



### Presence and Persistence of Salmonella in Water: The Impact on Microbial Quality of Water and Food Safety

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Salmonella ranks high among the pathogens causing foodborne disease outbreaks. According to the Centers for Disease Control and Prevention, Salmonella contributed to about 53.4% of all foodborne disease outbreaks from 2006 to 2017, and approximately 32.7% of these foodborne Salmonella outbreaks were associated with consumption of produce. Trace-back investigations have suggested that irrigation water may be a source of Salmonella contamination of produce and a vehicle for transmission. Presence and persistence of Salmonella have been reported in surface waters such as rivers, lakes, and ponds, while ground water in general offers better microbial quality for irrigation. To date, culture methods are still the gold standard for detection, isolation and identification of Salmonella in foods and water. In addition to culture, other methods for the detection of Salmonella in water include most probable number, immunoassay, and PCR. The U.S. Food and Drug Administration (FDA) issued the Produce Safety Rule (PSR) in January 2013 based on the Food Safety Modernization Act (FSMA), which calls for more efforts toward enhancing and improving approaches for the prevention of foodborne outbreaks. In the PSR, agricultural water is defined as water used for in a way that is intended to, or likely to, contact covered produce, such as spray, wash, or irrigation. In summary, Salmonella is frequently present in surface water, an important source of water for irrigation. An increasing evidence indicates irrigation water as a source (or a vehicle) for transmission of Salmonella. This pathogen can survive in aquatic environments by a number of mechanisms, including entry into the viable but nonculturable (VBNC) state and/or residing within free-living protozoa. As such, assurance of microbial quality of irrigation water is critical to curtail the produce-related foodborne outbreaks and thus enhance the food safety. In this review, we will discuss the presence and persistence of Salmonella in water and the mechanisms Salmonella uses to persist in the aquatic environment, particularly irrigation water, to better understand the impact on the microbial quality of water and food safety due to the presence of Salmonella in the water environment.

Keywords: irrigation water, agricultural water, Salmonella, produce safety, prevalence, pathogen detection, foodborne outbreaks

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### INTRODUCTION

Salmonella is a natural inhabitant in the gastrointestinal tract of many animals, including birds, reptiles, livestock, and humans (1–7). Salmonellosis caused by nontyphoidal Salmonella ranks among the highest in all gastroenteritis cases linked to food consumption, affecting the health of approximately one million people annually in the United States alone (8, 9), resulting in medical costs of \$3.7 billion. It is estimated that Salmonella species causes 93.8 million cases of gastroenteritis worldwide annually with 155,000 deaths (5-95th percentile, 39,000-303,000) (10). The causative source for salmonellosis has traditionally been attributed to animal origin (11, 12), including meat, eggs, and other poultry products, which has attracted considerable regulatory attention and enormous mitigation efforts (2, 9). In recent years, the number of foodborne outbreaks due to nontraditional sources of the pathogen such as domestic or imported fresh fruits, vegetables, spices, and nuts has been increasing (9, 13-15). A recent report from the Centers for Disease Control and Prevention (CDC) suggested nearly half (46%) of foodborne illnesses and 23% of deaths were associated with produce consumption (16). For instance, during July 2015 to February 2016, a Salmonella outbreak associated with consumption of imported Mexican cucumbers caused illness in at least 907 people, with six deaths in 40 states within the United States (17).

Consumption of more fresh fruits and vegetables has been advocated as a healthier diet habit because raw, or less processed, fruits and vegetables are good sources of vitamins, fiber, and other beneficial nutrients (18, 19). Since the early 1970s, the demand for fresh produce in the US has been on the rise continuously, and the estimated increase of per capita consumption of fresh fruits and vegetables from 1982 to 1997 reached approximately 32% (13). The consumption of vegetables grew more rapidly than fruits from 1976 to 2009 (20). A recent investigation by the US Department of Agriculture also suggested Americans consumed more fresh produce in 2015 than in 1970 (21) As a result, it is expected that more produce-related outbreaks of disease will occur even if the contamination sources and rates stay at the present levels. This has attracted the attention of the legislative branch and food safety regulatory agencies in the United States. The Food Safety Modernization Act (FSMA), which was enacted in 2011, emphasizes the significance of produce safety. In response to that, the US Food and Drug Administration (FDA) issued the Produce Safety Rule (PSR) in January 2013 to establish science-based standards for the growing, harvesting, packing, and holding by domestic and foreign farms of produce consumed in the United States (22).

