essential subject for future studies. Application of 713 techniques from other disciplines like ecology and medicine will be very useful.

714 Mining molecular data for new threats; metabolic capability models

715 New methods to screen sequenced genomes with the aim of understanding the physiological

716 capability of microbes have led to the identification of major differences at the genome level

717 between common laboratory strains of e.g. E. coli K12, enterohemorragic E. coli and

Confidential manuscript submitted to International Journal of Food Microbiology

718 uropathogenic E. coli (Brzuszkiewicz et al., 2006; Fraser-Ligget, 2005; see also Perna et al., 719 2001). Extending the analysis with state of the art rapid sequencing technique to other 720 pathogens such as bacilli and certain streptococci has led to the realization of the so-called 'pan' and 'core' genome concepts (see e.g. Hiller et al., 2007, Ara et al., 2007 and the 721 722 discussion in Medini et al., 2008). In this classification the pan-genome is seen as being 723 composed of three elements: the core genome, a set of non-essential genes shared by several 724 isolates (strains) of the species, and a set of genes unique to an isolate. The size of each can 725 significantly differ from species to species. The core genome generally gives the basic 726 metabolic requirements of a certain species whereas the genetic plasticity of strains is generated by the other sequences (Medini et al., 2008 for general concepts and Earl et al., 727 728 2008 for specifics regarding bacilli). Data on the core genome aid significantly in defining the 729 metabolic potential of an organism. Flux Balance Analysis is then often used as a modeling 730 approach to find the possible steady states that the organism can attain (Schilling, 2000). In a 731 next step such models may be detailed further to incorporate molecular signaling data at various levels of complexity (see Ropers et al., 2006 for a highly detailed model of carbon 732 733 starvation in *Escherichia coli*). Such extension should at all times be subject to scrutiny 734 though in assessing its use given the investment of effort needed versus the (food) microbiological problem at hand needs to be considered. 735

736 7. Interaction between micro-organisms and their environment (foods) and microbial 737 evolution

738 Understanding short-term adaptations of microbes

The recent genomics revolution has facilitated the interpretation of the molecular basis ofmicrobial behavior. Examples stem from many fields and range from bacteria to filamentous

fungi. The response to high-end temperature stress conditions often characteristic of the

Confidential manuscript submitted to International Journal of Food Microbiology

742 manufacturing process of savoury products is nowadays studied at the molecular level. This is 743 most relevant to aerobic bacterial spore formers. Various strains of bacilli produce spores 744 resistant to temperatures up to and well above those of classical sterilization at 121°C (Oomes et al., 2007). Keijser et al., (2007) showed that spores express specific stress response genes 745 746 during germination, some of which are likely responsible for repair of incurred thermal 747 damage (see also Setlow 2006). In order to aim at understanding spore behavior after a 748 thermal stress, in particular mechanisms of heat damage repair, we now have the possibility of utilizing the genome information for *Bacillus subtilis* 168 (Kunst et al., 1997). 749

750 Cells subjected to acidic food conditions or to low water activity environments have been 751 studied extensively in the context of food microbiology. Specific examples of food 752 preservative stresses are those where cells have to respond to the antimicrobial action of a 753 weak-organic acid such as sorbic acid. The latter is the most widely used food preservative. 754 The common view is that the cells initially use energy driven pumps to extrude the acid from 755 the cytosol while upon full adaptation they induce the synthesis of pumps specific for lipophilic weak acids (discussed for yeast extensively in Mollapour et al., 2008; for recent 756 757 original research on sorbic acid stress response in vegetative bacteria see Ter Beek et al., 758 2008).

Stress adaptation of microorganisms in foods or upon being exposed to food processing conditions may lead to the induction of survival systems and could even induce virulence in pathogens. Many phenomena i.e. resistance to preservatives, oxidizing agents and natural extracts in foods are as important for successful infection as are mechanisms operative in 'in host', actual infection, and survival. Erickson and Doyle (2007) have illustrated this extensively for Shiga toxin-producing *Escherichia coli* and its survival on fresh produce, meat and in unpasteurized juices. Successful activation of stress response systems by some but not

Confidential manuscript submitted to International Journal of Food Microbiology

all strains may be instrumental in letting some strains adapt to the 'adverse' conditions in thefood chain.

768 Presence and development of microbial virulence traits in non-human environments

769 The following section will provide some selected examples of microbial virulence traits of

relevance to man of organisms present in non-human environments.

771 Foodborne pathogens may survive well in the animal production chain. Classical examples

include Campylobacter jejuni, an organism not pathogenic to avian species but highly

pathogenic to man (reviewed by Poly and Guerry, 2008). The organism is widely spread, as

was demonstrated again by, for instance, the studies of Fearnley *et al.* (2008). These authors

demonstrated the occurrence of hyperinvasive *Campylobacter* strains in isolates both from

poultry and from human sources. There is not much data yet on the molecular basis of

777 infection, be it that information on the intracellular signal transduction cascades of the

778 organism becomes more and more available (Boyd *et al.*, 2007).

779 Salmonella species are well-known pathogens for animals and man. Callaway et al. (2008) 780 have recently described their occurrence in various types of cattle. Sternberg et al. (2008) 781 described an outbreak of Salmonella infection in a Swedish dairy herd. Schmidt et al. (2008) 782 reported on Salmonella enterica infections in Swiss cattle in the summer of 2008. The various 783 serovars have meanwhile been characterized at the molecular level (Edwards et al., 2002). 784 The use of such data can be in two non mutually exclusive directions. On the one hand the 785 data provide information for use in quality control settings and epidemiological surveillance. 786 On the other hand the gathered information can be used as a starting point in the formulation 787 of novel research questions such as the molecular physiological mechanisms behind the 788 observed microbial ecology. Studies aiming at answering such questions will require next to 789 kinetic data on microbial metabolism measured at the population level, also quantitative data

Confidential manuscript submitted to International Journal of Food Microbiology

790 on cell - cell variation in microbial stress response in order to allow incorporation in next 791 generation predictive food microbiology models (McMeekin et al., 2007). Both for non-792 pathogenic B. subtilis and for pathogenic B. cereus, spore formation of strains attached to naturally occurring biofilms is a well known phenomenon (Lindsay et al., 2006). It has also 793 794 been documented nicely that spore-formation in a biofilm-like environment, a complex 795 colony, leads to spores with a higher thermal resistance than that observed in spores 796 originating from liquid cultures (Veening et al., 2006). The (thermal) stress resistance of 797 spores is again not a direct virulence trait but is does contribute to survival in the animal chain 798 as well as transfer to the human food chain (Huck et al., 2008). As such it is a crucial 799 determinant of the likelihood of intoxication of the host. Other such virulence characteristics 800 of e.g. Bacillus cereus include the resistance of the spores (and vegetative cells) to acid 801 facilitating the 'settlement' and toxin production of the organism in the intestine (Wijnands, 802 2008; see also Stenfors Arnesen et al., 2008).

803 *Predictive modeling*

804 Crucial is the conversion of molecular physiological 'analogue' data at the population level to 805 data at the level of single cells relevant to the prediction of the behavior of low-numbers 806 exposed to stressful environments (discussed in McMeekin et al., 2007; see for original 807 research amongst others Den Besten et al., 2007). This provides insight into the link between 808 the genome, gene-expression, protein and metabolic functional cellular units (see for the 809 original physiology data Balaban et al., 2004).). Koutsoumanis (2008) provides a clear 810 example of the variability in growth limits of individual Salmonella Enteritidis cells subjected 811 to NaCl stress. Another highly relevant example is the quantification of the germination and 812 outgrowth processes operative in bacterial endospores (Stringer et al., 2005; Smelt et al., 813 2008). Both examples do not yet include a mechanistic analysis e.g. at the level of inclusion 814 of genome-wide expression data. The initial challenge for future research is to do just that i.e.

Confidential manuscript submitted to International Journal of Food Microbiology

study at single cell / spore level the molecular physiology in order to enable the generation of
mechanistic models that describe the cellular heterogeneity in genetically homogeneous
microbial populations. Phenotypic heterogeneity in microbial populations mediated by bistable signaling networks is a much discussed topic in general microbiology (e.g. Veening et
al., 2008, Locke and Elowitz, 2009). To be of relevance to (predictive) food microbiology this
will require performing experiments under the relevant physiological (food) conditions and as
much as possible with the relevant food isolates.

