Investing in Food Safety for Developing Countries: Opportunities and Challenges in Applying Whole-Genome Sequencing for Food Safety Management

Isabella Apruzzese,¹ Eunyeong Song,² Ernest Bonah,³ Vernadette S. Sanidad,⁴ Pimlapas Leekitcharoenphon,⁵ Julius John Medardus,⁶ Nagmeldin Abdalla,⁷ Hedayat Hosseini,⁸ and Masami Takeuchi⁹

Abstract

Whole-genome sequencing (WGS) has become a significant tool in investigating foodborne disease outbreaks and some countries have incorporated WGS into national food control systems. However, WGS poses technical challenges that deter developing countries from incorporating it into their food safety management system. A rapid scoping review was conducted, followed by a focus group session, to understand the current situation regarding the use of WGS for foodborne disease surveillance and food monitoring at the global level and identify key limiting factors for developing countries in adopting WGS for their food control systems. The results showed that some developed nations routinely use WGS in their food surveillance systems resulting in more precise understanding of the causes of outbreaks. In developing nations, knowledge of WGS exists in the academic/research sectors; however, there is limited understanding at the government level regarding the usefulness of WGS for food safety regulatory activities. Thus, incorporation of WGS is extremely limited in most developing nations. While some countries lack the capacity to collect and analyze the data generated from WGS, the most significant technical gap in most developing countries is in data interpretation using bioinformatics. The gaps in knowledge and capacities between developed and developing nations regarding use of WGS likely introduce an inequality in international food trade, and thus, relevant international organizations, as well as the countries that are already proficient in the use of WGS, have significant roles in assisting developing nations to be able to fully benefit from the technology and its applications in food safety management.

Keywords: whole-genome sequencing, next-generation sequencing, foodborne disease surveillance, food safety, developing countries, Food and Agriculture Organization of the United Nations

Introduction

N EW FOODBORNE PATHOGEN analytical technologies often hold the promise of improving food safety, and wholegenome sequencing (WGS) is one of them (Deng *et al.*, 2016). This technology positively contributes to epidemiological investigations of foodborne outbreaks, to the identification of emerging health threats, to genome characterization of bacterial isolates, and to identify virulence, antimicrobial resistance (AMR), and other relevant genes in complex samples (Nadon *et al.*, 2017; Taboada *et al.*, 2017). As WGS evolves from a research tool to a practical food safety management instrument,

¹Franco Prattico Masters' Course in Science Communication, Trieste, Italy.

²Department of Chemistry, College of Chemistry and Chemical Engineering, Xiamen University, Fujian, China.

³Food and Drugs Authority, Northern Regional Office, Accra, Ghana.

⁴National Meat Inspection Service, Manila, Philippines.

⁵National Food Institute, Technical University of Denmark, Copenhagen, Denmark.

⁶Department of Veterinary Anatomy and Pathology, College of Veterinary Medicine and Biomedical Sciences, Sokoine University of Agriculture, Morogoro, Tanzania.

⁷Sudanese Standard and Metrology Organization, Khartoum, Sudan.

⁸National Nutrition & Food Technology Research Institute, Shahid Beheshti University of Medical Sciences, Teheran, Iran.

⁹Food and Agriculture Organization of the United Nations, Rome, Italy.

^{© 2019} Food and Agriculture Organization of the United Nations; Published by Mary Ann Liebert, Inc. This Open Access article is distributed under the terms of the Creative Commons License (http://creativecommons.org/licenses/by/4.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

The views expressed in this publication are those of the author(s) and do not necessarily reflect the views or policies of the Food and Agriculture Organization of the United Nations.

a number of bioinformatic software and tools have been developed to enable seamless analyses of the sequenced foodborne pathogen data (Langmead and Nellore, 2018). WGS is already a routine tool to identify and characterize pathogens in developed countries (Jackson *et al.*, 2016), and significant amounts of relevant data have been stored systematically and the results of use of such data can be found on various websites, such as GenomeTrakr (US FDA, 2019).

New technologies are often perceived as too advanced by developing countries; some have stated that WGS is too sophisticated and data generated from the technology can be overwhelming as capacity development activities and trainings on rather basic microbiological analyses are still their fundamental needs (FAO, 2016). However, many researchers state that WGS itself is straightforward to perform, and this knowledge needs to be properly communicated (Allard *et al.*, 2016).

With the world population reaching a number of 9.1 billion in 2050 (UN/DESA, 2017), agricultural trade is expected to continue expanding (FAO, 2015). The global food supply chain has become extremely complex and it is not uncommon that one food item's ingredients come from various suppliers in multiple countries (McCullough *et al.*, 2008). This means even a single local contamination could affect a large portion of the food supply chain, thus affecting a large number of people on a global scale (Gharehgozli *et al.*, 2017).

In addition to the already problematic informal transportation of food items through smuggling and alike, the everincreasing online food trade and direct e-food purchasing may create a sideline traffic of food items, which do not necessarily go through established border control systems (McCullough *et al.*, 2008). Consequently, it is increasingly necessary to strengthen food safety collaborations among countries regardless of their geographical location (Fukuda, 2015) and sharing data obtained through WGS has a potential in creating a functional global environment to enable such collaboration for effective foodborne disease outbreak management (Sasaki and Burr, 2000).

Incorporation of WGS into regulatory frameworks in developed countries has demonstrated its effectiveness in food safety management (Edmond-Rheault *et al.*, 2017), but whether WGS would be feasible and effective in developing countries needs to be assessed. A rapid scoping review, a keyword-based research synthesis that outlines the areas of research on a particular topic, was conducted followed by a focus group session to understand how realistically WGS can bring benefits to developing countries; how this technology can help improve the current situation; whether developing nations can meet essential requirements in terms of knowledge and infrastructures; and how introduction of WGS could affect international trade.

Materials and Methods

A rapid scoping review was conducted using a standard framework (Arksey and O'Malley, 2005). A set of hypotheses was developed with primary, secondary, and tertiary keywords (Table 1). A series of searches were conducted using the various combinations of keywords in two bibliographic databases: ScienceDirect* and PubMed.[†] An addi-

tional search was conducted online for general publications and information. All searches were limited to publications/documents written in English and published up to July 2018. Keyword combinations were expressed using Boolean positional operators to refine the results (Spink *et al.*, 2001). To maintain the focus on WGS applications regarding microbiological hazards, the results on other types of contamination such as chemical and physical hazards were not considered.

Publications and information made available by key international organizations, including the United Nations (UN) specialized agencies, Organization for Economic Cooperation and Development and World Organization for Animal Health, were reviewed. Further review was conducted on the websites of some food safety competent authorities, including the Canadian Food Inspection Agency; European Centre for Disease Prevention and Control; European Food Safety Authority; United States Centers for Disease Control and Prevention; United States Food and Drug Administration; Food Safety Authority of Ireland; Food Safety Commission of Japan; United Kingdom Food Standards Agency; Food Standards Australia New Zealand; and Ministry for Food and Drug Safety in the Republic of Korea.