The causative pathogens for produce contaminations include viruses, bacteria, and parasites. Contamination may occur at various stages during the production process (2). Salmonella has been regarded as the primary pathogen for causing produce-related foodborne outbreaks. Pathogenic bacteria carried by produce that can lead to an outbreak may involve multiple external sources and production stages. For instance, at the pre-harvest stage, Salmonella can come from specific agricultural practices such as using animal manure as fertilizer (1) and others

may include using contaminated water for irrigation, pesticide spraying, or anti-frost spraying (2, 23, 24).

Water has been shown as a source of microbial contamination of fresh produce and a vehicle for pathogen transmission (23, 25). FSMA defines "agricultural water" partially as water that is "intended to, or is likely to contact covered produce or food contact surfaces" (22). Irrigation water consists of a major component of agricultural water. The sources of irrigation water can come from ground water, surface water, or municipal water. Irrigation water can be applied to produce through various ways including drip tape, furrows, and overhead sprinklers (26). In this review, we will primarily focus on the presence, survival, persistence, and source of *Salmonella* in surface water, particularly irrigation water, to help us better understand the impact on the microbial quality of water and food safety due to the presence of *Salmonella* in the water environment.

## PRODUCE AS A VEHICLE FOR TRANSMISSION OF SALMONELLA

Foodborne outbreaks associated with fresh produce in the United States have been on the rise in the last few decades; and Salmonella has been recognized as the primary causative pathogen (1, 11, 13, 27). According to a report from the CDC, 31 Salmonella outbreaks from 2002 to 2003 were associated with fresh produce; while 29 were poultry related (2, 28). Also, Salmonella contributes to 53.4% (55/103) of the foodborne disease outbreaks documented by CDC among all pathogens investigated from 2006 to 2017 (Figure 1A); 32.7% (18/55) of the multistate foodborne Salmonella outbreaks were associated with produce (Figure 1B); and 60% (18/60) of the fresh producerelated outbreaks were attributed to Salmonella among all the pathogens involved (Figure 1C) (29). Many common types of produce have been implicated in Salmonella-related outbreaks, including beans, alfalfa sprouts, tomatoes, hot peppers, lettuce, cucumbers, cantaloupes, water melons, papayas, and mangoes (1, 2, 29-32).

Salmonella is originally from, and has adapted to, the microenvironment of the gastrointestinal tract of animals. Once released from the animal gastrointestinal tract or other sources with feces or exudates, Salmonella can be carried to surface waters through rainfall and surface runoffs, survive many challenges such as ultraviolet (UV) radiation from sunlight, poor nutrients, the changes in pH, and temperature (33-35). After attaching to produce, Salmonella must persist under these adverse environmental conditions at a sufficient concentration to cause human illness (13). The surface of various fruits and vegetables provides a niche for numerous bacterial species, such as epiphytes, plant pathogens, and opportunistic human pathogens, such as Salmonella (36). Interestingly, some plant pathogenic bacteria, such as Pectobacterium carotovorum, can promote the growth of Salmonella by macerating plant tissue and providing nutrients (9, 37), whereas other bacterial species such as Panebacillus spp. on tomatoes and cilantro may inhibit or even kill Salmonella (38, 39). Thus Salmonella has to survive

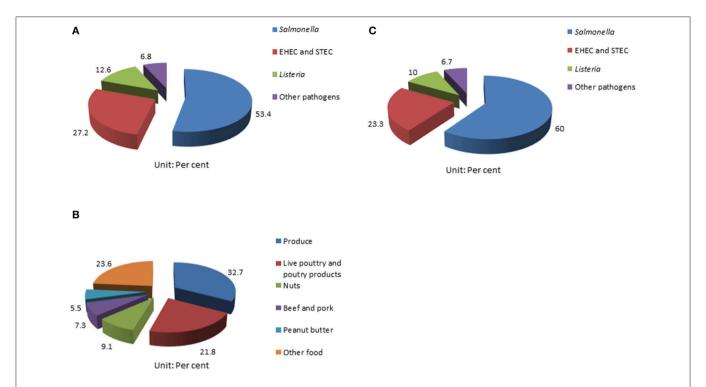


FIGURE 1 | Summary of surveillance of foodborne pathogen outbreaks from CDC (29). (A) Multistate foodborne outbreaks caused by Salmonella rank highest in all the pathogens surveilled; (B) the percentage of multistate Salmonella outbreaks linked to produce from 2006 to 2017; and (C) and Salmonella contributes to 60% of all produce-related outbreaks caused by different pathogens surveilled.

the inhibition from these microbes and outcompete them for the acquisition of nutrients and space (33).