822 Taxonomy

823 Regulators in the US use the GRAS approach (generally recognized as safe) to assess the 824 safety of microbes used in food production, while the EFSA employs the QPS (qualified 825 presumption of safety) system (Anonymous, 2007d). Basically it is assumed that a micro-826 organism that has been used for a considerable time in food manufacturing without causing 827 problems can be considered "safe". The difficulty in establishing these regulatory systems is the breadth of the taxonomic unit for which QPS or GRAS status can be conferred. If it only 828 829 can be conferred at strain level, a full risk assessment will still have to be performed for any 830 other strain, even those that are closely related to the ones having GRAS or QPS status. If it is 831 applied at an overly high level, e.g. genus, it can happen that pathogenic cousins of a safe 832 strain are wrongly considered harmless. The taxonomic units concerned are defined in the 833 OPS list and the list is reviewed every year to take into account changes in the taxonomy and 834 other considerations. If the history of safe use and the body of knowledge concerns only one 835 strain, this property cannot be generalized to apply to the entire species, (an example is the 836 *Enterococcus* spp). That is, if there is a risk that closely related strains of the safe strains are 837 pathogenic, the taxonomic unit, species or genus, is not given QPS status, unless the 838 pathogenic strains can be specifically identified, for instance by the presence of virulence

Confidential manuscript submitted to International Journal of Food Microbiology

- 839 factors. In the latter case the unit can be QPS, but with additional qualification. This is, for
- 840 instance, the case for some *Bacillus* species.
- 841 Guidelines for the selection of "safe" cultures for biopreservation exist in the area of feed, but
- 842 not for food (apart from the probiotics area). For general biopreservation, screening of
- 843 cultures for virulence factors or other genes coding for undesirable properties would be
- 844 relevant as well as studies of cultures possibly acquiring resistance.
- 845 For the approval of specific biopreservation agents (e.g. bacteriocins such as nisin)
- 846 governmental as well as academic and industry views on the criteria to be applied differ
- around the world. Inconsistencies can cause problems, especially where criteria for safety
- 848 evaluation or for an agent's effectiveness are too lenient or because they do not take sufficient
- 849 account of ill-informed use of such agents. The inverse, when a harmless strain is considered
- 850 pathogenic is less problematic as the only consequence is that the strain undergoes an
- 851 unnecessary safety assessment. Safe and robust use would need to follow general guidelines
- 852 on the steps to be taken, as defined in the safe design of the product or the process.
- 853 **8. R**i

8. Risk communication and education

854 Collaboration/communication between scientists

Historically, food systems used to be fairly simple; most foods were produced and eaten 855 locally. Nowadays, a large share of our diet is produced in another country, and not 856 857 uncommonly, several food ingredients come from different parts of the world (Käferstein et 858 al., 1997). Furthermore, we prefer to eat the food as fresh as possible (Doyle and Erickson, 859 2008). This trend increases the need for worldwide food safety systems and thus collaboration 860 and communication between all players in the food chain. As food chains extend or expand 861 from local chains into worldwide chains, more and different factors may affect food safety. This implies that for food safety management knowledge or information from different 862

Confidential manuscript submitted to International Journal of Food Microbiology

scientific disciplines needs to be combined. Furthermore, the format in which information ismade available needs to be standardized.

Food safety starts at primary production. To reduce the risk of foodborne gastroenteritis, 865 866 especially for foods to be eaten raw, such as fresh produce and shellfish, knowledge about 867 contamination routes and preventive measures is of great importance. In certain cases 868 interventions could be effective early in the primary production phase, including the production environment. For fresh produce, grown in the open field, Escherichia coli (in 869 870 particular the verocytotoxin producing strains VTEC) is, amongst others, a food safety 871 hazard. This is underlined by a massive foodborne infection outbreak in the USA in 2006 872 (Anonymous, 2007e) caused by baby spinach eaten raw. The probable source of the VTEC in 873 this outbreak was either irrigation water, or feces from cattle or wild boar. Cattle are a known 874 source of this pathogenic bacterium (Hussein and Sakuma, 2005), yet cow manure continues 875 to be used as a main soil fertilizer in organic farming (Anonymous, 2000b). Preventive 876 measures could be a change of feeding diet in order either to reduce numbers of VTEC shed by cattle (Diez-Gonzalez et al., 1998; Synge, 2000) or to reduce their survival in manure-877 878 amended soils (Franz et al., 2005). Thus, produce safety can be increased in this case by 879 combining agricultural science and (food) microbiology.

In other outbreaks, preventive measures are straighter foreword. In 2008, a *C. jejuni* outbreak in Alaska was linked to the consumption of raw peas contaminated on the field by Sandhill crane feces. The outbreak investigation identified a lack of chlorine residual in pea-processing water, which could have been easily prevented (Gardner and McLaughlin, 2008). Introducing buffer zones, set-back distances and fences to restrict wildlife access to the production environment of produce such as leafy greens may prevent problems with feral swine or deer (Atwill, 2008). Although these solutions seem sometimes fairly simple, they could only be

Confidential manuscript submitted to International Journal of Food Microbiology

taken due to a proper outbreak investigation that elucidated the (probable) source of the foodcontamination.

889 In order to stop an ongoing outbreak, such outbreak investigations should be carried out 890 quickly, which relies on close collaboration between different (scientific) disciplines such as 891 microbiologists, epidemiologists, wildlife control specialists, risk communicators, etc, often 892 represented by different organisations or institutes, which may hamper the investigation. 893 Although not standing on its own, the lack of proper communication between different scientific disciplines and/or institutes/departments during an outbreak investigation is shown 894 895 in the 2009 U.S.A. Salmonella Saintpaul outbreak caused by raw jalapeño and serrano pepper 896 in which more than 1400 cases were registered. This outbreak continued to spread due to 897 malfunctioning at the level of policy, the public-health system's organization and outbreak response, and its communications with the media and the public as concluded by the post-898 899 mortem investigation into this outbreak (Anonymous, 2008). This clearly shows that 900 collaboration of scientific disciplines is eminent to increase the safety of our food system. 901 Other examples that demonstrate the added value of combining different expertises to limit 902 the risk of foodborne illness are for instance Vibrio spp. in shellfish and mycotoxins in grain 903 (products). The quality of shellfish depends on the quality of the water they are grown in. 904 Pathogenic bacteria of importance in these types of products are *Vibrio* spp. The number of 905 vibrios present in the water is positively related to the water temperature (Motes et al., 1998) 906 and models that predict ocean water temperatures can be used to predict the level of these 907 pathogens in shellfish, thus combining (food) microbiology and oceanography, Ford et al., 908 2009). This knowledge was used by Californian lawmakers in order to ban the sale of raw 909 oysters from certain waters during the warmer months of the year (Anonymous, 2003). The 910 presence of mycotoxins on grain products is affected by weather conditions during growth 911 and harvest. As humidity increases, growth increases of the moulds that produce mycotoxins

Confidential manuscript submitted to International Journal of Food Microbiology

on these products. Thus meteorological data are a reliable indicator of the risk of the
concentration of mycotoxins on grain products (Schaafsma and Hooker, 2007; Van der FelsKlerx *et al.*, 2008).

915 To determine which intervention strategy is the best or most cost effective to be taken to 916 reduce risk of foodborne illness, a detailed risk assessment needs to be conducted and 917 combining with a proper economic analysis may prove necessary. This requires a highly multidisciplinary approach, involving microbiologists, epidemiologists, risk modelers, 918 919 economists and social scientists (Havelaar et al., 2007a). An example of this approach is the 920 CARMA project, carried out in the Netherlands, to evaluate options to reduce the risk of 921 campylobacteriosis due to consumption of chicken meat (Havelaar et al., 2007b). For risk assessment studies, many data are needed like prevalence and numbers of foodborne 922 923 pathogens in food. However, these data are not always available or are not in the correct 924 format. For instance, data on the occurrence of Salmonella in food is mainly available as "presence or absence in 25 g of product" as legislation requires testing based on this criterion 925 926 (Anonymous, 2005a). For risk assessment studies, however, the exact level of contamination 927 is important (Malorny et al., 2008). Even when data are available in the preferred format, 928 problems may arise with nomenclature of foods or lack of other necessary details. For 929 instance, a meatball can be either made from beef or pork or a combination of both. And 930 problems increase with multiple-ingredient products, such as in lasagna. Uniformity in 931 nomenclature of foods is, therefore, of great importance and a tool such as LanguaL, a food 932 description thesaurus (www.langual.org), can be very useful.

In many food consumption surveys foods are categorized as, for example "beef with or without sauce", whereas the degree of cooking of the meat is of greater microbiological relevance. To improve the use of data generated in food consumption surveys, closer collaboration between risk assessors, food microbiologists and nutritionists is needed.

Confidential manuscript submitted to International Journal of Food Microbiology

937 In the Netherlands, the Dutch Food and Consumer Product Safety Authority (VWA) works 938 closely together with the National Institute for Public Health and the Environment (RIVM) to 939 improve the quality and usefulness of the data obtained from routine monitoring programs of 940 the microbiological quality of foods. Some recent studies focused on the relative 941 microbiological risk to consumers associated with the consumption of fresh vegetables 942 (Pielaat and Wijnands, 2008) and prevalence of potentially pathogenic Bacillus cereus in food (Wijnands et al., 2006). A simple, spreadsheet-based tool is being developed to assess 943 consumer risks associated with such products using available data (Evers and Chardon, 2008). 944 These examples clearly show the benefits of inter- and intra- scientific collaboration. Close 945 946 personal collaboration may not always be necessary when data can be shared by other means. 947 In the scientific literature, many data are published that can be used by others. However, 948 translation of these data to a uniform data set is time consuming. In order to improve sharing 949 of data on microbial growth and inactivation, the ComBase Initiative was established, a collaboration between the Food Standards Agency and the Institute of Food Research from 950 951 the United Kingdom, the USDA Agricultural Research Service and its Eastern Regional 952 Research Center from the United States and the Australian Food Safety Centre of Excellence 953 (Combase Consortium, 2008). ComBase is a combined database of microbial responses to 954 food environments and data can be used for predictive modelling. Recently, Combase started 955 a collaboration with the Journal of Food Protection, which request authors to submit their data 956 to the Combase database. A clearinghouse of interdisciplinary data is offered by Foodrisk.org, 957 an initiative of the Joint Institute for Food Safety and Applied Nutrition (JIFSAN), and which 958 is a collaboration between the University of Maryland (UM) and the Food and Drug 959 Administration (FDA). The clearinghouse provides data and methodology on food safety risk analysis offered by the private sector, trade associations, federal and state agencies, and 960 961 international sources (www.foodrisk.org).