In order for the study to highlight the opportunities and drawbacks in incorporating WGS in countries with challenges in terms of both financial resources and technical capacities, the UN country classifications (UN/DESA, 2018) were used and the terminologies of "developed countries" and "developing countries" were applied.

Accordingly, a separate set of searches was conducted to find information on situations in developing countries and the sources included the following: National Service of Agri-Food Health and Quality in Argentina; the Ministry of Health and the Ministry of Agriculture, Livestock and Food Supply in Brazil; Agency for Quality and Food Safety in Chile; China Food and Drug Administration; National Food Safety Coordination Committee in Kenya; Ministry of Agriculture and Ministry of Health in Malaysia; Food and Drug Administration in Myanmar; Department of Agriculture in South Africa; and Department of Health and Department of Trade and Industry in South Africa.

To cover the possibility that the scoping review had an information gap regarding developing countries, a focus group session was conducted. Specific objectives of the focus group session were as follows: to confirm one of the key results of the scoping review that developing countries have not generally used WGS for food safety management; to assess the availability and accessibility of global WGS data; to collect feedback on specific challenges for developing countries in generating, storing, and sharing such data; to assess the capacity level for prerequisite activities necessary for use of WGS, such as isolating pathogens; and to identify the capacity needs in terms of infrastructure, knowledge, and technical capacities such as bioinformatics. Due to the geographical separation of participants, the focus group was conducted online.

Experts from Ghana, Iran, the Philippines, Sudan, Tanzania, and Thailand (n=6) participated in the focus group session. All are members of the FAO informal network of developing countries to share information, knowledge, and experience in WGS (FAO, 2018). All confirmed their agreement to be recorded during the focus group session. The focus group

^{*}http://www.sciencedirect.com

[†]http://www.ncbi.nlm.nih.gov/pubmed

No.	Primary keyword	Secondary keyword	Tertiary keyword	<i>ScienceDirect</i> ^a	PubMed ^b
1	WGS	Outbreak		671	292
			Investigation	546	105
			Identification	486	59
			Detection	468	59
		Incident	Detection	678	5
		meldent	Investigation	544	5
			Identification	277	2
			Detection	212	2
		Casa	Detection	207	202
		Case	• .• .•	11,530	202
			Investigation	86/3	38
			Identification	3639	34
			Detection	3275	29
	NGS	Outbreak		833	89
			Investigation	633	19
			Identification	567	26
			Detection	614	25
		Incident	Detection	830	23
		meldent	Investigation	500	1
				257	1
			Identification	337	1
		~	Detection	406	1
		Case		17,554	815
			Investigation	11,478	37
			Identification	7455	145
			Detection	7777	146
2	Developing countries ^c	Microbial safety		21.019	97
			NGS	225	
			WGS	174	
		Outbrook	W05	60.840	2627
		Outbreak	NCC	221	2027
			INUS INUS	331	1
			WGS	3/5	10.10
		Food safety		99,188	1040
			NGS	433	1
			WGS	342	1
3	WGS	Data share		2990	10
			Trace	741	
			Routine surveillance	250	
			Transmission	761	1
	NGS	Data share	Tunshinssion	5608	27
	1105	Data share	Trace	1080	27
			Douting ourseillenge	1080	
				93	2
			Transmission	1194	3
	Genotyping	Data share	-	/2,162	132
			Trace	10,255	I
			Routine surveillance	3670	1
			Transmission	19,277	19
4	Developing country	WGS		2337	8
			Source attribution	89	
			Interpretation	850	1
		NGS	1	2582	25
		1.00	Source attribution	66	
			Interpretation	917	3
5	Trada	WGS	interpretation	1356	1
5	ITade	1105	Trace	250	4
			Irace	350	1
			Reference	962	1
			Frontline tool	4	0
			Phylogenetic	88	0
		NGS		1277	27
			Trace	289	
			Reference	888	1
			Frontline tool	12	Ō
			Phylogenetic	184	1
			1 11/10/2010/10	104	1

TABLE 1. SCOPING REVIEW KEYWORDS AND NUMBERS OF RELEVANT HITS

(continued)

No.	Primary keyword	Secondary keyword	Tertiary keyword	<i>ScienceDirect</i> ^a	PubMed ^t
6	WGS	Prediction		3215	79
			Management	1128	6
			Risk assessment	621	1
		Forecast		609	1
			Management	366	0
			Risk assessment	197	0
7	WGS	Developing country		2337	8
		1 6 9	Retrospective investigation	170	0
			Microbiology investigation	305	0
			Epidemiology investigation	287	0

 TABLE 1. (CONTINUED)

A rapid scoping review was conducted to outline the key thematic areas on the topic of WGS applications in food safety management. As the relevance was verified in the process, the higher number of hits indicates that the topic has been widely discussed in published literature and the lower number of hits indicates that the supporting evidence to the topic is not sufficiently available in published literature. ^aScienceDirect. Available at: https://www.sciencedirect.com

^bPubMed—NCBI. Available at: https://www.ncbi.nlm.nih.gov/pubmed

^cBased on the UN country classification and terminology. Available at: https://www.un.org/development/desa/dpad/wp-content/uploads/ sites/45/publication/WESP2018_Full_Web-1.pdf

NGS, next-generation sequencing; WGS, whole-genome sequencing.

questions were developed following a standardized pattern (Krueger, 2002). The recording was transcribed and analyzed with five stages: familiarization with the data, identification of thematic ideas, indexing of data, charting of quotes, and mapping and interpretation of data (Krueger and Casey, 2000).

Results

The rapid scoping review successfully outlined key thematic areas of the topic, and as anticipated, the relatively low number of hits was compensated with the search strategy of repeating the combinations of synonyms of each keyword (Table 1). The scoping review identified that, compared with the previously confirmed number of countries (n=4) that used WGS for food safety management in 2016 (FAO, 2016), more countries (n > 10) are using WGS for regulatory purposes. However, the review also confirmed that all those additional countries are developed nations. There were no data that any developing country had initiated using WGS in the government system and this was further confirmed by the focus group session with experts from developing countries.

The search combining keywords with the technology, "case" and "investigation" resulted in a high number of hits, indicating that many articles report the positive impacts that implementation of WGS has had in surveillance systems. Most of these reports describe how introduction of WGS has increased the number of detected outbreaks and surveillance (den Bakker *et al.*, 2014; Dallman *et al.*, 2015; Kwong *et al.*, 2015; FAO, 2016; Jackson *et al.*, 2016). Nonexperts may wrongly perceive that introduction of WGS is creating more food safety problems rather than solving them, but experts state that the technology brings major public health benefits by minimizing the scale of the outbreaks and preventing recurrence of the problem from the same source (Kwong *et al.*, 2015).

In general, the scoping review generated a very low number of hits for any searches with the keyword "developing countries," indicating that this type of information may not be usually reported in peer-reviewed journals. There is some evidence that WGS is being introduced at the research level in various developing countries and many scientists have good understanding of its potential (Healy *et al.*, 2016). Some research institutes, such as Institut Pasteur, have conducted WGS-related studies in developing countries to promote the technology (Weill *et al.*, 2019).