The mechanisms for how Salmonella are carried in irrigation water or from other sources interact with plants, survive or persist on these intermediate hosts have become an interesting topic in recent years. Studies indicate Salmonella can be internalized into tomato plants through different routes and may possibly colonize and employ the plant as an alternative host (2, 9, 36). The formation of biofilm-like structures on the surface of roots, the colonizing regions of emerging lateral roots and wounded tissues has been reported (40, 41). Moreover, Salmonella inoculated into soil or blossoms can be recovered from the fruit at low internal levels, suggesting its ability to colonize and internalize tomato plants, and however, survival ability with different serovars varies in the soil and in different parts of the plants (42). Salmonella was also recovered from tomato fruits after it was introduced into the plant by stem injection or by flower inoculation (43). Stine et al. demonstrated that Salmonella was still detectable for at least 14 days after inoculation, indicating the bacteria probably developed a mechanism to adapt and survive within this hostile environment (44).

Recent studies have further suggested *Salmonella* may not only passively use plants as a "shelter" for survival, but a growing body of evidence has indicated it also may have evolved mechanisms for active adhesion and escape of plant immunity systems and actively internalize and proliferate in some plants, such as the tomato (9, 42, 45–47). It was found that *Salmonella* 

populations inoculated onto the exterior of tomatoes can increase in numbers at suitable humidity and temperature and such bacterial growth is serovar dependent (48). This may explain why some serovars, such as Newport, Montevideo, and Saintpaul, are more frequently linked to *Salmonella* outbreaks (2). A latest study suggested that the mechanisms *Salmonella* requires to colonize tomato plants are similar to phytopathogens, such as biosynthesis of amino acids, lipopolysaccharides (LPS), and nucleotides, indicating the flexibility of this pathogen to fit different hosts (49). The studies on *Salmonella* entry into inner leaves indicated that while trichomes are postulated as preferential colonization sites (50, 51), stomata are shown to be entry points *Salmonella* utilize for penetration of lettuce leaves (52, 53). The interaction of *Salmonella* with *Arabdopsis*, potato, sprouts and other plants are also reported and reviewed by Schikora et al. (47).

Preservation of taste, nutrients, and other desirable qualities of produce demand minimum processing, including avoidance of heating and sanitation with disinfectant to leafy vegetables and sprouts (2, 54), if washing is inadequate, the contamination acquired in field or production stage and postharvest easily transit from field to the table. Washing with chlorinated water, water-dip disinfection procedures may be applied to fruits such as melons, mango, and papaya. But if the disinfection efficiency of chlorinated water is not adequately monitored, the contaminated *Salmonella* can still be attached; and the rough surfaces for some types of melon like Cantaloupes can increase the adherence of bacteria and compromise the effect of washing (15, 30, 31,

55). Nowadays, the precut ready-to-eat vegetables and fruits are widely available to consumers in grocery stores; and poor hygiene and sanitary practice in preparation of these foods can bring additional chances for bacterial contamination (15, 56, 57). Therefore, the elimination of *Salmonella* contamination of fresh produce from fields to forks is almost impossible because of the above-mentioned loopholes. In general, the frequent occurrence of *Salmonella* outbreaks associated with produce and the scarcity of approaches available for sanitizing contaminated produce underscore the urgency for development of more efficient preventive control measures that can be applied at an earlier stage in the produce production.