Confidential manuscript submitted to International Journal of Food Microbiology

962 In conclusion, although interdisciplinary collaboration sounds very promising, it must be

noted that it is generally not straightforward as, for instance, the data thus made available may

be limited or be in the wrong format. Despite this, with some extra effort more progress may

965 be achieved than would otherwise be the case.

966 Education

967 New trends in food consumption patterns, e.g. the consumption of raw foods, and the

968 introduction of a wide variety of newly developed food products with each their specific way

969 of preparation (e.g. ready-to-eat; ready-to-heat), in combination with an increase in the

970 number of vulnerable people, require clear communication about food safety aspects,

971 communication between industry, consumer and government.

Risk managers should be aware and understand public concerns about food safety as this must
be the basis of a risk management strategy (Frewer, 2004). Whether such a strategy will be
judged as effective by the public, will depend on the expertise of food risk managers (Van
Kleef *et al*, 2007) and cultural variation: what is effective in one country, is not always as
effective in another (Van Dijk *et al*, 2008).

977 Industry

When introducing new food products or new food preparation techniques, it is crucial to
provide pertinent information concerning safe food handling and preparation. The food
industry can contribute to food safety in several ways. Labeling, providing information about
correct storage conditions and ways of preparation, can contribute to food safety, although the
addition of more information is at odds with providing clear food labels (Mills *et al*, 2004).
Icons can be used. The food industry can further contribute to food safety by educating
professionals working along the food production chain. As an example, the efforts of the

Confidential manuscript submitted to International Journal of Food Microbiology

985 public-private partnership formed by the Industry Council for Development with FAO and986 WHO may be noteworthy (Motarjemi, 2006).

987 Consumers

988 Presence of pathogenic bacteria on raw food materials, such as meat and fresh produce is in 989 most cases not totally avoidable, therefore intervention strategies are also needed in 990 subsequent steps of the food chain to reduce the risk of foodborne illness for the consumer. At 991 the other side of the farm-to fork continuum the consumers also play an important role in 992 maintaining food safety. Different information campaigns therefore focus on improving home hygiene (Anonymous, 2008d, 2008e). However, the impact of such campaigns is often not 993 994 evaluated. A combined research project undertaken by social scientists, food microbiologists 995 and risk assessors showed the limited effect of such campaigns on reducing the actual level of 996 bacteria present in a meal, and on the associated risk of human illness (Nauta et al., 2008). Slovic (1987) developed a psychometric paradigm, which demonstrated that psychological 997 998 factors determine a person's response to different hazards, including those in the area of food 999 safety. Do consumers know what they should do in order to prepare a safe meal, in particular 1000 how to avoid cross-contamination and proper heating of reused food (Fischer *et al*, 2007)? 1001 According to Nauta et al (2008), consumers already possess the necessary knowledge 1002 regarding hygiene practices; this knowledge only needs to be activated. How to achieve this? 1003 What can we learn in this respect from other education campaigns, (e.g. smoking, alcohol, 1004 and fat), and from communication of medical product risks (Goldman, 2004), or from 1005 programs focusing on disease prevention and control (O'Loughlin et al., 1995; Sarraf-1006 Zadegan et al., 2003).

Health related behaviors are often differentially distributed across socioeconomic groups.Close collaboration with social scientists therefore seems logical. Communication with

Confidential manuscript submitted to International Journal of Food Microbiology

1009 different consumer groups requires different approaches and media. Information should be 1010 targeted to specific groups at risk, like single households, pregnant women or elderly people. 1011 This is because different groups have different food preparation and cooking habits and 1012 therefore are exposed to different levels of risk. Kornelis et al. (2007) showed that different 1013 consumers prefer different information sources when posing questions about food safety. 1014 Two-thirds of all consumers prefer information from either institutional or social sources. 1015 Your life, a free magazine containing articles about fashion, lifestyle and entertainment 1016 published by the UK National Health Service, is an illustrative example of such a social 1017 source. Members of the lower socioeconomic groups are more likely to respond to 1018 information from their direct social environment (Weenig and Midden, 1997). Apparently, 1019 communication with different consumer groups requires different approaches and media. The 1020 importance of educating children through the school system cannot be emphasized enough.

1021 Government

Food preparation and cooking practices are based on habits. This goes for consumers and 1022 1023 often also for food professionals in small food establishments, such as food services and 1024 restaurants. Since such behavior is difficult to change, education should be a life long learning 1025 process on general aspects of food safety as hygiene and contamination routes, starting at a 1026 young age, making use of all types of media, including video-gaming. Education in food 1027 safety aspects could be combined with nutritional information. Concurrent with the 1028 development of education programs, strategies should be developed to measure the impact of 1029 such programs (Nauta et al., 2008). In addition, governmental organizations should consider 1030 introducing and supporting specific education programs for consumers and professionals in 1031 small food preparation enterprises as well as for producers, especially for producers of fresh 1032 produce, as all have their responsibility with regard to food safety.

Confidential manuscript submitted to International Journal of Food Microbiology

- 1033 Finally, while education might be expected to improve food safety in the developed world, in
- 1034 developing countries, economic growth rather than education might be the best way to
- 1035 minimalize food-related mortality and morbidity amongst children.

1036 Acknowledgements

- 1037 The authors wish to thank all participants of the conference for their willingness to share their
- 1038 insights in plenary and group discussions. Special thanks are due to the plenary speakers (see
- 1039 contributions elsewhere in this Special Issue), the chairs and rapporteurs of breakout groups
- 1040 and the members of the Scientific Committee. These include, apart from the authors, Andrea
- 1041 Ammon, József Baranyi, Bob Buchanan, Patrice Buche, John D. Collins, Paul Cook, Mike
- 1042 Doyle, Aamir Fazil, Leon Gorris, Colin Hill, Marta Hugas, Marion Koopmans, Hilde Kruse,
- 1043 Hein van Lieverloo, Pia Makelä, Tom McMeekin, Diane Newell, Wim Ooms, Tom Quested,
- 1044 Peter Raspor, Tom Ross, Moez Sanaa, Don Schaffner, Claudia Stein, Mark Tamplin, Rob
- 1045 Tauxe, Mieke Uyttendaele, Yvonne Van Duynhoven, Didier Verloo, and Martin Wagner.

Confidential manuscript submitted to International Journal of Food Microbiology

1047 References

- 1048 Adak, G.K., Meakins, S.M., Yip, H., Lopman, B.A., O'Brien, S.J., 2005. Disease risks from foods, England and
- 1049 Wales, 1996-2000. Emerging Infectious Diseases 11, 365-372.
- 1050 Amar, C. F., East, C. L., Gray, J., Iturriza-Gomara, M., Maclure, E. A., and McLauchlin, J. (2007). Detection by
- 1051 PCR of eight groups of enteric pathogens in 4,627 faecal samples: re-examination of the English case-
- 1052 control Infectious Intestinal Disease Study (1993-1996). European Journal of Clinical Microbiology and
- 1053 Infectious Diseases 26, 311-23.
- 1054 Anonymous, 2000a. Dorland's Illustrated Medical Dictionary. 29th ed. WB Saunders Co, Philadelphia, PA.
- 1055 Anonymous, 2000b. National Organic Program. American Society for Microbiology. Available at

1056 http,//www.asm.org/Policy/index.asp?bid=3585. Accessed 9 December 2008.

- 1057 Anonymous, 2002. Regulation (EC) No 178/2002 of the European Parliament and of the Council of 28 January
- 1058 2002 laying down the general principles and requirements of food law, establishing the European Food
- Safety Authority and laying down procedures in matters of food safety. Official Journal of the European
 Communities, L31/1.
- 1061 Anonymous, 2003. Raw gulf oysters, labeling, written warnings and additional requirements. California

1062 Department of Public Health. CCR, Title 17, Section 13675, Available at

- 1063 http://www.cdph.ca.gov/services/Documents/fdb%20Raw%20Gulf%20Oyst%20Regs.pdf. Accessed 9
- 1064 December 2008.
- Anonymous, 2005a. Commission regulation (EC) No 2073/2005 of 15 November 2005 on microbiological
 criteria for foodstuffs. Official Journal of the European Communities, L 338/1.
- 1067 Anonymous, 2006. Development of practical risk management strategies based on microbiological risk
- assessment outputs. FAO and WHO, Rome and Geneva.
- 1069 Anonymous, 2007a. The Community Summary Report on trends and sources of zoonoses, zoonotic agents,
- antimicrobial resistance and foodborne outbreaks in the European Union in 2006. The EFSA Journal 130, 1-352.
- 1072 Anonymous, 2007b. Opinion of the Scientific Panel on Biological Hazards on microbiological criteria and
- 1073 targets based on risk analysis. The EFSA Journal 462, 1-29.

Confidential manuscript submitted to International Journal of Food Microbiology

1074 Anonymous, 2007c. Multistate outbreak of *Salmonella* serotype Tennessee infections associated with peanut

1075 butter--United States, 2006-2007. MMWR Morbidity Mortality Weekly Report 56, 521-524.