During the focus group session, the most consistent findings were that scientific knowledge, understanding of potential applications and benefits, and strong motivation to use WGS exist, but they expressed an immediate need to increase the level of understanding by high-level government officials. Therefore, it can be concluded that a systematic incorporation of WGS into the government systems in developing nations has not yet occurred, and thus, the technology still is not used as an official food safety management tool (Pekdemir, 2018). Understanding both the reality and the perceptions from experts in such developing countries could significantly contribute to the formulation of a global strategy on WGS and food safety (Pekdemir, 2018). Table 2 presents the summary of the focus group results.

The scoping review also had a relatively low number of hits when searching about the food safety situation in developing countries, which implies that developing countries are currently facing significant challenges in detecting foodborne outbreaks and/or in reporting outbreaks in publishable articles (Ahmed *et al.*, 2015; Odeyemi, 2016). Assuming that it is already costly for developing countries to fully implement traditional methodologies to detect foodborne outbreaks, it is necessary to question whether incorporating WGS would be cost-effective in improving outbreak detection/reporting situations.

Conventional methodologies to detect foodborne pathogens often require laboratories to obtain an official accreditation for species-specific identification and typing protocols. The universality of WGS has a benefit in efficiency and the cost-per-sample has been significantly decreasing. Although WGS could contribute to cost savings for identification of foodborne pathogens, the overall cost may still be high as WGS requires relevant infrastructures and functioning equipment/personnel. In addition, the cost-per-run of sequencing is still perceived as costly in many countries (FAO, 2016). The issue of WGS cost being both a benefit and a

		TABLE 2. SUMMARY OF THE FOCUS GROUP RESULTS
No.	Topic	Responses
	Use of WGS for official food safety management activities	 WGS is not being used for food safety management in any countries that participants are from.^a Priority conflict exists as many countries focus on food security. Awareness of the potential benefit of WGS in the area of food safety is low.@
7	WGS data collection and accessibility to the global database	 No budget to consider the new technology to be employed in the area of 1000 safety. No national database of WGS for food safety exists in any countries participants are from. All were aware of the global databases that they can access, but the access from the government sector is extremely limited.
\mathfrak{c}	WGS data generation	 Two countries have genomic center and relevant research institute that can host WGS data for food safety, but currently no accessible data are available. Three participants are involved in WGS data generation and the data are shared on some global databases. There is a recognition that WGS data generation from countries such as theirs would be useful in strengthening their surveillance systems.
4	Food safety data sharing	 A multiplication of a state of the state of the
S	Basic laboratory capacity	from their countries with international community is rare. • Participants confirmed that their countries have reference laboratories that can isolate pathogens, thus basic microbiological capacity exists.
9	Food and environmental monitoring for foodborne pathogens	 The work is not currently aimed at application of WGS. All participants confirmed that clinical samples are often checked on a regular basis, but food and environmental monitoring for foodborne pathogens is usually done on an <i>ad hoc</i> basis, when some incidents are suspected.
٢	Technical challenges to improve food monitoring systems	 Budgetary constraints exist to regularly conduct such acuvities. As food safety is a cross-cutting issue, communication and collaboration with different sectors (agriculture, health, trade and commerce) are a challenge. General awareness on the importance of microbiological food safety issues is low.
8	Bioinformatic capacity	 Financial and number resources are number. Capacities exist at the research level, but all participants confirmed that the capacity is not sufficient at the government level.
6	Needs in improving WGS knowledge and capacity	 Some countries confirmed that bioinformatics is not in the curricula of major universities, and thus, the number of bioinformaticians in their countries is very low. There is a strong need in raising the awareness of the potential of WGS for food safety in developing countries. The focus for knowledge and capacity improvement should be in the government and regulatory sectors. A long-term vision is necessary to initiate the awareness camaaisn to follow the technology development and trend in
10	Potential of WGS in developing countries	 food safety analysis. Participants strongly agreed that WGS would increase the number of detected food safety incidences and outbreaks. If the number of outbreaks is properly reported, food safety will become a priority for their countries, and thus, the technology would contribute to protecting public health.
$A_{dev(a)}^{a}$	focus group session was conducted with technica eloping countries. Based on a standard pattern of an mapped interpretations of quotes. Jhana, Iran, the Philippines, Sudan, Tanzania, and MR, antimicrobial resistance; INFOSAN, Internati	experts ($n = 6$) from developing countries to assess the current situation regarding WGS applications in food safety management for alyzing a focus group study, the items in the left column represent thematic ideas and the items in the right column represent the indexed Thailand. Thailand.

drawback for developing countries has been also documented in detail in a previous study (FAO, 2016).

Another challenge that developing countries face when considering use of WGS for foodborne disease surveillance is regarding data handling and sharing. WGS generates a large amount of data, characterized by a high volume of information, high-speed production of data, and a wide variety of extremely precise information—all of which are useful for risk managers to strengthen their understanding of the complexity and diversity of a food safety problem (Bergholz *et al.*, 2014). However, significant data mining efforts are necessary to attribute the source of transmission of a foodborne pathogen and provide insights on control strategies (Langmead and Nellore, 2018), and such expertise in bioinformatics is not readily available in developing nations (Pongor and Landsman, 1999).

A number of user-friendly cloud-based tools are now available that can be used by people without formal bioin-formatic education. Use of these tools may alleviate availability of bioinformaticians as a critical need for developing countries (NCBI, 2018) and (Fierro *et al.*, 2018).

WGS-generated data can be especially important for international food trade (Zankari *et al.*, 2013). While only a limited amount of information on the status of WGS incorporation in developing countries was available in literature, many industrialized nations have already incorporated this technology in their food safety regulatory framework. The imbalance in use of WGS may pose compliance difficulties to the developing countries who often export food products to developed countries (Franz *et al.*, 2014).

Application of WGS would reduce the need of multiple tests at the borders (Yoshida, 2014) and its accuracy, together with its timely results, is likely to make a positive impact on international food trade in the future (Zankari *et al.*, 2013). However, if this technological advancement is only happening in developed countries, it may create inequality in trade. Many developed countries have been incorporating new approaches and new technologies to upgrade their national food control systems[‡] in the last decade (Gilchrist *et al.*, 2015), and as a consequence, a possibly large gap between developed and developing nations might have been induced. Such a gap indirectly imposes the pressure to some exporting developing countries in needing to increase their technical capacities to the ones of the importing developed countries (Veggeland and Borgen, 2005).

A number of articles explore the predictive power of WGS data (Gordon *et al.*, 2014), demonstrating that its use in outbreak investigations can support decision makers and therefore can reduce the misidentification of food sources (Parkhill *et al.*, 2000). However, the scoping review revealed that this advantage has not been discussed much in developing nations. Also, since there were almost no bioinformatic-related studies found in the scoping review on WGS and food safety in developing countries, it can be concluded that there

may not be a sufficient number of bioinformaticians available in developing countries. To sustain effective national food control systems where WGS data are regularly used, many bioinformatic-related challenges must be addressed in developing countries (Fricke and Rasko, 2014).