# IRRIGATION WATER AS A SOURCE OR A VEHICLE FOR TRANSMISSION OF SALMONELLA

The original source of Salmonella on produce may come from soil, manure, irrigation water, and contact with reptiles, birds, or other small animals (23, 24, 58). Irrigation water has drawn considerable attention in recent years, and studies have implicated irrigation water as a source of Salmonella contamination (23, 24, 59). This hypothesis has been previously reviewed (5, 23, 24, 27, 60, 61) and appears to be supported by trace-back investigations of produce-related Salmonella outbreaks (62, 63). The 2005 multistate Salmonella outbreak involving tomatoes was caused by Newport serotype, and a traceback investigation linked this strain to an outbreak associated with tomatoes that occurred 3 years earlier. Furthermore, the isolates from these two outbreaks were shown to be genotypically identical by pulsed-field gel electrophoresis (PFGE), and the same bacterial strains were found present in a pond the farm used for irrigation (62). A case control study of a 2008 multistate outbreak identified Salmonella Saintpaul on serrano and jalapeño peppers and was also present in irrigation ponds (63). Another study indicated that the PFGE patterns of some of the isolates from irrigation ponds of produce farms in southern Georgia were indistinguishable from strains that were associated with the Salmonella Thompson outbreaks in 2010, 2012, and 2013, Salmonella Enteritidis outbreaks in 2011 and 2013, and a Salmonella Javiana outbreak in 2012 (61). The investigation of papaya Salmonella outbreak happened from 2006 to 2007 in Australia also found that the river water used for washing papayas was contaminated with Salmonella (64).

# PREVALENCE AND SOURCE OF SALMONELLA IN IRRIGATION WATER

The fresh produce consumption over the last 40 years and number of foodborne disease outbreaks associated with fresh produce has been increasing (15, 18). Irrigation is a critical factor for the production of fruits and vegetables, *Salmonella* present in irrigation waters has been regarded as one of the major source for fresh produce contamination (2, 23, 24), and this has become a public health concern and drawn more attention of food safety regulatory agencies. Irrigation is a critical factor

for the production of fruits and vegetables. As identified above, the source of irrigation water can include groundwater from wells, surface water (rivers and irrigation ponds), and treated wastewater (24, 27, 65, 66). Groundwater in wells is naturally filtered by soil and generally has a higher microbial quality (i.e., less microbes present). But it may be compromised by inferior construction or insufficient depth of the well, and may be contaminated from nearby latrines, septic tanks leaching fields, land application of waste water, and rainfalls (27, 67). Incidences of *Salmonella* contamination of ground water is mainly a concern of developing countries, especially in the rural areas, due to poor hygienic conditions, deficiently-structured water supply systems, and inadequate disinfection treatment (68), but occasionally occurs in developed countries as well (69, 70).

Surface waters, which include ponds, lakes, rivers, and streams, account for nearly half of the water used for irrigation in the United States (26). They are more exposed to environmental events such as discharge of sewage, rainfall, animal husbandry, and wildlife, and thus are more susceptible to contamination as compared to groundwater (23, 67). Rivers have been widely used as an irrigation source for agricultural practice (71, 72); river water, however, has been shown to be one of the largest reservoirs of viable Salmonella (2, 73). An estimation of Salmonella loads from a coastal Mediterranean river of a 16-month period (74) indicated Salmonella occurrence up to 95% during high waterflow (21% of the year) such as storm events. The subsequent study discovered S. Typhimurium accounted for 33.1% of all isolates recovered from the river in that period (73). In another study (12), Salmonella were detected from 57 of 72 (79.2%) water samples monthly collected from six stations of Little River in upper Suwannee Bain of Southern Georgia State of the United States. The recovery of Salmonella from rivers exhibited seasonality pattern, with summer time being highest, which is similar to the prevalence of Salmonella in irrigation ponds (6, 61). This increased prevalence of Salmonella during summer time may be related to multiple environmental factors, such as host shedding, enhanced persistence of Salmonella in warm temperature, and increase of storm events (12). In the Suwannee River watershed of southern Georgia, open surface pond waters are the main source of irrigation water. The case rates in this region have been observed to be 1.5-fold higher than the national average (6, 12). Presence of Salmonella in irrigation ponds within this region has been surveyed by Li et al. and showed that Salmonella was recovered from 50 of 170 (29.4%) water samples collected monthly over 27 months from 10 selected irrigation ponds that serve as water sources for irrigation of vegetables (61). Furthermore, more than half of the isolates were identified as S. Newport and antimicrobial susceptibility testing confirmed 16 S. Newport isolates were multidrug resistant (MDR), exhibiting resistance to ampicillin, chloramphenicol, streptomycin, sulfamethoxazole, and tetracycline (ACSSuT) and to the 1st, 2nd, and 3rd generations of cephalosporins (cephalothin, amoxicillin-clavulanic acid ceftriaxone) (61). Another monthly survey by Luo et al. on 10 ponds in the same area isolated Salmonella from 28.2% of all samples (n = 635)and the most common serotypes were Hadar, Montevideo, and Newport. In addition, 98.9% of the strains were reported to

be resistant to streptomycin and about 20% were MDR strains (6). The contamination of fresh produce by these nontyphi *Salmonella* species present in irrigation waters could generate detrimental clinical and public health consequence because the increasing antibiotic resistance limits options of treatment after microbial diagnosis. In foodborne disease outbreak settings, the infection of MDR strains may bring excess of mortality and morbidity (75, 76). Moreover, the encodings genes of MDR may be horizontally transferred to other pathogenic bacteria resulting the spreading of antibiotic resistance (75).