- 1076 Anonymous, 2007d. EFSA Scientific Committee, Introduction of a Qualified Presumption of Safety (QPS)
- 1077 approach for assessment of selected microorganisms referred to EFSA. The EFSA Journal 587, 1-16.
- 1078 Anonymous, 2007e. Investigation of an *Escherichia coli* O157,H7 outbreak associated with Dole pre-packaged
- 1079 spinach. California Food Emergency Response Team Sacramento/Alameda, CA, USA. Available at
- 1080 http://www.marlerclark.com/2006_Spinach_Report_Final_01.pdf. Accessed 22 December 2008.
- 1081 Anonymous, 2007f. Scientific Opinion of the Panel on Biological Hazards on a request from the European
- Commission on public health risks involved in the human consumption of reptile meat. The EFSA Journal
 578, 1-55.
- 1084 Anonymous, 2008a. Outbreak of *Salmonella* serotype Saintpaul infections associated with multiple raw produce
- 1085 items--United States, 2008. MMWR Morbidity Mortality Weekly Report 57, 929-934.
- Anonymous, 2008b. Climate change, Implications for food safety. Food and Agriculture Organization of the
 United Nations, Rome.
- 1088 Anonymous, 2008c. Report of the Task Force on Zoonoses Data Collection on the Analysis of the baseline
- 1089 survey on the prevalence of *Salmonella* in slaughter pigs, in the EU, 2006-2007[1] Part A, *Salmonella*
- 1090 prevalence estimates. The EFSA Journal 135, 1-11.
- Anonymous, 2008d. Fight Bac! Educating consumers about safe food handling. Partnership for Food Safety
 Education. Available at http://www.fightbac.org/. Accessed 9 December 2008.
- 1093 Anonymous, 2008e. Hygiëne. Voedingscentrum. Available at
- 1094 http://www.voedingscentrum.nl/EtenEnVeiligheid/Hygiëne/. Accessed 9 December 2008.
- 1095 Anonymous, 2009. Multistate outbreak of Salmonella infections associated with peanut butter and peanut butter-
- 1096 containing products--United States, 2008-2009. Morbidity Mortality Weekly Report 58, 85-90.
- 1097 Ara, K., Ozaki, K., Nakamura, K., Yamane, K., Sekiguchi, J. Ogasawara, N., 2007. Bacillus minimum genome
- 1098 factory, effective utilization of microbial genome information. Biotechnology and Applied Biochemistry. 46,
 1099 169-178.
- 1100 Arvanitoyannis, I.S. Traikou, A., 2005. A comprehensive review of the implementation of hazard analysis
- 1101 critical control point (HACCP) to the production of flour and flour-based products. Critical Reviews in Food

Confidential manuscript submitted to International Journal of Food Microbiology

- 1102 Science and Nutrition 45, 327-370.
- Balaban, N.Q., Merrin, J., Chait, R., Kowalik, L., Leibler, S., 2004. Bacterial persistence as a phenotypic switch.
 Science 305, 1622-1625.
- 1105 Batz, M.B., Doyle, M.P., Morris, G., Painter, J., Singh, R., Tauxe, R.V., Taylor, M.R., Lo Fo Wong, D.M.A.,
- 1106 2005. Attributing Illness to Food . Emerging Infectious Diseases 11, 993-999.
- 1107 Beckmann, M., Enot, D.P., Overy, D.P. and Draper, J., 2007. <u>Representation, comparison, and interpretation of</u>
- 1108 metabolome fingerprint data for total composition analysis and quality trait investigation in potato cultivars.
 1109 Journal of Agricultural and Food Chemistry. 55, 3444-3451.
- 1110 Boschi-Pinto, C., Velebit, L., Shibuya, K., 2008. Estimating child mortality due to diarrhoea in developing

1111 countries. Bulletin of the World Health Organization 86, 710-717.

- 1112 Boyd, A., Philbin, V.J., Smith, A.L., 2007. Conserved and distinct aspects of the avian Toll-like receptor (TLR)
- system, implications for transmission and control of bird-borne zoonoses. Biochemical Society Transactions
 35, 1504-1507.
- Boyle, R.J., Robins-Browne, R.M., Tang, M.L., 2006. Probiotic use in clinical practice: what are the risks?
 American Journal of Clinical Nutrition 83, 1256-64.
- 1117 Brzuszkiewicz, E., Bruggemann, H., Liesegang, H., Emmerth, M., Olschlager, T., Nagy, G., Albermann, K.,
- 1118 Wagner, C., Buchrieser, C., Emody, L., Gottschalk, G., Hacker, J., Dobrindt, U., 2006. How to become a
- 1119 uropathogen, comparative genomic analysis of extraintestinal pathogenic *Escherichia coli* strains.
- 1120 Proceedings of the National Acadamy of Sciences USA 103, 12879–12884.
- 1121 Callaway, T.R., Edrington, T.S., Anderson, R.C., Byrd, J.A., Nisbet, D.J., 2008. Gastrointestinal microbial
- ecology and the safety of our food supply as related to *Salmonella*. Journal of Animal Science 86, E 163-
- 1123 172.
- 1124 Combase Consortium, 2008. ComBase. Available at http://www.combase.cc/, Accessed 9 December 2008.
- Cretikos, M., Telfer, B., McAnulty, J., 2008. Enteric disease outbreak reporting, New South Wales, Australia,
 2000 to 2005. New South Wales Public Health Bulletin 19, 3-7.
- 1127 Dalton, C.B., Austin, C.C., Sobel, J., Hayes, P.S., Bibb, W.F., Graves, L.M., Swaminathan, B., Proctor, M.E.,
- 1128 Griffin, P.M., 1997. An outbreak of gastroenteritis and fever due to *Listeria monocytogenes* in milk. New
- England Journal of Medicine 336, 100-105.

- 1130 Den Besten, H.M.W., Ingham, C.J., Van Hylckama Vlieg, J.E.T., Beerthuyzen, M.M., Zwietering, M.H., Abee,
- 1131 T., 2007. Quantitative analysis of population heterogeneity of the adaptive salt stress response and growth
- 1132 capacity of *Bacillus cereus* ATCC 14579. Applied and Environmental Microbiology 73, 4797-4804.
- 1133 De Wit, M.A.S., Koopmans, M.P.G., Kortbeek, L.M., van Leeuwen, N.J., Bartelds, A.I.M., van Duynhoven,
- 1134 Y.T.H.P., 2001a. Gastroenteritis in sentinel general practices, The Netherlands. Emerging Infectious
- 1135 Diseases 7, 82-91.
- 1136 De Wit, M.A.S., Koopmans, M.P.G., Kortbeek, L.M., Van Leeuwen, N.J., Vinjé, J., Van Duynhoven, Y.T.H.P.,
- 2001b. Etiology of gastroenteritis in sentinel general practices in the Netherlands. Clinical Infectious
 Diseases 33, 280-288.
- 1139 De Wit, M.A.S., Koopmans, M.P.G., Kortbeek, L.M., Wannet, W.J., Vinjé, J., Van Leusden, F., Bartelds,
- 1140 A.I.M., Van Duynhoven, Y.T.H.P., 2001c. Sensor, a population-based cohort study on gastroenteritis in the
- 1141 Netherlands, incidence and etiology. American Journal of Epidemiology 154, 666-674.
- Diez-Gonzalez, F., Callaway, T.R., Kizoulis, M.G., Russell J.B., 1998. Grain feeding and the dissemination of
 acid-resistant *Escherichia coli* from cattle. Science 281, 1666-1668.
- 1144 Doyle, M.P., Erickson, M.C., 2008. Summer meeting 2007 the problems with fresh produce: an overview.
- 1145 Journal of Applied Microbiology105, 317-330.
- Earl, A.M., Losick, R., Kolter, R., 2008. Ecology and genomics of *Bacillus subtilis*. Trends in Microbiology 16,
 269-275.
- Edwards, R.A., Olsen, G.J., Malov, S.R., 2002. Comparative genomics of closely related salmonellae. Trends in
 Microbiology 10, 94-99.
- 1150 Ellis, D.I., Broadhurst, D., Rowland, J.J. and Goodacre, R., 2007. Rapid detection method for microbial spoilage
- 1151 using FT-IR and machine learning. In: Van Amerongen, A., Barug, D and Lauwaars, M. (eds.). Rapid
- Methods (for Food and Feed Quality Determination). Wageningen Academic Publishers, Wageningen, the
 Netherlands, pp. 73-84.
- Erickson, M.C., Doyle, M.P., 2007. Food as a vehicle for transmission of Shiga toxin-producing *Escherichia coli*. Journal of Food Protection 70, 2426-2449.
- 1156 Evers. E.G., Chardon, J.E., 2008. A swift quantitative microbiological risk assessment (sQMRA) tool. Abstract
- 1157 K7, Food Micro 2008, 1-4 September, Aberdeen, Scotland.