Results of the focus group session provided remarkable information that these experts from developing countries are confident about WGS and that scientific capacity to develop the technology exists. All participants demonstrated awareness about the potential of the technology, some of them have had extensive experience with it, and two of the countries represented by those interviewed currently operate genomic research centers.

The main issue they are facing is connected to priority conflicts that result in national governments not dedicating financial/human resources for the introduction of WGS. All focus group participants stated that they are more than ready to use the technology to contribute to their national food control systems, but lack of commitment at higher levels of government has been a barrier to their use of WGS. To fill the gap, knowledge, and capacity, collaboration between developed and developing countries and technical assistance from international organizations are needed. However, factors such as the current movement toward proactive global sharing of AMR data, awareness of experts about WGS, and existence of university degree programs in bioinformatics in some developing countries, together with the high level of interest and strong motivation expressed by focus group participants, suggest developing countries are getting ready to introduce WGS.

Discussion

WGS is a powerful tool to identify and characterize foodborne pathogens, and if routinely used, it can prevent and control outbreaks (Joensen *et al.*, 2014). Currently more than 10 countries use the technology for food safety management, making WGS an essential tool to more completely understand food microbiology (Allard *et al.*, 2018). As of July 2018, between 11,000 and 18,000 scientific articles discuss the use of genomic technologies for identification, investigation, and/or prevention of foodborne disease outbreaks. In fact, one of the major advantages of WGS is the speedy subtyping of foodborne bacterial pathogens worldwide (Nadon *et al.*, 2017).

A few case studies from the United States, Denmark, and England have shown how WGS can be incorporated into the food safety regulatory system for outbreak investigations with benefits, such as specificity, allowing improved case definition to enhance the outbreak management; sensitivity, enabling linkage of apparently sporadic diseases occurring under the outbreak surveillance radar; and precision, determining the root cause of complex outbreaks (FAO, 2016).

The major success of WGS is that it enables sharing of high-resolution data of the genomes, giving the possibility to identify genetic variants, and to explore the effects of gene expression and regulation (Gilad *et al.*, 2009). In addition, the large quantity of high-quality data produced in a short amount of time makes this technique efficient as well as accurate (Daetwyler *et al.*, 2014). Latest sequencers are equipped with a function to combine many steps, including template preparation, sequencing, imaging, genome alignment, and assembling the sequenced data, making the

[‡]Codex Alimentarius Commission provides principles and guidelines for national food control systems (CAC/GL 82-2013. Available at: http://www.fao.org/input/download/standards/13358/ CXG_082e.pdf) and the terminology of "national food control systems" has been widely used among food safety competent authorities worldwide to indicate the overall food-related systems to protect the health of consumers and ensure fair practices in the food trade.

laboratory procedures drastically simple and fast (Metzker, 2009).

Hence, WGS has positively contributed to timely and effective foodborne outbreak investigations in those countries that utilize the technology, confirming that it will be an excellent "next-generation" tool that will play a significant role in the area of food safety (Wang *et al.*, 2016). However, many developing countries may not agree that it is efficient and inexpensive in their circumstance. Countries without established surveillance systems may not see the cost/benefit of adding WGS capability and implementation of WGS may divert essential resources from more pressing priorities (FAO, 2016).

Underreporting of foodborne diseases and microbial contamination cases is common in both developed and developing countries; however, the extremely low number of outbreaks reported by developing nations likely does not reflect the actual situation in these countries, especially if compared with the large numbers of outbreaks reported on a regular basis by many developed countries (Fierro *et al.*, 2018) as well as with the estimated global burden of foodborne diseases (WHO, 2015). A major reason for this extreme underreporting in developing countries is often linked to limited capacities in their national food control systems with insufficient resources and infrastructures (Grace, 2015).

The availability of diagnostic services to report infections is also key, because without it countries have to rely only on syndromic surveillance, without sufficient sensitivity and specificity. Proper implementation of internationally harmonized measures and regulations (WHO, 2005) might help developing countries address this problem. Before the introduction of WGS, it is essential for countries to have a systematic mechanism to collect isolates and their metadata from both clinical samples and food/environment samples (EFSA, 2008; FAO, 2016). Unless these matters are addressed, introduction of WGS will remain a tool used in research projects only.

While the focus group participants declared that wellequipped in-country laboratories are available for analysis of isolates, WGS analysis needs to be conducted routinely if it is used as the basis for food surveillance systems. Therefore, assessing the capacity in developing countries to confirm the feasibility in establishing WGS-based surveillance systems for both food supply and clinical infections is important, taking into consideration the current isolate and data collection mechanisms (FAO, 2016). However, a question remains whether the unavailability of a complete surveillance system should stop integration of WGS in developing countries-and the focus group participants suggested that the opportunity is to be presented equally to anyone to advance knowledge. Also, Helmy et al. (2016) discussed that, in their study in developing countries, health risk identification, diagnoses, treatment, and prevention would likely be improved with new tools and technologies.

Improvement of pathogen detection capacity could lead countries to have a food safety monitoring situation that more accurately reflects the reality of their situation to undertake food safety surveillance activities. For example, the introduction of WGS in Kenya drew attention of decision makers to the overall importance of food safety and thus provided the basis for the development of a national food control system (FAO, 2016). The Kenya Medical Research Institute introduced the use of WGS to sequence strains from selected pathogens from clinical samples. With the detailed data obtained through WGS, the Kenyan Government was able to map disease hotspots to revise existing treating regimens and identify high-risk foods. As a result, interest in investing more on food safety was increased and the usefulness of WGS was recognized for analysis of microbial food contamination for regulatory interventions (FAO, 2016).

WGS is innovative because it does not require targeting primers (Burall *et al.*, 2017), but it allows molecular subtyping (Bal *et al.*, 2016) and further state-of-the-art analyses using sophisticated approaches (Kase *et al.*, 2017). Moreover, WGS has shown to provide advantages, compared with traditional methods, because of its suitability to identify AMR and to detect the emergence of new foodborne pathogen strains (Baker *et al.*, 2017). AMR is already a global concern (Paterson, 2006) and WGS can accurately identify resistant genotypes and predict resistant phenotypes (Tyson *et al.*, 2015). Resistance levels have increased in pathogens that are more common in developing countries, and this trend is increasing (Okeke *et al.*, 2005).

Many authorities in developing nations have shown interest in addressing AMR, because their countries show increasing numbers of cases of AMR-associated illnesses (WHO, 2001). When developing countries utilize WGS for microbiological risk assessment related to AMR, it can also benefit food safety management as food is one of the major vehicles of AMR (FAO/WHO, 2018). The topic of AMR was spontaneously raised by the focus group participants who agreed that the accuracy of WGS, in particular, the application of single-nucleotide polymorphism-based AMR identification, could play a key role in addressing the problem.