The microbial quality of treated municipal water depends on the efficiency of treatments to remove pathogenic enteric bacteria, viruses, and parasites. Prevalence of *Salmonella* in treated effluent has been reported (77). In the United States, only a limited scale of treated municipal water is used for irrigation of crops because of the concern of potential bacterial contamination (24). However, in countries or regions with a shortage of fresh water, wastewater treated to a suitable level can be used as a substitute for ground water or surface water for irrigation purposes, and the microbial quality guidelines adopted for treated wastewater must be the strictest (23).

It is generally accepted that Salmonella present in water can be traced back to its animal origins. This pathogen may directly be transported from feces or exudates of wild animals by rain water runoff to rivers or ponds used for irrigation (12). Manure of domesticated animals has long been used to fertilize soil because it is economical and beneficial to the environment. However, studies have indicated that Salmonella in manure can survive as long as 231 days and may eventually contaminate produce by rain water splashing and/or by surface irrigation water (78). Sewage effluents contain waste water from human toilets with a high concentration of bacteria and other pathogens (79). The average concentration of Salmonella can reach as high as  $2.7 \times 10^2$  CFU/100 ml (80, 81), which could become a major source of contamination if discharged directly or with inadequate treatment. Salmonella and other bacteria in sewage water can be effectively reduced to very low levels with modern treatment methods, but it is not practical to eliminate all the bacteria. When discharged, this will pose another contamination source of Salmonella to surface waters (79, 82) as illustrated in the proposed model of Salmonella transmission in the environment (5).

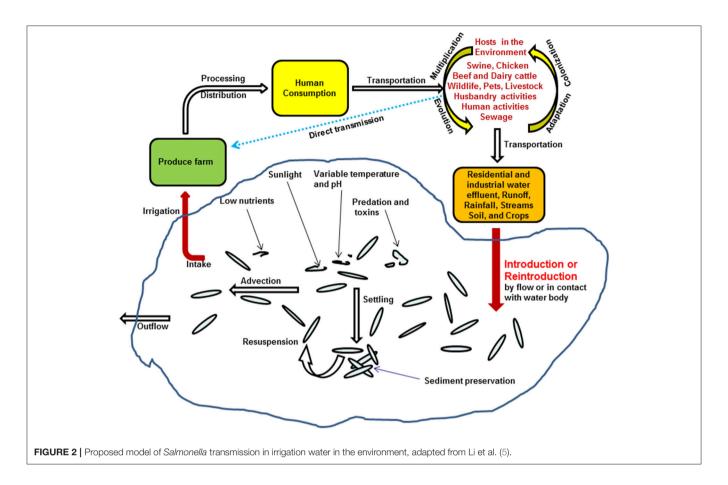
# SALMONELLA SURVIVAL AND PERSISTENCE IN IRRIGATION WATER

The gastrointestinal tract of vertebrates is generally regarded as the natural habitat for *Salmonella* (83), and bacteria released from feces could be transported into aquatic systems by sewage discharges, rainfall events or associated surface runoffs (84). *Salmonella* may survive in an environment with a broad range of pH (4.05–9.5) and can multiply in a broad range of temperatures (7–48°C) (34). In a closed environment at room temperature (25°C), it has been shown that *Salmonella* can survive for up to 5 years in sterile water or a phosphate buffered solution (85). However, the eco-environment of irrigation water, such as river or pond water, is harsher and more complicated