- 1158 Fearnley, C., Manning, G., Bagnal, M., Javed, M.A., Wassenaar, T.M., Newell, D.G., 2008. Identification of
- 1159 hyperinvasive *Campylobacter jejunii* strains isolated from poultry and human clinical sources. Journal of
- 1160 Medical Microbiology 57, 570-580.
- 1161 Fischer, A.R., de Jong, A.E., de Jonge, R., Frewer, L.J., Nauta, M.J., 2005. Improving food safety in the
- domestic environment, the need for a transdisciplinary approach. Risk Analysis 25,503-517.
- 1163 Fischer, A.R., De Jong, A.E., Van Asselt, E.D., De Jonge, R., Frewer, L.J., Nauta, M.J., 2007. Food safety in the
- domestic environment, an interdisciplinary investigation of microbial hazards during food preparation. Risk
 Analysis 27, 1065-1082.
- 1166 Flint, J.A., Van Duynhoven, Y.T., Angulo, F.J., DeLong, S.M., Braun, P., Kirk, M., Scallan, E., Fitzgerald, M.,
- Adak, G.K., Sockett, P., Ellis, A., Hall, G., Gargouri, N., Walke, H., Braam, P., 2005. Estimating the burden
- 1168 of acute gastroenteritis, foodborne disease, and pathogens commonly transmitted by food, an international
- review. Clinical Infectious Diseases 41, 698-704.
- Ford, T.E., Colwell, R.R., Rose, J.B., Morse, S.S., Rogers, D.J., Yates, T.L., 2009. Using Satellite Images of
 Environmental Changes to Predict Infectious Disease Outbreaks. Emerging Infectious Diseases 15, 13411172 1345.
- 1173 Franz, E., van Diepeningen, A.D., De Vos, O.J., van Bruggen A.H., 2005. Effects of cattle feeding regimen and
- soil management type on the fate of *Escherichia coli* O157:H7 and *Salmonella enterica* serovar
- 1175 Typhimurium in manure, manure-amended soil, and lettuce. Applied and Environmental Microbiology 71,1176 6165-6174.
- Fraser-Liggett, C. M., 2005. Insights on biology and evolution from microbial genome sequencing. Genome
 Research 15, 1603–1610.
- 1179 Frewer, L., 2004. The public and effective risk communication. Toxicology Letters 149, 391-397.
- 1180 Gebreyes, W.A., Bahnson, P.B., Funk, J.A., McKean, J., Patchanee, P. 2008. Seroprevalence of Trichinella,
- 1181 Toxoplasma, and Salmonella in antimicrobial-free and conventional production systems. Foodborne
- 1182 Pathogens and Disease 5, 199-203.
- Gerner-Smidt, P., Hise, K., Kincaid, J., Hunter, S., Rolando, S., Hyytia-Trees, E., Ribot, E.M., Swaminathan, B.,
 2006. PulseNet USA, a five-year update. Foodborne Pathogens and Disease 3, 9-19.
- 1185 Ginsberg, M.M., Granzow, L., Teclaw, R.F., Gaul, L.K., Bagdure, S., Cole, A., Drumgoole, R., Barzilay, E.J.,

Confidential manuscript submitted to International Journal of Food Microbiology

- 1186 Biggerstaff, M.S., Lynch, M.F., Maslanka, S.E., Williams, I.T., Juliao, P.C., Barton Behravesh, C., Olson,
- 1187 C.K., 2007. Botulism associated with commercially canned chili sauce--Texas and Indiana, July 2007.
- 1188 MMWR Morbidity Mortality Weekly Report 56, 767-769.
- 1189 Goldman, S.A., 2004. Communication of medical product risk, how effective is effective enough? Drug Safety
- 1190 27, 519-534.
- Hall, G., Kirk, M.D., Becker, N., Gregory, J.E., Unicomb, L., Millard, G., Stafford, R., Lalor, K., 2005.
- 1192 Estimating foodborne gastroenteritis, Australia. Emerging Infectious Diseases 11, 1257-1264.
- 1193 Hald, T., Vose, D., Wegener, H.C., Koupeev, T., 2004. A Bayesian approach to quantify the contribution of
- animal-food sources to human salmonellosis. Risk Analysis 24, 255-269.
- Hansen, J., Holm, L., Frewer, L., Robinson, P., Sandøe, P., 2003. Beyond the knowledge deficit, recent research
- 1196 into lay and expert attitudes to food risks. Appetite 41, 111-121.
- 1197 Havelaar, A.H., Mangen, M.J., de Koeijer, A.A., Bogaardt, M.J., Evers, E.G., Jacobs-Reitsma, W.F., van Pelt,
- 1198 W., Wagenaar, J.A., de Wit, G.A., van der Zee, H., Nauta, M.J., 2007a. Effectiveness and efficiency of

1199 controlling *Campylobacter* on broiler chicken meat. Risk Analysis 27: 831-844.

- 1200 Havelaar, A.H., Bräunig, J., Christiansen, K., Cornu, M., Hald, T., Mangen, M.J., Molbak, K., Pielaat, A., Snary,
- 1201 E., van Pelt, W., Velthuis, A., Wahlstrom, H., 2007b. Towards an integrated approach in supporting

1202 microbiological food safety decisions. Zoonoses and Public Health 54: 103-117.

- 1203 Havelaar, A.H., van Duynhoven, Y.T.H.P., van Pelt, W., 2008. Microbiologische ziekteverwekkers in voedsel.
- 1204 Omvang van het probleem . Hoe vaak komt ziekte als gevolg van microbiologische ziekteverwekkers in
- 1205 voedsel voor? In: Volksgezondheid Toekomst Verkenning, Nationaal Kompas Volksgezondheid. Bilthoven,
- 1206 RIVM. Available at http://www.rivm.nl/vtv/object_document/o3617n22451.html. Accessed 12 September
 1207 2008.
- Herwaldt, B.L., 2000. *Cyclospora cayetanensis*, a review, focusing on the outbreaks of cyclosporiasis in the
 1209 1990s. Clinical Infectious Diseases 31, 1040-57.
- Hiller, N. L. *et al.* 2007. Comparative genomic analyses of seventeen *Streptococcus pneumoniae* strains, insights
 into the pneumococcal supragenome. Journal of Bacteriology 189, 8186–8195.
- 1212 Hoogenboom, L. A.; Bokhorst, J. G.; Northolt, M. D.; van de Vijver, L. P.; Broex, N. J.; Mevius, D. J.; Meijs, J.
- 1213 A., and Van der Roest, J., 2008. Contaminants and microorganisms in Dutch organic food products: a

- 1214 comparison with conventional products. Food Additives & Contaminants. Part A: Chemistry, Analysis,
- 1215 Control, Exposure & Risk Assessment. 25:1195-1207.
- 1216 Hounsome, N., Hounsome, B., Tomos, D., Edwards-Jones, G., 2008. Plant metabolites and nutritional quality of
- 1217 vegetables. Journal of Food Science 73, R48-65.
- 1218 Huck, J.R., Sonnen, M., Boor, J.K., 2008. Tracking heat resistant, cold-thriving fluid milk spoilage bacteria from
- 1219 farm to packaged product. Journal of Dairy Science 91, 1218-1228.
- Hussein, H.S., Sakuma. T., 2005. Prevalence of shiga toxin-producing *Escherichia coli* in dairy cattle and their
 products. Journal of Dairy Science 88, pp. 450-465.
- Jones, T.F., Scallan, E., and Angulo, F.J., 2007. FoodNet, overview of a decade of achievement. Foodborne
 Pathogens and Disease 4, 60-66.
- Käferstein, F. K., Motarjemi, Y., Bettcher, D. W., 1997. Foodborne disease control: a transnational challenge.
 Emerging Infectious Diseases 31, 503-510.
- Keijser B.J.F., Ter Beek, A., Rauwerda, H., Schuren, F., Montijn, R., van der Spek, H., Brul, S. 2007. Analysis
 of temporal gene expression during *Bacillus subtilis* spore germination and outgrowth. Journal of
 Bacteriology 189, 3624-3634.
- 1229 Kirk, M.D., Little, C.L., Lem, M., Fyfe, M., Genobile, D., Tan, A., Threlfall, J., Paccagnella, A., Lightfoot, D.,
- 1230 Lyi, H., McIntyre, L., Ward, L., Brown, D.J., Surnam, S., Fisher, I.S., 2004. An outbreak due to peanuts in
- 1231 their shell caused by *Salmonella enterica* serotypes Stanley and Newport--sharing molecular information to
- solve international outbreaks. Epidemiology and Infection 132, 571-517.
- 1233 Kirk, M.D., McKay, I., Hall, G.V., Dalton, C.B., Stafford, R., Unicomb, L., Gregory, J., 2008. Food safety,
- 1234 foodborne disease in Australia, the OzFoodNet experience. Clinical Infectious Diseases 47, 392-400.
- 1235 Kornalijnslijper, J.E., Rahamat-Langendoen, J.C., Van Duynhoven, Y.T.H.P., 2008. Volksgezondheidsaspecten
- 1236 van veehouderijmegabedrijven in Nederland: zoönosen en antibioticumresistentie. Bilthoven, the
- 1237 Netherlands: RIVM. Briefrapportnr. 215011002. Available at
- 1238 http://www.rivm.nl/bibliotheek/rapporten/215011002.pdf.
- 1239 Kornelis, M., de Jonge J., Frewer, L., Dagevos, H., 2007. Consumer selection of food-safety information
- 1240 sources. Risk. Analysis 27, 327-335.
- 1241 Kort, R., Keijser, B.J., Caspers, M.P.M., Schuren, F., Montijn, R., 2008. Transcriptional activity around bacterial