While WGS holds technological advantages, the data have to be interpreted, stored, and shared to make a comparison with data from other sources (Bergholz *et al.*, 2014). Results obtained from sequencing will only be useful for interpretation after they are put into bioinformatic databases (Franz *et al.*, 2014). Therefore, not having access to relevant comparable global data makes the outputs obtained through WGS almost useless. Such restriction is partially due to IT-related issues that involve high-capacity computers (Karunaratne *et al.*, 2018) or fast, reliable internet connections (Karunaratne *et al.*, 2018), which constitute a serious problem in developing countries and without which even easy-to-use, online bioinformatic tools (NCBI, 2018) cannot be used.

The availability of high-speed wireless internet connections such as 5G is expected to address this issue in most developing countries, but the full benefit of this may be realized several more years later. Moreover, lack of international standards regarding the quality of the sequenced data represents another important issue that needs to be solved. In addition, if there is no previously sequenced genome on which to perform the bioinformatic alignment, the identification process through WGS is neither simple nor fast (Yang *et al.*, 2017).

A number of countries have established WGS data-sharing mechanisms and there are some key databases and platforms developed for routine use. These include the US FDA GenomeTrakr (Allard *et al.*, 2016), which is a subset database of the Sequence Read Archive at the National Center for Biotechnology Information (NCBI), the European Nucleotide Archive, and the deoxyribonucleic acid (DNA) Data Bank of Japan (FAO, 2016). Although accessibility to the real-time global databases is highly desirable, microbial evolution

makes it hard to obtain up-to-date, complete information, since only a small fraction of microbes will ever be cultured and sequenced (Selifonova *et al.*, 2001). Even more challenging is the reluctance of the countries to share their data in real time.

Some initiatives seek to facilitate global data sharing: a scientific consortium, the global microbial identifier (GMI) envisions a global system of DNA genome databases for microbial and infectious disease identification and diagnostics (GMI, 2018); an international laboratory network, the PulseNet International works toward the standardized use of WGS methods and data to identify and subtype foodborne bacterial pathogens worldwide (Nadon *et al.*, 2017); and GenomeTrakr provides an open source WGS database for microbial pathogens collected and publically shared by agencies in real time (US FDA, 2018). These initiatives are supported by technical experts from many countries and Taboada *et al.* (2017) stated that more effort needs to be made to develop or unify a global tool to make relevant WGS data as the one-of-a-kind interoperable resource.

None of the countries involved in the focus group session has its own national WGS database. Some foodborne diseaserelated data exist, but the data are currently being obtained through traditional techniques; they are often not systematically collected and are only occasionally published in scientific journals. Scientists in developing countries use the abovementioned global databases to publish their data, but the reports do not constitute a national data set, because there is no system to consolidate them. Further work needs to be done in order for such countries to systematically use WGS as an integrated part of their national food control systems.

International market trends demonstrate that there are many opportunities for developing countries in export of food products (Henson and Jaffee, 2006). These markets are driven by competition based on safety and quality of products, and therefore, the associated characteristics are codified in standards (Ponte and Gibbon, 2005). Such standards can act both as a barrier and a catalyst to upgrade food management capacities, because they can facilitate or constrain the access of developing nations to high-value markets for agricultural and food products (Henson and Jaffee, 2006). Use of WGS data to assay for foodborne pathogens creates trade opportunities for developing countries and can be advantageous for them to meet the quality assurance criteria requested by industrialized countries (Henson and Jaffee, 2006). Moreover, WGS data could be used as a supportive evidence for decision makers in case of food trade disputes.

To mitigate the food safety challenges associated with food trade, a global collaborative effort is required (Buzby, 2003). For example, an economical estimation study on possible harmonization for a standard on aflatoxin B1 in nuts and cereals showed a potential increase of world exports by USD 38.8 billion for nuts and USD 6.1 billion for cereals compared with total exports when using divergent national standards (Wilson and Otsuki, 2001). This study further revealed that in addition to the gain in safety of the products, a healthy global competition would likely promote fair practices in global food trade. Technical assistance to developing countries provided by neutral international organizations would help them effectively access international markets in a proactive way (Henson and Jaffee, 2006).

The potential economic benefit of using WGS for detection of contaminated sources is worthwhile to discuss further with developing countries. Food safety problems often result in large-scale financial losses due to product recalls, disposals, and penalties, along with potential damage to the overall reputation of food producers, companies, and countries, with the resulting decrease in consumers' confidence (Hussain and Dawson, 2013). In theory, use of WGS with its precision could contribute in avoiding unnecessary food recalls due to possible source-attribution mistakes and even eventually contribute in reducing food waste. The specificity and sensitivity of WGS could provide more targeted approaches to regulatory authorities (FAO, 2016), and thus, WGS can become a good tool to reduce the economic burden for developing countries.

Developing countries may need feasibility studies to assess their priorities, their requirements, and their readiness to incorporate WGS in food safety management (FAO, 2016). In particular, WGS may not be the most effective tool in the situation where food safety policies or national food control systems do not function well. Technical capacities to manage the technology and relevant tools of WGS will eventually be needed, but to assure implementation of WGS, government officials who have an understanding of how WGS works are a key factor (Healy et al., 2016). Developed nations need to pay attention to the emerging needs of developing countries because the challenges the latter has to face may have been imposed by the richer nations (Henson and Jaffee, 2006). Major benefits for all could arise from assuring equal opportunities and having globally comparable WGS systems (Sansone et al., 2012).

Conclusion

Results from the scoping review and the focus group session indicated that in developing countries, WGS is rarely used in nonresearch settings. Governmental authorities do not always see food safety as a priority, and thus, they do not invest in tools such as WGS to test foods for contamination or to investigate foodborne diseases.

The gap in capacity between developing and developed countries when applying WGS may create inequalities in ensuring fair food trade. In developing nations, infrastructure and technical capacities, including laboratory conditions and bioinformatic capacities, still constitute a challenge to realistically apply WGS (Ahmed *et al.*, 2015). This can be of a global concern because the core nature of WGS largely lies in data sharing to have a real-time access to global data to match with the relevant local data sets (Kaye *et al.*, 2009) and partial collection of data from limited geographical areas will not reach the maximum results that WGS could provide.

Collecting and globally sharing such data in a comparable format benefit all to have a better food safety situation from the angles of both public health and global food trade (Pekdemir, 2018). In order for all countries to benefit from WGS, the work of various global initiatives aiming at building knowledge around WGS-related capacities, such as GMI, GenomeTrakr, and PulseNet, will be essential.

Timely scientific advice provided by international organizations to facilitate knowledge transfer and discussions on global data management is important to ensure the technology will benefit all. Technical support to developing nations is the immediate need to be provided by international organizations, experienced countries, and global technical alliances

WGS FOR FOOD SAFETY IN DEVELOPING COUNTRIES

with capacity development activities focusing on practical aspects of WGS application in food safety management.