and dynamic. Survival and persistence of Salmonella in water depends on multiple environmental factors, such as temperature, pH, salt, dissolved oxygen concentration, nutrient availability, interaction with other microorganisms, and exposure to UV light radiation (23, 27, 86, 87). As a result, Salmonella viability will decrease in water over time and could survive generally less than 30 days (23). However, many studies now show that biofilms can facilitate the survival of Salmonella in water and invertebrates such as free-living protozoa and various vertebrate animals can serve as reservoirs for Salmonella and other pathogenic bacteria. For example, Gaertner et al. detected Salmonella from water biofilms and crayfish samples from the headwater spring of Spring Lake, Texas and found that Salmonella isolates from biofilms collected 23 days apart shared the same Rep-PCR profile, suggesting Salmonella infrequently washed into an aquatic system might take up water biofilms as dwellings for long term persistence (88). Sha et al. further demonstrated Salmonella isolated from biofilms on a concrete surface exhibited significant microheterogeneity, but remained pathogenic (89). Additionally, Salmonella were recovered from sediments in water, and the concentration was higher than in the overlying water. Similar phenomena have also been observed in survival of other bacteria in water. This might be caused by sedimentation and absorption (90-92), which may help explain why continuous prevalence of Salmonella over a long period of time was found in river and surface ponds in multiple studies (5, 12).

Besides the elongation of *Salmonella* viability and survivability in water environment, recently, Li et al. also suggested that frequent detection of *Salmonella* in irrigation water might be due to numerous reintroduction events associated with several different hosts in the environment. This was supported by genomic microarray analysis on *Salmonella* Newport isolates (5). On the one hand, produces growing in the field can be directly contaminated by *Salmonella* excreted from proximate environmental hosts, such as swine, chicken, beef, dairy cattle, husbandry or human, other wild lives, or inadequately composited manures applied in the fields (4, 5, 9); on the other hand, these bacteria could also be repeatedly introduced to the irrigation ponds through storm events and rainfalls, survive in water, and reach fresh produce through irrigation, and circulate back to human and other animals (**Figure 2**) (5).

Furthermore, the prevalence and distribution of *Salmonella* in rivers, streams, or ponds exhibit seasonality, which has been documented in multiple studies (12, 61, 93). Thus, running water formed by rainfall may act as an important vehicle for *Salmonella* transportation. In addition, a number of studies suggest that bacteria can enter a viable but nonculturable (VBNC) state under stress conditions (94), such as low temperature (95), osmotic stress (96), pH changes (97), or nutrient starvation (98). Bacteria in the VBNC state cannot be recovered with routine growth media, but are alive (i.e., they metabolic activity) and these cells can become culturable again on resuscitation (99). *Salmonella* is one of the bacteria reported using VNBC as a survival strategy in harsh environments, including aquatic systems (100–102). The existence of VBNC has been controversial, since although it was reported 30 years ago, there is still little direct conclusive



information on the molecular mechanisms underlying VNBC state induction and resuscitation (92, 103).

As mentioned above, many bacterial pathogens are known to survive within free-living protozoa, particularly amoebae (104). In fact, more than 20 species of pathogenic bacteria have been reported to associate with a single species (i.e., Acanthamoeba) of amoebae (105). This provides a potential mechanism for bacterial survival in the aquatic environment with the amoebae serving as an environmental reservoir. Salmonella Typhimurium was shown to replicate and survive in Acathamoeba spp.; however, it was cytotoxic and killed the amoebae (106-108). Douesnard-Malo and Daigle investigated the interactions between Acathamoeba castellanii and Salmonella Typhi, the etiologic agent of typhoid fever (109). They showed that S. Typhi could survive at least 3 weeks when grown in coculture with A. castellanii as opposed to less than 10 days when grown as singly cultured bacteria under the same conditions. Additionally, growth rates of amoebae after 14 days were similar in cocultures or when amoebae were singly cultured, suggesting that S. Typhi was not cytotoxic to A. castellanii (109). These studies suggest that certain species of free-living amoebae can serve as an environmental reservoir for pathogenic Salmonella species.

With an increasing number of outbreaks associated with consumption of fresh produce, understanding the mechanisms of produce contamination by foodborne pathogens is useful

to develop preventative and processing measures to curtail the microbial populations in produce. A relatively new topic regarding the mechanisms of produce contamination is the concept of internalization of produce by pathogens, which has given rise to debate in the field in the last two decades (43, 110-113). Internalization is defined as the uptake of pathogens through the roots into the intercellular spaces between plant cells and in the plant vasculature tissues, xylem and phloem (114). Internalized pathogens cannot be removed by washing or disinfection and thus can pose a risk for human health if contaminated produce is eaten uncooked (115). The uptake of different foodborne microbes including bacteria (Salmonella and E. coli) (42, 116-121), viruses (norovirus) (122-124), fungi (125), and protozoa (126) through roots into produce has been reported. However, the presence of pathogens including Salmonella inside plant cells remains controversial (35), and further comprehensive and in-depth study is needed on the internalization of produce by pathogens.