Confidential manuscript submitted to International Journal of Food Microbiology

- 1242 cell death reveals molecular biomarkers for cell viability. BMC Genomics 9, 590 (Published ahead of print).
- 1243 Koutsoumanis, K., 2008. A study on the variability in the growth limits of individual cells and its effect on the
- behavior of microbial populations. International Journal of Food Microbiology 128, 116-121.
- 1245
- 1246 Kroneman, A.; Verhoef, L.; Harris, J.; Vennema, H.; Duizer, E.; van Duynhoven, Y.; Gray, J.; Iturriza, M.;
- 1247 Bottiger, B.; Falkenhorst, G.; Johnsen, C.; von Bonsdorff, C. H.; Maunula, L.; Kuusi, M.; Pothier, P.;
- 1248 Gallay, A.; Schreier, E.; Hohne, M.; Koch, J.; Szucs, G.; Reuter, G.; Krisztalovics, K.; Lynch, M.;
- 1249 McKeown, P.; Foley, B.; Coughlan, S.; Ruggeri, F. M.; Di Bartolo, I.; Vainio, K.; Isakbaeva, E.; Poljsak-
- 1250 Prijatelj, M.; Grom, A. H.; Mijovski, J. Z.; Bosch, A.; Buesa, J.; Fauquier, A. S.; Hernandez-Pezzi, G.;
- 1251 Hedlund, K. O., Koopmans, M., 2008 Analysis of integrated virological and epidemiological reports of
- 1252 norovirus outbreaks collected within the foodborne viruses in Europe Network from 1 July 2001 to 30 June
- 1253 2006. Journal of Clinical Microbiology 46,2959-2965.
- 1254 Kunst, F., Ogasawara, N., Moszer, I., Albertini, A.M., Alloni, G., Azevedo, V., Bertero, M.G., Bessieres, P.,
- 1255 Bolotin, A., Borchert, S., Borriss, R., Boursier, L., Brans, A., Braun, M., Brignell, S.C., Bron, S., Brouillet,
- 1256 S., Bruschi, C.V., Caldwell, B., Capuano, V., Carter, N.M., Choi, S.K., Codani, J.J., Connerton, I.F.,
- 1257 Danchin, A., *et al.*, 1997. The complete genome sequence of the gram-positive bacterium *Bacillus subtilis*.

1258 Nature 390, 249-256.

- 1259 Lindsay, D., Brözel, V.S., Von Holy, A., 2006. Biofilm-spore response in *Bacillus cereus* and *Bacillus subtilis*
- 1260 during nutrient limitation. Journal of Food Protection 69, 1168-1172.
- Locke, J.C. and Elowitz, M.B. 2009. Using movies to analyse gene circuit dynamics in single cells. Nature
 Reviews Microbiology 7, 383-392.
- 1263 Magnino S., Colin P., Dei-Cas E., Madsen M., McLauchlin J., Nockler K., Maradona, M. P., Tsigarida, E.,
- Vanopdenbosch, E., Van Peteghem, C., 2009. Biological risks associated with consumption of reptile
 products. International Journal of Food Microbiology 134,163-175.
- 1266 Majowicz, S.E., Edge, V.L., Fazil, A., McNab, W. B., Dore, K. A., Sockett, P. N., Flint, J. A., Middleton, D.,
- 1267 McEwen, S. A., Wilson, J. B, 2005. Estimating the under-reporting rate for infectious gastrointestinal illness
- in Ontario. Canadian Journal of Public Health 96, 178-181.
- 1269 Malorny, B., Löfström, C., Wagner, M., Krämer, N., Hoorfar, J., 2008. Enumeration of Salmonella bacteria in

Confidential manuscript submitted to International Journal of Food Microbiology

- 1270 food and feed samples by real-time PCR for quantitative microbial risk assessment. Applied and
- 1271 Environmental Microbiology 74, 1299-1304.
- 1272 McMeekin, T.A., Baranyi, J., Bowman, J., Dalgaard, P., Kirk, M., Ross, T., Schmid, S., Zwietering, M.H., 2006.
- 1273 Information systems in food safety management. International Journal of Food Microbiology 112, 181-194.
- 1274 McMeekin, T.A., Mellefont, L.A., Ross, T., 2007. Predictive microbiology, past, present and future. In,
- 1275 Modelling microorganisms in food Brul, S., van Gerwen, S., Zwietering, M. (eds.), Woodhead, Cambridge
- 1276 (UK). pp. 7-21.
- McMeekin, T.A., Ross, T., 2002. Predictive microbiology, providing a knowledge-based framework for change
 management. International Journal of Food Microbiology 78, 133-153.
- 1279 Medini, D., Serruto, D., Parkhill, J., Relman, D.A., Donati, C., Moxon, R., Falkow, S., Rappuoli, R., 2008.
- 1280 Microbiology in the post-genomic era. Nature Reviews Microbiology 6, 419-430.
- 1281 Mevius D.J., Van Pelt, W. (editors), 2006. Monitoring of antimicrobial resistance and antibiotic usage in animals

in the Netherlands in 2005. Lelystad, the Netherlands: Central Institute for Animal Disease Control.

1283 Available on-line: http://www.cvi.wur.nl/UK/publications/otherpublications/maran/.

- 1284 Michel, P., Wilson, J.B., Martin, S.W., Clarke, R.C., McEwen, S.A., Gyles, C.L., 2000. Estimation of the under-
- reporting rate for the surveillance of *Escherichia coli* O157:H7 cases in Ontario, Canada. Epidemiology and
 Infection 125, 35-45.
- 1287 Mills, E.N.C., Valovirta, E., Madsen, C., Taylor, S.L., Vieths, S., Anklam, E., Baumgartner, S., Koch, P., Crevel,
- R.W.R., Frewer, L., 2004. Information provision for allergic consumers-where are we going with food
 allergen labeling? Allergy 59, 1262-1268.
- 1290 Mollapour, M., Shepherd, A., Piper, P.W., 2008. Novel stress responses facilitate Saccharomyces
- *cerevisiae* growth in the presence of the monocarboxylate preservatives. Yeast 25, 169-177.
- 1292 Monday, S.R., Beisaw, A. and Feng, P.C.H. 2007. Identification of Shiga toxigenic Escherichia coli
- seropathotypes A and B by multiplex PCR. Molecular and Cellular Probes 21, 308-311.
- 1294 Motarjemi, Y., 2006. ICD in perspective: putting social responsibility into practice. Food Control 17, 1018-1022.
- 1295 Motes, M.L., DePaola, A., Cook, D.W., Veazey, J.E., Hunsucker, J.C., Garthright, W.E., Blodgett, R.J., Chirtel,
- 1296 S.J., 1998. Influence of water temperature and salinity on *Vibrio vulnificus* in Northern Gulf and Atlantic

- 1297 Coast oysters (*Crassostrea virginica*). Applied and Environmental Microbiology 64, 1459-1465.
- 1298 Nauta, M., Fischer, A., van Asselt, E., de Jong, A., Frewer, L., de Jonge, R., 2008. Food safety in the domestic
- 1299 environment, the effect of consumer risk information on human disease risk. Risk Analysis 28, 179-192.
- 1300 Noteborn, H.P.J.M., Ooms, W., De Prado, M., 2005. Emerging risks identification in food and feed for human
- 1301 health. Food and Consumer Product Safety Authority, The Hague.
- 1302 O'Brien, S.J., Gillespie, I.A., Sivanesan, M.A., Elson, R., Hughes, C., Adak, G.K., 2006. Publication bias in
- 1303 foodborne outbreaks of infectious intestinal disease and its implications for evidence-based food policy.
- England and Wales 1992-2003. Epidemiology and Infection 134, 667-74.
- 1305 O'Loughlin, J., Paradis, G., Kishchuk, N., Gray-Donald, K., Renaud, L., Fines, P., Barnett, T., 1995. Coeur en
- 1306 santé St-Henri- a heart health promotion programme in Montreal, Canada, design and methods for
- evaluation. Journal of Epidemiology and Community Health 49, 495-502.
- 1308 Oomes, S.J., van Zuijlen, A.C., Hehenkamp, J.O., Witsenboer, H., van der Vossen, J.M. and Brul, S., 2007. The
- 1309 characterisation of *Bacillus* spores occurring in the manufacturing of (low acid) canned products.
- 1310 International Journal of Food Microbiology 120, 85-94.
- 1311 Pielaat, A., Wijnands, L., 2008. Survey analysis of microbiological contamination in the fresh vegetables food
- 1312 chain and the associated relative risk to consumers. Food Micro 2008, Aberdeen, Scotland, p. 85.
- Pires, S.M., Evers, E.G., Van Pelt, W., Ayers, T., Scallan, E., Angulo, F.J., Havelaar, A.H., Hald, T., 2009.
 Attributing the human disease burden of foodborne infections to specific sources. Foodborne Pathogens and Disease 6, 417-423.
- 1316 Poly, F., Guerry, P., 2008. Pathogenesis of *Campylobacter*. Current Opinion in Gastroenterology 24, 27-31.
- 1317 Perna, N.T., Plunkett, G., 3rd, Burland, V., Mau, B., Glasner, J.D., Rose, D.J., Mayhew, G.F., Evans, P.S.,
- 1318 Gregor, J., Kirkpatrick, H.A., Posfai, G., Hackett, J., Klink, S., Boutin, A., Shao, Y., Miller, L., Grotbeck,
- 1319 E.J., Davis, N.W., Lim, A., Dimalanta, E.T., Potamousis, K.D., Apodaca, J., Anantharaman, T.S., Lin, J.,
- 1320 Yen, G., Schwartz, D.C., Welch, R.A., Blattner, F.R., 2001. Genome sequence of enterohaemorrhagic
- 1321 *Escherichia coli* O157:H7. Nature 409, 529–533.
- 1322 Pontrelli, G., Boccia, D., Di Renzi, M., Massari, M., Giugliano, F., Celentano, L.P., Taffon, S., Genovese, D., Di
- 1323 Pasquale, S., Scalise, F., Rapicetta, M., Croci, L., Salmaso, S., 2008. Epidemiological and virological
- 1324 characterization of a large community-wide outbreak of hepatitis A in southern Italy. Epidemiology and
- 1325 Infection 136, 1027-1034.