Acknowledgments

At the time of article preparation, I.A. and E.S. were interns and Takeuchi designed and led the overall research. The views expressed in this publication are those of the author(s) and do not necessarily reflect the views or policies of the Food and Agriculture Organization of the UN. Song collected and organized the information/data, I.A. prepared the first draft, and E.B., V.S.S., P.L., J.J.M., N.A., and H.H. participated in the focus group session. A written consent from the focus group members was obtained to follow the relevant rules and procedures for the ethical clearance review. Content and editorial review provided by Virginia Val Hillers, a former Extension Specialist and Professor at Washington State University, is gratefully acknowledged.

Disclosure Statement

No competing financial interests exist.

Disclaimer

The views expressed in this publication are those of the author(s) and do not necessarily reflect the views or policies of the Food and Agriculture Organization of the United Nations.

References

- Ahmed SS, Alp E, Ulu-Kilic A, Doganay M. Establishing molecular microbiology facilities in developing countries. J Infect Public Health 2015;8:513–525.
- Allard MW, Bell R, Ferreira CM, Gonzalez-Escalona N, Hoffmann M, Muruvanda T, Ottesen A, Ramachandran P, Reed E, Sharma S, Stevens E, Timme R, Zheng J, Brown EW. Genomics of foodborne pathogens for microbial food safety. Curr Opin Biotechnol 2018;49:224–229.
- Allard MW, Strain E, Melka D, Bunning K, Musser SM, Brown EW, Timme R. Practical value of food pathogen traceability through building a whole-genome sequencing network and database. J Clin Microbiol 2016;54:1975–1983.
- Arksey H, O'Malley L. Scoping studies: Towards a methodological framework. Int J Soc Res Methodol 2005;8:19–32.
- Baker KS, Campos J, Pichel M, Della Gaspera A, Duarte-Martínez F, Campos-Chacón E, Bolaños-Acuña HM, Guzmán-Verri C, Mather AE, Diaz Velasco S, Zamudio Rojas ML, Forbester JL, Connor TR, Keddy KH, Smith AM, López de Delgado EA, Angiolillo G, Cuaical N, Fernández J, Aguayo C, Morales Aguilar M, Valenzuela C, Morales Medrano AJ, Sirok A, Weiler Gustafson N, Diaz Guevara PL, Montaño LA, Perez E, Thomson NR. Whole genome sequencing of *Shigella sonnei* through PulseNet Latin America and Caribbean: Advancing global surveillance of foodborne illnesses. Clin Microbiol Infect 2017;23:845–853.
- Bal AM, Coombs GW, Holden MTG, Lindsay JA, Nimmo GR, Tattevin P, Skov RL. Genomic insights into the emergence and spread of international clones of healthcare-, community-, and livestock-associated methicillin-resistant *Staphylococcus aureus*: Blurring of the traditional definitions. J Glob Antimicrob Resist 2016;6:95–101.

- Burall LS, Chen Y, Macarisin D, Pouillot R, Strain E, De Jesus AJ, Laasri A, Wang H, Ali L, Tatavarthy A, Zhang G, Hu L, Day J, Kang J, Sahu S, Srinivasan D, Klontz K, Parish M, Evans PS, Brown EW, Hammack TS, Zink DL, Datta AR. Enumeration and characterization of *Listeria monocytogenes* in novelty ice cream samples manufactured on a specific production line linked to a listeriosis outbreak. Food Control 2017;82:1–7.
- Buzby JC. International Trade and Food Safety: Economic Theory and Case Studies, (Agricultural Economic Report No. 828), Electronic Report from the Economic Research Service, United States Department of Agriculture, 2003.
- Daetwyler HD, Capitan A, Pausch H, Stothard P, van Binsbergen R, Brøndum RF, Liao X, Djari A, Rodriguez SC, Grohs C, Esquerré D, Bouchez O, Rossignol MN, Klopp C, Rocha D, Fritz S, Eggen A, Bowman PJ, Coote D, Chamberlain AJ, Anderson C, VanTassell CP, Hulsegge I, Goddard ME, Guldbrandtsen B, Lund MS, Veerkamp RF, Boichard DA, Fries R, Hayes BJ. Whole-genome sequencing of 234 bulls facilitates mapping of monogenic and complex traits in cattle. Nat Genet 2014;46:858–865.
- Dallman TJ, Byrne L, Ashton PM, Cowley LA, Perry NT, Adak G, Petrovska L, Ellis RJ, Elson R, Underwood A, Green J, Hanage WP, Jenkins C, Grant K, Wain J. Wholegenome sequencing for national surveillance of Shiga toxinproducing *Escherichia coli* O157. Clin Infect Dis 2015;61: 305–312.
- den Bakker HC, Allard MW, Bopp D, Brown EW, Fontana J, Iqbal Z, Kinney A, Limberger R, Musser KA, Shudt M, Strain E, Wiedmann M, Wolfgang WJ. Rapid whole-genome sequencing for surveillance of *Salmonella enterica* Serovar Enteritidis. Emerg Infect Dis 2014;20:1306–1314.
- Deng X, den Bakker HC, Hendriksen RS. Genomic epidemiology: Whole-genome-sequencing-powered surveillance and outbreak investigation of foodborne bacterial pathogens. Annu Rev Food Sci Technol 2016;7:353–374.
- Edmond-Rheault JG, Jeukens J, Freschi L, Kukavica-Ibrulj I, Boyle B, Dupont MJ, Colavecchio A, Barrere V, Cadieux B, Arya G, Bekal S, Berry C, Burnett E, Cavestri C, Chapin TK, Crouse A, Daigle F, Danyluk MD, Delaquis P, Dewar K, Doualla-Bell F, Fliss I, Fong K, Fournier E, Franz E, Garduno R, Gill A, Gruenheid S, Harris L, Huang CB, Huang H, Johnson R, Joly Y, Kerhoas M, Kong N, Lapointe G, Larivière L, Loignon S, Malo D, Moineau S, Mottawea W, Mukhopadhyay K, Nadon C, Nash J, Ngueng Feze I, Ogunremi D, Perets A, Pilar AV, Reimer AR, Robertson J, Rohde J, Sanderson KE, Song L, Stephan R, Tamber S, Thomassin P, Tremblay D, Usongo V, Vincent C, Wang S, Weadge JT, Wiedmann M, Wijnands L, Wilson ED, Wittum T, Yoshida C, Youfsi K, Zhu L, Weimer BC, Goodridge L, Levesque RC. A Syst-OMICS Approach to ensuring food safety and reducing the economic burden of Salmonellosis. Front Microbiol 2017;2:996.
- [EFSA] European Food Safety Authority Working Group on Developing Harmonised Schemes for Monitoring Antimicrobial Resistance in Zoonotic Agents. Harmonised monitoring of antimicrobial resistance in *Salmonella* and *Campylobacter* isolates from food animals in the European Union. Clin Microbiol Infect 2008;14:522–533.
- Fierro RG, Thomas-Lopez D, Deserio D, Liebana E, Rizzi V, Guerra B. Outcome of EC/EFSA questionnaire (2016) on use

of Whole Genome Sequencing (WGS) for food- and waterborne pathogens isolated from animals, food, feed and related environmental samples in EU/EFTA countries. Eur Food Saf Author Support Publ 2018;15:1–44.