# INDIRECT DETECTION OF SALMONELLA IN IRRIGATION WATER

Rapid and accurate detection/estimation of the levels of *Salmonella* and other pathogens is prerequisite for understanding the dynamics of microbial populations and determination of

microbial quality of irrigation water (102). The major source of microbial contamination of irrigation water is from fecal origin (4, 127, 128), and thus the microbial quality guidelines of irrigation water are through testing total counts of coliforms, fecal coliforms, E. coli, fecal streptococci, and nematode eggs (23, 129). Because total counts of coliforms and fecal coliforms cannot exclude bacteria from nonfecal origin, the presence of E. coli is now regarded as a better indicator of microbial quality of irrigation water (23, 129-131). The PSR establishes microbial quality criteria for various uses of agricultural water using generic E. coli as an indicator for fecal pollution. However, the use of generic E. coli as an index organism for the presence of human pathogens in water sources has been discussed in the literature as well. The PSR also requires that agricultural water must be safe and of adequate sanitary quality for its intended use. It has been shown that the concentration of E. coli can predict the level of Salmonella present in water (6, 132). However, Salmonella could be identified even if E. coli counts were below the actionable levels, raising a concern that the satisfaction of the current limit of generic E. coli may not necessarily represent the absence of Salmonella (6). Accurate estimation/enumeration and isolation of Salmonella in water are still indispensable in source-tracking investigations of produce associated outbreaks.

# DIRECT DETECTION OF SALMONELLA IN IRRIGATION WATER

Traditional methods for detection, isolation, and identification of Salmonella in water involve nonselective and selective preenrichment in liquid culture followed by isolation using selective and differential agar plates. Such methods are laborious and time-consuming, which may take 4-5 days to complete (133). To overcome these limitations, immunoassays such as enzymelinked immunosorbent assay (ELISA) have been combined with culture methods for detection of Salmonella (134), but the application of ELISA is greatly hampered by its poor performance in sensitivity and specificity. Fortunately, a combination of culture-based methods with DNA-based technologies has given a great boost for Salmonella detection in various food matrices. DNA-based methods, including conventional and real-time PCR (qPCR), have become the most common methods for the detection of Salmonella (135-137), with qPCR being more advantageous due to its specificity, sensitivity, and short turnround time.

There are two different approaches used in qPCR detection. One approach uses SYBR Green dye to nonspecifically bind to synthesized double DNA (dsDNA). This dye only fluoresces when bound to dsDNA, thus the intensity of fluorescence quantitatively reflects the amount of the newly synthesized dsDNA (138, 139). The other approach is specific, employing a DNA probe with a fluorescent reporter incorporated at one end and a quencher of fluorescence at the other end to prevent detection of fluorescence. Degradation of the probe by 5'-3' nuclease activity of TaqMan polymerase will allow the unquenched emission of fluorescence, and the probe is also complementary to the DNA target and can anneal in each

cycle. Thus, the increase of fluorescence can proportionately reflect the amplification of the DNA product. The advantage of this method is that it can be used to detect multiple targets simultaneously with high specificity and sensitivity and is widely used in detection and identification of microorganisms (140–142). The Loop-Mediated Isothermal Amplification (LAMP) can be carried out at a constant temperature and the amplicons are detected by measuring turbidity or fluorescence (143–147). It is a good option for detection of *Salmonella* from waters in rural area or developing countries where thermocycler is not equipped or budget is limited.

Most conventional PCR and qPCR assays for Salmonella target the invasive gene (invA) because it is an important virulence factor gene (148). This locus is considered to be present in all Salmonella spp., including a wide range of Salmonella serotypes and absent in other closely related bacterial species and genera (149-151). The biological confirmation of positive PCR results of irrigation water samples with convention methods should proceed, especially when the results are to serve the regulatory purpose. This can be achieved by plating on conventionally-used selective growth agars, including XLD (Xylose Lysine Deoxycholate) (152), HE (Hektoen eteric) (153), and BS (Bismuth sulfite, a modification of Wilson and Blair agar) (154) agar. The presumed Salmonella colonies on these plates based on morphology may undergo further biochemical or molecular identification process. Salmonella is urease-negative, hence, this trait can also be used for differentiation from some urease-positive bacteria such as Proteus and Citrobacter species on urea medium with phenol red as in indicator (155).