- 1326 Roy, S.L., Scallan, E., Beach, M.J., 2006. The rate of acute gastrointestinal illness in developed countries.
- 1327 Journal of Water Health 4, Supplement 2, 31-69.
- Ropers, D., de Jong, H., Page, M., Schneider, D., Geiselmann, J., 2006. Qualitative simulation of the carbon
 starvation response in *Escherichia coli*. BioSystems 84, 124–152.
- 1330 Rossmanith, P., Süss, B., Wagner, M., Hein, I., 2007. Development of a matrix lysis for concentration of gram
- positive bacteria from food and blood. Journal of Microbiological Methods 69, 504-511.
- 1332 Sarraf-Zadegan, N., Sadri, G., Malek Afzali, H., Baghaei, M., Mohammadi Fard, N., Shahrokhi, S., Tolooie, H.,
- 1333 Poormoghaddas, M., Sadeghi, M., Tavassoli, A., Rafiei, M., Kelishadi, R., Rabiei, K., Bashardoost, N.,
- Boshtam, M., Asgary, S., Naderi, G., Changiz, T., and Yousefie, A., 2003. Isfahan healthy heart program, a
- 1335 comprehensive community-based programme for cardiovascular disease prevention and control. Design,
- 1336 methods and initial experience. Acta Cardiologica 58, 309-320.
- 1337 Scaria, J., Palaniappan, R.U., Chiu, D., Phan, J.A., Ponnala, L., McDonough, P., Grohn Y.T., Porwollik, S.,
- 1338 McClelland, M., Chioa, C.S., Chu, C., Chang, Y.F., 2008. Microarray for molecular typing of Salmonella
- 1339 *enterica* serovars. Molecular and Cellular Probes. 22, 238-243.
- 1340 Schaafsma, A.W., Hooker, D.C., 2007. Climatic models to predict occurrence of *Fusarium* toxins in wheat and
- 1341 maize. International Journal of Food Microbiology 119, 116-125.
- 1342 Schilling, C.H., Edwards, J.S., Letscher, D., Palsson B.O., 2000. Combining pathway analysis with flux balance
- analysis for the comprehensive study of metabolic systems, Biotechnology and Bioengineering 71, 286–306.
- 1344 Schmidt, H., Hachler, H., Stephan, R., Baumgartner, A. and Boubaker, K., 2008. Outbreak of Salmonella
- *enterica* serovar Typhimurium in Switzerland, May-June 2008, implications for production and control of
 meat preparations. Eurosurveillance 13, 44, 19020.
- 1347 Senhaji, A.F., 1977. The protective effect of fat on the heat resistance of bacteria (II). International Journal of
- 1348Food Science and Technology 12: 217-230.
- Setlow, P., 2006. Spores of *Bacillus subtilis*, their resistance to radiation, heat and chemicals. Journal of Applied
 Microbiology 101, 514–525.
- Siguier, P., Filee, J., Chandler, M., 2006. Insertion sequences in prokaryotic genomes. Current Opinion in
 Microbiology 9, 526-31.
- 1353 Simonsen, J., Strid, M.A., Molbak, K., Krogfelt, K.A., Linneberg, A., Teunis, P., 2008. Sero-epidemiology as a

Confidential manuscript submitted to International Journal of Food Microbiology

- tool to study the incidence of *Salmonella* infections in humans. Epidemiology and Infection 136, 895-902.
- 1355 Slovic, P., 1987. Perception of risk. Science 236, 280-285.
- 1356 Smelt, J.P.P.M., Bos, A.P., Kort, R. and Brul, S., 2008. Modelling the effect of sub(lethal) heat
- 1357 treatment of *Bacillus subtilis* spores on germination rate and outgrowth to exponentially growing vegetative
- cells. International Journal of Food Microbiology 128, 34-40.
- 1359 Stein, C., Kuchenmuller, T., Hendrickx, S., Pruss-Ustun, A., Wolfson, L., Engels, D., Schlundt, J., 2007. The
- 1360 Global Burden of Disease Assessments-WHO Is Responsible? PLoS Neglected Tropical Diseases 1, e161.
- Stenfors Arnesen, L.P., Fagerlund, A., Granum, P.E., 2008. From soil to gut, *Bacillus cereus* and its food
 poisoning toxins. FEMS Microbiological Reviews 32, 579-606.
- 1363 Sternberg, S., Johnsson, A., Aspan, A., Bergström, K., Kallay, T.B., Szanto, E., 2008. Outbreak of Salmonella

1364 Thompson infection in a Swedish dairy herd. Veterinary Record 163, 596-599.

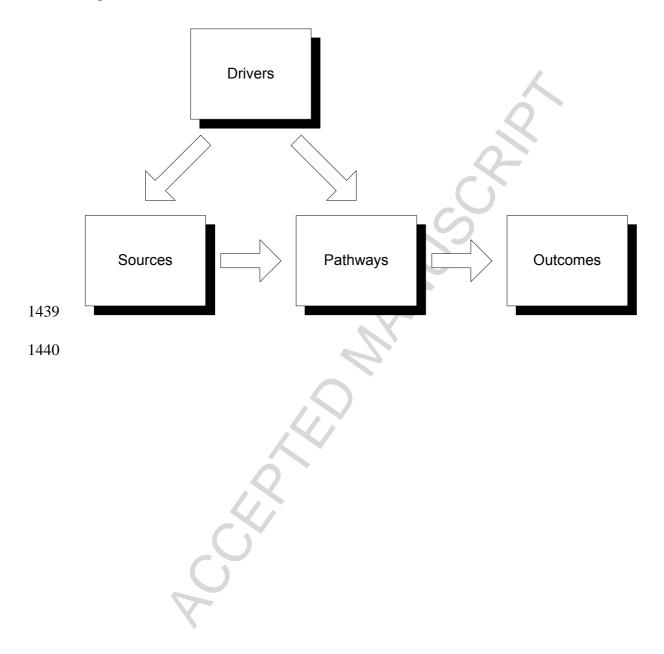
- 1365 Stringer, S.C., Webb, M.D., George, S.M., Pin, C., Peck, M.W., 2005. Heterogeneity of times required for
- germination and outgrowth from single spores of nonproteolytic *Clostridium botulinum*. Applied and
 Environmental Microbiology 71, 4998-5003.
- Synge, B.A., 2000. Verocytotoxin-producing *Escherichia coli*, a veterinary view. Journal of Applied
 Microbiology 88, p. 31S-378.
- Tait, J., Meagher, L., Lyall, C., Suk, J., 2006. Foresight. Infectious Diseases: preparing for the future. T2: Risk
 Analysis. Office of Science and Innovation, London.
- Taylor, E., Taylor, J. Z., 2004. Using qualitative psychology to investigate HACCP implementation barriers.
 International Journal of Environmental Health Research 14, 53-63.
- 1374 Ter Beek, A., Keijser, B.J.F., Boorsma, A., Zakrzewska, A., Orij, R., Smits, G.J., Brul, S., 2008. Transcriptome
- 1375 analysis of sorbic acid stressed *Bacillus subtilis* reveals a nutrient limitation response and indicates cell
- 1376 membrane remodeling Journal of Bacteriology 190, 1751-1761.
- Teunis, P.F., Ogden, I.D., Strachan, N.J., 2008. Hierarchical dose response of *E. coli* O157:H7 from human
 outbreaks incorporating heterogeneity in exposure. Epidemiology and Infection 136, 761-770.
- 1379 Thomas, M.K., Majowicz, S.E., Sockett, P.N., Fazil, A., Pollari, F., Dore, K., Flint, J.A., Edge, V.L., 2006.
- 1380 Estimated numbers of community cases of illness due to Salmonella, Campylobacter and verotoxigenic