- [FAO] Food and Agriculture Organization of the United Nations. The State of Agricultural Commodity Markets Trade and Food Security: Achieving a Better Balance Between National Priorities and the Collective Good. Rome: FAO, 2015.
- [FAO] Food and Agriculture Organization of the United Nations. Applications of WGS in Food Safety Management. Rome: FAO, 2016.
- [FAO/WHO] Food and Agriculture Organization of the United Nations, World Health Organization. FAO/WHO Expert Meeting on Foodborne Antimicrobial Resistance: Role of Environment, Crops and Biocides. Rome, June 11–15, 2018.
- [FAO] Food and Agriculture Organization of the United Nations Food Safety and Quality, Whole Genome Sequencing (WGS) and Food Safety 2018. Available at: http://www.fao .org/food/food-safety-quality/a-z-index/wgs, accessed June 2, 2019.
- Franz E, Delaquis P, Morabito S, Beutin L, Gobius K, Rasko DA, Bono J, French N, Osek J, Lindstedt BA, Muniesa M, Manning S, LeJeune J, Callaway T, Beatson S, Eppinger M, Dallman T, Forbes KJ, Aarts H, Pearl DL, Gannon VP, Laing CR, Strachan NJ. Exploiting the explosion of information associated with whole genome sequencing to tackle Shiga toxin-producing *Escherichia coli* (STEC) in global food production systems. Int J Food Microbiol 2014;187:57–72.
- Fricke WF, Rasko DA. Bacterial genome sequencing in the clinic: Bioinformatic challenges and solutions. Nat Rev Genet 2014;15:49–55.
- Fukuda K. Food safety in a globalized world. Bull World Health Organ 2015;93:212.
- Gharehgozli A, Iakovou E, Chang Y, Swaney R. Trends in global E-food supply chain and implications for transport: Literature review and research directions. Res Transport Busin Manage 2017;25:2–14.
- Gilad Y, Pritchard JK, Thornton K. Characterizing natural variation using next-generation sequencing technologies. Trends Genet 2009;25:463–471.
- Gilchrist CA, Turner SD, Riley MF, Petri WA Jr, Hewlett EL. Whole-genome sequencing in outbreak analysis. Clin Microbiol Rev 2015;28:541–563.
- [GMI] Global Microbial Identifier. *Global Microbial Identifier* 2018. Available at: http://www.globalmicrobialidentifier.org, accessed June 2, 2019.
- Gordon NC, Price JR, Cole K, Everitt R, Morgan M, Finney J, Kearns AM, Pichon B, Young B, Wilson DJ, Llewelyn MJ, Paul J, Peto TE, Crook DW, Walker AS, Golubchik T. Prediction of *Staphylococcus aureus* antimicrobial resistance by whole-genome sequencing. J Clin Microbiol 2014;52:1182– 1191.
- Grace D. Food safety in low and middle income countries. Int J Environ Res Public Health 2015;12:10490–10507.
- Healy MJ, Tong W, Ostroff S, Eichler HG, Patak A, Neuspiel M, Deluyker H, Slikker W. Regulatory bioinformatics for food and drug safety. Regul Toxicol Pharmacol 2016;80:342–347.
- Helmy M, Awad M, Mosab KA. Limited resources of genome sequencing in developing countries: Challenges and solutions. Appl Transl Genom 2016;9:15–19.
- Henson S, Jaffee S. Food safety standards and trade: Enhancing competitiveness and avoiding exclusion of developing countries. EJDR 2006;18:593–621.

- Hussain MA, Dawson CO. Economic impact of food safety outbreaks on food businesses. Foods 2013;2:585–589.
- Jackson BR, Tarr C, Strain E, Jackson KA, Conrad A, Carleton H, Katz LS, Stroika S, Gould LH, Mody RK, Silk BJ, Beal J, Chen Y, Timme R, Doyle M, Fields A, Wise M, Tillman G, Defibaugh-Chavez S, Kucerova Z1, Sabol A, Roache K, Trees E, Simmons M, Wasilenko J, Kubota K, Pouseele H, Klimke W, Besser J, Brown E, Allard M, Gerner-Smidt P. Implementation of nationwide real-time whole-genome sequencing to enhance listeriosis outbreak detection and investigation. Clin Infect Dis 2016;63:380–386.
- Joensen KG, Scheutz F, Lund O, Hasman H, Kaas RS, Nielsen EM, Aarestrup FM. Evaluation of real-time WGS for routine typing, surveillance and outbreak detection of verotoxigenic *Escherichia coli*. J Clin Microbiol 2014;52:1501–1510.
- Karunaratne T, Peiris C, Hansson H. Implementing small scale ICT projects in developing countries—How challenging is it?. IJEDICT 2018;14:118–140.
- Kase JA, Zhang G, Chen Y. Recent foodborne outbreaks in the United States linked to atypical vehicles—Lessons learned. Curr Opin Food Sci 2017;18:56–63.
- Kaye J, Heeney C, Hawkins N, de Vries J, Boddington P. Data sharing in genomics—Re-shaping scientific practice. Nat Rev Genet 2009;10:331–335.
- Krueger RA. Focus Group Interviewing: Designing and Conducting Focus Group Interviews. Saint Paul, MN, 2002. Available at: https://www.eiu.edu/ihec/Krueger-FocusGroup Interviews.pdf, accessed June 2, 2019.
- Krueger RA, Casey MA. Focus Groups: A Practical Guide for Applied Research (third edition). Thousands Oaks, CA: Sage, 2000.
- Kwong JC, McCallum N, Sintchenko V, Howden BP. Whole genome sequencing in clinical and public health microbiology. Pathology 2015;47:199–210.
- Langmead B, Nellore A. Cloud computing for genomic data analysis and collaboration. Nat Rev Genet 2018;19:208–219.
- McCullough EB, Pingali PL, Stamoulis KG. The Transformation of Agri-Food Systems: Globalization, Supply Chains and Smallholder Farmers. Sterling, VA: The Food and Agriculture Organization and Earthscan, 2008.
- Metzker ML. Sequencing technologies—The next generation. Nat Rev Genet 2009;31–46.
- Nadon C, Van Walle I, Gerner-Smidt P, Campos J, Chinen I, Concepcion-Acevedo J, Gilpin B, Smith AM, Man Kam K, Perez E, Trees E, Kubota K, Takkinen J, Nielsen EM, Carleton H, FWD-NEXT Expert Panel. PulseNet International: Vision for the implementation of whole genome sequencing (WGS) for global food-borne disease surveillance. Euro Surveill 2017;22:1–12.
- [NCBI] National Center for Biotechnology Information, NCBI Home, 2018. Available at: https://www.ncbi.nlm.nih.gov/, accessed June 2, 2019.
- Odeyemi OA. Public health implications of microbial food safety and foodborne diseases in developing countries. Food Nutr Res 2016;60:29819.
- Okeke IN, Laxminarayan R, Bhutta ZA, Duse AG, Jenkins P, O'Brien TF, Pablos-Mendez A, Klugman KP. Antimicrobial resistance in developing countries. Part I: Recent trends and current status. Lancet Infect Dis 2005;5:481–493.
- Parkhill J, Wren BW, Mungall K, Ketley JM, Churcher C, Basham D, Chillingworth T, Davies RM, Feltwell T, Holroyd S, Jagels K, Karlyshev AV, Moule S, Pallen MJ, Penn CW, Quail MA, Rajandream MA, Rutherford KM, van Vliet AH, Whitehead S, Barrell BG. The genome sequence of the food-