# ENUMERATION OF SALMONELLA IN IRRIGATION WATER

In general, the level of Salmonella cells in food, poultry, produce, or other food matrices is low, and direct enumeration of Salmonella cell number has always been a challenge (156). A variety of methods have been established, including direct plating, fluorescence in situ hybridization (157), most probable number (MPN) (158), modified MPN methods (132, 156), and qPCR (135, 159, 160). These methods can be applied for enumeration of Salmonella populations in irrigation water, however, direct plating sometimes is inefficient due to low level of Salmonella cells in water and competitiveness from natural microbiota (161). Direct counting by immunofluorescence-based methods are also not widely used due to low sensitivity in enumeration, problems of antibody quality and linkage of fluorochrome (162). The MPN method is still widely used, in particular, for determination of low concentration of Salmonella samples (156), and it has been improved by combination with serology or multiplex PCR in the confirmation step in the MPN method (158, 163). McEgan et al. have developed a modified MPN method for irrigation water. This modified MPN includes a three-by-three MPN dilution test, selective enrichment, plating, biological confirmation, and PCR confirmation, and it normally takes about a week to complete the entire process (132).

Detection of low level of Salmonella in food or water samples with molecular methods typically requires a pre-enrichment step to increase the number of the target cells, but the high background microbiota in some foods, such as fresh produce, will also multiply in this process and consequentially complicates the subsequent steps (164). Methods for quantification of Salmonella in foods have been developed by using qPCR to detect the DNA of the samples in order to estimate the copy number of a target gene that reflects the quantity of bacterial cells (135, 165). Elimination of the enrichment process presumably may improve the accuracy of the quantification, but this approach is not without drawbacks. First, only a very small amount of DNA template can be added to the PCR reaction mixture, and DNA purity and proportion of target DNA in the total DNA may affect the accuracy of the results. Therefore, it becomes critical to select the appropriate preparation method for separation of bacteria in water or food. The other issue involves the inability of PCR to discriminate DNA from viable cells and dead cells, which may undermine the reliability of the assay. One way to overcome this issue is to detect the presence of RNA (166, 167), which is laborious and can result in false-negative results due to degradation of RNA. The other approach involves using a photoreactive dye propidium monoazide (PMA), which has been incorporated into the PCR to differentiate dead from live bacterial cells in foods (142, 168–170). The principle for PMA's selectivity in detection is based on its ability to penetrate only cell membranes of dead cells, covalently bind to DNA upon light exposure, and subsequently inhibit the modified DNA from amplification by PCR. But caution should be exercised because in some cases inhibition of amplification of PMA-bound DNA from dead cells was found incomplete (171-174).

### REDUCING SALMONELLA IN IRRIGATION WATER

Irrigation water is extensively applied during the produce growth stage, and thus the assurance of microbial quality of irrigation water is critical in the protection of produce

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safety. In the United States, the standards for drinking water (the Safe Drinking Water Act, SDWA) was enacted in 1974, and subsequently the Environmental Protection Agency set the drinking water standards, which requires total coliforms (including the fecal coliforms and E. coli) to be 0 per 100 ml water (EPA) (https://www.epa.gov/ground-water-anddrinking-water/national-primary-drinking-water-regulations). Developing general standards for water used for irrigation is rather complicated because the source of water for irrigation can vary greatly with region, season, and climate. Even within the same region, water microbial quality can be quite dynamic. Precipitation, the distance to domesticated animal raising facilities, and the number of wild or domestic animals in the proximity of the water can also constitute substantial changes to the microbial quality of the water (61). However, with the passage of FSMA, and more specifically, the PSR, many growers producing covered crops must now meet certain minimum requirements for the safe use of agricultural water. This may serve as a good solution for improving microbial quality of irrigation water reduction of foodbonre outbreaks assocaited to fresh produces.

#### **CONCLUSIONS**

Salmonella is frequently detected in surface water, which accounts for nearly half of the water used for irrigation. Traceback investigations of outbreaks often implicate irrigation water as a source (or a vehicle) for transmission of Salmonella. In addition, the bacterium can survive in these aquatic environments by a number of mechanisms, including entry into the VBNC state and/or residing within free-living protozoa. As such, assurance of microbial quality of irrigation water is vital to the mitigation of produce-related foodborne outbreaks.

#### **AUTHOR CONTRIBUTIONS**

BL conceived this project. HL, CW, BL wrote the manuscript. All authors have read and approved the manuscript.

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