- *Escherichia coli*, Pathogen-specific community rates. Infectious Diseases and Medical Microbiology 17,
 229-234.
- 1383 Tompkins, D.S., Hudson, M.J., Smith, H.R., Eglin, R.P., Wheeler, J.G., Brett, M.M., Owen, R.J., Brazier, J.S.,
- 1384 Cumberland, P., King, V., Cook, P.E., 1999. A study of infectious intestinal disease in England,
- 1385 microbiological findings in cases and controls. Communicable Diseases and Public Health 2, 108-113.
- 1386 Van Der Fels-Klerx, H.J., Kandhai, M.C., Booij, C.J.H., 2008. A conceptual model for identification of
- emerging risks, applied to mycotoxins in wheat-based supply chains. World Mycotoxin Journal 1, 13-23.
- 1388 Van Der Giessen, J., Fonville, M., Bouwknegt, M., Langelaar, M., Vollema, A., 2007. Seroprevalence of
- *Trichinella spiralis* and *Toxoplasma gondii* in pigs from different housing systems in The Netherlands. Vet
 Parasitol 148:371-374.
- 1391 Van Der Spiegel, M., Luning, P.A., Ziggers, G.W., Jongen, W.M., 2004. Evaluation of performance
- measurement instruments on their use for food quality systems. Critical Reviews in Food Science and
 Nutrition 44, 501-512.
- Van Dijk, H., Houghton, J., van Kleef, E., van der Lans, I., Rowe, G., Frewer, L.J., 2008. Consumer responses to
 communication about food risk management. Appetite 50, 340-352.
- 1396 Van Kleef, E., Houghton, J.R., Krystallis, A., Pfenning, U., Rowe, G., van Dijk, H., van de Lans, I.A., Frewer,
- L.J., 2007. Consumer evaluation of food risk management quality in Europe. Risk Analysis 27, 1565-1580.
- 1398 Van Kreijl, C.F., Knaap, A.G.A.C., Van Raaij, J.M.A., Busch, M.C.M., Havelaar, A.H., Kramers, P.G.N.,
- 1399 Kromhout, D., Van Leeuwen, F.X.R., Van Leent-Loenen, H.M.J.A., Ocké, M.C., Verkleij, H., 2006. Our
- food, our health healthy diet and safe food in the Netherlands. National Institute for Public Health and the
 Environment, Bilthoven.
- 1402 Van Pelt, W., Van De Giessen, A.W., Van Leeuwen, W.J., Wannet, W., Henken, A.M., Evers, E.G., De Wit,
- 1403 M.A.S., Van Duynhoven, Y.T.H.P., 1999. Oorsprong, omvang en kosten van humane salmonellose. Deel 1.
- 1404 Oorsprong van humane salmonellose met betrekking tot varken, rund, kip, ei en overige bronnen.
- 1405 Infectieziekten Bulletin 10, 240-243.
- Veening, J.W., Kuipers, O.P., Brul, S., Hellingwerf, K.J., Kort, R. 2006. Effects of phosphorelay perturbations
 on architecture, sporulation, and spore resistance in biofilms of *Bacillus subtilis*. Journal of Bacteriology
 1408 188, 3099-3109.

- 1409 Veening, J.W., Smits, W.K. and Kuipers, O.P. 2008. Biostability, epigenetics and bet-hedging in bacteria.
- 1410 Annual Reviews in Microbiology 62, 193-210.
- Verhoef, L. P.; Kroneman, A.; van Duynhoven, Y.; Boshuizen, H.; van Pelt, W., Koopmans, M., 2009. Selection
 tool for foodborne norovirus outbreaks. Emerging Infectious Diseases 15,31-38.
- 1413 Voetsch, A.C., Van Gilder, T.J., Angulo, F.J., Farley, M. M., Shallow, S., Marcus, R., Cieslak, P. R., Deneen,
- 1414 V.C., Tauxe, R. V., 2004. FoodNet estimate of the burden of illness caused by nontyphoidal Salmonella
- 1415 infections in the United States. Clinical Infectious Diseases 38, S127-S134.
- 1416 Wagner, M., Dahl, A., 2008. Direct molecular quantification of food-borne pathogens. In: Proceedings of the
- 1417 'Future challenges to microbial food safety' symposium, Wolfheze, The Netherlands.
- 1418 Wang, S., Duan, H., Zhang, W., Li, J.W., 2007. Analysis of bacterial foodborne disease outbreaks in China
- between 1994 and 2005. FEMS Immunology and Medical Microbiology 51, 8-13.
- 1420 Wattiau, P., Weijers, T., Andreoli, P., Schliker, C., Veken, H.V., Maas, H.M., Verbruggen, A.J., Heck, M.E.,
- 1421 Wannet, W.J., Imberechts, H., Vos, P., 2008. Evaluation of the Premi Test Salmonella, a commercial low
- density DNA microarray system intended for routine identification and typing of *Salmonella enterica*.
- 1423 International Journal of Food Microbiology 123, 293-298.
- 1424 Weenig, M.W.H., Midden, C.J.H., 1997. Mass-media information campaigns and knowledge-gaps effects.
- 1425Journal of Applied Social Psychology 27, 945-958.
- 1426 Wheeler, J.G., Sethi, D., Cowden, J.M., Wall, P.G., Rodrigues, L.C., Tompkins, D.S., Hudson, M.J., Roderick,
- 1427 P.J., 1999. Study of infectious intestinal disease in England, rates in the community, presenting to general
- 1428 practice, and reported to national surveillance. The Infectious Intestinal Disease Study Executive. British
- 1429 Medical Journal 318, 1046-1050.
- 1430 Wijnands, L.M., 2008. Bacillus cereus associated foodborne disease. Quantitative aspects of exposure
- assessment and hazard characterization. PhD thesis, Wageningen University.
- 1432 Wijnands, L.M., Dufrenne, J.B., Rombouts, F.M., In 't Veld, P.H., van Leusden, F.M., 2006. Prevalence of
- potentially pathogenic *Bacillus cereus* in food commodities in The Netherlands. Journal of Food Protection
 69, 2587-2594.
- 1435 Wilson, D.J., Gabriel, E., Leatherbarrow, A.J., Cheesbrough, J., Gee, S., Bolton, E., Fox, A., Fearnhead, P., Hart,
- 1436 C.A., Diggle, P.J., 2008. Tracing the source of campylobacteriosis. PLoS Genetics 4, e1000203.

Confidential manuscript submitted to International Journal of Food Microbiology

1438 Figure 1. Basic risk framework for infectious diseases (from Tait *et al.*, 2006)



Confidential manuscript submitted to International Journal of Food Microbiology

1441 **Table 1. Different aggregation levels for evaluation of food safety.**

MICROORGANISM-RELATED FACTORS	HUMAN-RELATED FACTORS
Ecosystems	Global systems
Food chains	Regions
Food products	Countries
Food products	Consumer
Populations of micro-organisms	Human populations
Individual cells of micro-organisms	Human individuals
Cellular and molecular processes	Cellular and molecular processes

s of ..

Confidential manuscript submitted to International Journal of Food Microbiology

Table 2. A systems approach to food safety.

Bold red font: Source increases risk to food safety;

	ct on risk to food safety unc ice reduces risk on food safe				
Green names fort. Sour		<i></i>	S		
DRIVERS	SOURCES	PATHWAYS Farm Processing/distribution Preparation/Consumption			OUTCOMES
	Pathogens				Public health
Economy			2		1
Globalization	Reduced geographical barriers to spread (of new variants)	Inadequate sanitation: higher pathogen loads Global sourcing Intensified contact structures	Long and complex supply chains Varying hygiene levels		Increased risk
Food price / income level Science and technology	, and industry	Less profit margins; decreased investment in food safety		Preference for cheaper alternatives (e.g. less meat and butter; discounters; home brands)	Risk not clear
Minimal processing	Adaptation		Less kill steps		Increased risk if not well controlled
Innovation	New food animal species	Step change food innovation	Smart labels	Risk not clear	
		Smart packaging			
		Bacteriophages			

DRIVERS	SOURCES	PATHWAYS			OUTCOMES
	Pathogens	Farm	Processing/distribution	Preparation/Consumption	Public health
Laboratory methods	Discovery of new pathogens or variants		Q		Increased observed risk
	Omics approaches		2		
Culture and demograph	hy	I	^o		
Population growth		Polluted environments	S	Increased demand	Increased risk
Migration			$\langle \rangle$	New food habits	
Age structure		×.	Increase in elderly	Increased	
		<pre></pre>		More premature babies	risk
Nature and environment	nt	Q,			
Climate change and	Changing ecology	Droughts, floods		Population displacement	Changing spatial patterns of risk
regional differences		Competition for land resources Movement of farms to new areas		Increased difficulties to maintain cold chain	
Water, waste and		Irrigation water	Water/energy savings		Increased risk
energy		quality Waste recycling	cleaning, process and ingredient water quality		
Evolution Emergence and transfer of virulence factors Antimicrobial resistance	transfer of virulence	New reservoirs	Increased survival	Increased infectivity	Increased risk
Population contact structures	Species jumps (spill- over from epizootics or exploitation of new	Contact zoonoses (MRSA, Q-fever)			Increased risk

DRIVERS	SOURCES	PATHWAYS			OUTCOMES
Pathogens	Pathogens	Farm	Processing/distribution	Preparation/Consumption	Public health
	agricultural areas)				
Consumer behavior				L	4
Food choice Food handling	Psychrotrophs Re-emerging pathogens	Exotic/ethnic foods Regional products	No or mild processing, less heat treatment Increased pre-processing and -packaging No acceptance of	Convenience foods Year round availability Healthy foods (fish, vegetables & fruits) Less fat/salt/sugar Eating outside home Storage: inadequate	Increased risk Increased
Technologies Attitudes/education			irradiation	time/temp control	risk
Information					
Surveillance	Identification of new pathogens Detection of unexpected events	No.	Effectiveness of current controls	Changes in consumption patterns: who, what, where, why?	Increase in observed risk
Education		Professional education		Hygiene campaigns	Reduced risk
			Attitude changes to accept safe technologies		
Government and poli	cies	1		1	1
Regulations		Standardisation	GHP/HACCP		Reduced risk
	Ban of antibiotic growth promoters	Fraudulent behavior			

	SOURCES		PATHWAYS		OUTCOMES
	Pathogens	Farm	Processing/distribution	Preparation/Consumption	Public health
Risk (-benefit) assessment		Targets for pathogen reduction	Risk-based targets		Reduced risk
Food defense		Agro/bioterrorism		Increased risk	
Agriculture			^o		
Animal friendly and organic production	Reduced AMR	Re-emergence (Trichinella, Toxoplasma)	22		Risk not clear
		Higher (<i>Campylobacter</i>) or <i>lower prevalence</i> (<i>Salmonella</i>)			
Aquaculture		More farmed fish			Risk not clear
Antimicrobial use	Resistance development	Increased therapeutic use			Increased risk
		A CY			