WGS FOR FOOD SAFETY IN DEVELOPING COUNTRIES

borne pathogen *Campylobacter jejuni* reveals hypervariable sequences. Nature 2000;403:665–668.

- Paterson DL. Resistance in Gram-negative bacteria: Enterobacteriaceae. Am J Med 2006;119:S20–S28.
- Pekdemir C. On the regulatory potential of regional organic standards: Towards harmonization, equivalence, and trade? Glob Environ Change 2018;50:289–302.
- Pongor S, Landsman D. Bioinformatics and the developing world. Biotechnol Dev Monit 1999;40:10–13.
- Ponte S, Gibbon P. Quality standards, conventions and the governance of global value chains. Econ Soc 2005;37:1–31.
- Sansone SA, Rocca-Serra P, Field D, Maguire E, Taylor C, Hofmann O, Fang H, Neumann S, Tong W, Amaral-Zettler L, Begley K, Booth T, Bougueleret L, Burns G, Chapman B, Clark T, Coleman LA, Copeland J, Das S, de Daruvar A, de Matos P, Dix I, Edmunds S, Evelo CT, Forster MJ, Gaudet P, Gilbert J, Goble C, Griffin JL, Jacob D, Kleinjans J, Harland L, Haug K, Hermjakob H, Ho Sui SJ, Laederach A, Liang S, Marshall S, McGrath A, Merrill E, Reilly D, Roux M, Shamu CE, Shang CA, Steinbeck C, Trefethen A, Williams-Jones B, Wolstencroft K, Xenarios I, Hide W. Toward interoperable bioscience data. Nat Genet 2012;44:121–126.
- Sasaki T, Burr B. International Rice Genome Sequencing Project: The effort to completely sequence the rice genome. Curr Opin Plant Biol 2000;3:138–142.
- Selifonova O, Valle F, Schellenberger V. Rapid evolution of novel traits in microorganisms. Appl Environ Microbiol 2001;67:3645–3649.
- Spink A, Wolfram D, Major Jansen BJ, Saracevic T. Searching the web: The public and their queries. J Am Soc Inf Sci Technol 2001;52:226–234.
- Taboada EN, Graham MR, Carriço JA, Van Domselaar G. food safety in the age of next generation sequencing, bioinformatics, and open data access. Front Microbiol 2017; 8:909.
- Tyson GH, McDermott PF, Li C, Chen Y, Tadesse DA, Mukherjee S, Bodeis-Jones S, Kabera C, Gaines SA, Loneragan GH, Edrington TS, Torrence M, Harhay DM, Zhao S. WGS accurately predicts antimicrobial resistance in *Escherichia coli*. J Antimicrob Chemother 2015;70: 2763–2769.
- [UN/DESA] United Nations Department of Economic and Social Affairs Population Division. World Population Prospects: The 2017 Revision. New York: United Nations, 2017.
- UN/DESA. World Economic Situation and Prospects 2018. New York: United Nations, 2018.
- [US FDA] United States Food and Drug Administration. Whole Genome Sequencing (WGS) Program. 2018. Available at: https://www.fda.gov/Food/FoodScienceResearch/ WholeGenomeSequencingProgramWGS/default.htm, accessed June 2, 2019.
- US FDA. GenomeTrakr fast facts. 2019. Available at: https:// www.fda.gov/Food/FoodScienceResearch/WholeGenome SequencingProgramWGS/ucm403550.htm

- Veggeland F, Borgen SO. Negotiating International Food Standards: The World Trade Organization's Impact on the Codex Alimentarius Commission. Governance 2005;18:675– 708.
- Wang S, Weller D, Falardeau J, Strawn LK, Mardones FO, Adell AD, Moreno AI. Food safety trends: From globalization of whole genome sequencing to application of new tools to prevent foodborne diseases. Trends Food Sci Technol 2016;57A:1–47.
- Weill FX, Domman D, Njamkepo E, Almesbahi AA, Naji M, Nasher SS, Rakesh A, Assiri AM, Sharma NC, Kariuki S, Pourshafie MR, Rauzier J, Abubakar A, Carter JY, Wamala JF, Seguin C Bouchier C, Malliavin T, Bakhshi B, Abulmaali HHN, Kumar D, Njoroge SM, Malik MR, Kiiru J, Luquero FJ, Azman AS, Ramamurthy T, Thomson NR, Quilici ML. Genomic insights into the 2016–2017 cholera epidemic in Yemen. Nature 2019;565:230–233.
- [WHO] World Health Organization. International Health Regulations. 3rd ed. 2005. Available at: https://www.who.int/ihr/ publications/9789241580496, accessed June 2, 2019.
- WHO. WHO Estimates of the Global Burden of Foodborne Diseases: Foodborne Disease Burden Epidemiology Reference Group 2007–2015. Geneva, Switzerland: World Health Organization, 2015.
- WHO. Interventions and Strategies to Improve the Use of Antimicrobials in Developing Countries: A Review. Geneva, Switzerland: World Health Organization, 2001.
- Wilson J, Otsuki T. Global Trade and Food Safety: Winners and Losers in a Fragmented System—Policy Research Working Papers. Washington, DC: The World Bank, 2001.
- Yang Z, Mammel M, Papafragkou E, Hida K, Elkins CA, Kulka M. Application of next generation sequencing toward sensitive detection of enteric viruses isolated from celery samples as an example of produce. Int J Food Microbiol 2017;261: 73–81.
- Yoshida M. Global harmonization of food safety regulations: Perspectives from Japan after the Fukushima nuclear accident. J Sci Food Agric 2014;94:1938–1940.
- Zankari E, Hasman H, Kaas RS, Seyfarth AM, Agersø Y, Lund O, Larsen MV, Aarestrup FM. Genotyping using whole-genome sequencing is a realistic alternative to surveillance based on phenotypic antimicrobial susceptibility testing. J Antimicrob Chemother 2013;68:771– 777.

Address correspondence to: Masami Takeuchi, PhD Food and Agriculture Organization of the United Nations (FAO) Viale delle Terme di Caracalla 00153 Rome Italy

E-mail: masami.takeuchi@fao.org