Article

Presence of enterobacteria in agricultural inputs in La Comarca Lagunera

Claudia Zamantha Chavarría Hernández¹ Miguel Ángel Gallegos Robles^{2§} Manuel Fortis Hernández³ Uriel González Salas² María Gabriela Cervantes Vázquez² Edmundo Castellanos Pérez²

¹Sciences in Sustainability of Agricultural Resources-UJED. Highway Gómez Palacio-Tlahualilo km 28, Ejido Venice, Durango. CP. 35170. Tel. 01 (871) 7118918. (zamantha-chavarria@hotmail.com). ²Faculty of Agriculture and Zootechnics-Juarez University of the State of Durango. Highway Gómez Palacio-Tlahualilo km 28, Ejido Venice, Durango. CP. 35170. Tel. 01 (871) 7118918. (urgosa87@hotmail.com; cevga@hotmail.com; ecastellmx@yahoo.com.mx). ³Division of Postgraduate Studies-Technological Institute of Torreón. Road Torreón-San Pedro km 7.5, Ejido Ana, Torreón, Coahuila, Mexico. CP. 27070. Tel. (871) 7507199. (fortismanuel@hotmail.com).

[§]Corresponding author: garoma64@hotmail.com.

Abstract

The global food trade situation forces exporting countries to strengthen their control systems and adopt and monitor safety control strategies based on the risk of contamination. The objective of this research was to isolate enterobacteria in various inputs used in agricultural production in La Comarca Lagunera and to determine the risk involved in its use in agricultural activities. Samples were microbiologically analyzed from different sources of water, manures and soil, used in the agricultural production of La Comarca Lagunera, Mexico, during the years 2014-2016. *Klebsiella* spp. to was identified as the main risk factor of bacterial contamination in water that is used for irrigation of different crops with a prevalence of 46.95%. In the soil samples, only *Klebsiella* spp. (40%), *Salmonella* spp. (14%) in goat manure, and *Salmonella* spp. (100%) in chicken manure. The above represents a risk factor for the health of the final consumer, in the case of producers who intend to export their agricultural products, they would reduce competitiveness and could close international markets. Further investigation is necessary along the fruit and vegetable production chain in order to reduce this contamination.

Keywords: E. coli, Klebsiella spp., Salmonella spp., agricultural safety.

Reception date: April 2019 Acceptance date: June 2019

Introduction

Fruits and vegetables provide essential nutrients such as vitamins, fiber, minerals and have multiple health benefits, however, they are also recognized as potential vehicles of foodborne diseases (ETA) (Mesbah *et al.*, 2017). It is currently considered that to ensure the food safety of agricultural products it is necessary to prevent contamination in the field and minimize cross contamination during post-harvest management, however, preventing contamination in fields or greenhouses is a challenge and even Good agricultural practices (GAP) are insufficient to ensure that human pathogens are not introduced into the chain of agricultural products (Francis *et al.*, 2012).

The open-field production of horticultural products makes them susceptible to contamination by multiple sources. Several reviews have been made on the source of contamination with soil, water, biological amendments and the activity of wild animals, all considered as routes by which human pathogens can be introduced (Warriner *et al.*, 2009; Olaimat and Holley, 2012; Goodburn and Wallace, 2013; Nuesch-Inderbinen and Stephan, 2016).

Even in greenhouses, which are considered closed environments, pathogens such as *Salmonella* can be established (Holvoet *et al.*, 2014). It is documented that enteric pathogens can survive in manure, water and soil for prolonged periods (Franz *et al.*, 2005; Ferens and Hovde, 2011; Brandl *et al.*, 2013) and bind tenaciously to a variety of seeds of plants, roots and leaves (Tyler and Triplett, 2008; Teplitski *et al.*, 2011).

In Mexico, vegetable production represents 1.6% of the total value of national agricultural production, and in La Laguna of Coahuila and Durango, it amounts to 16.12 billion pesos (SIAP, 2016). In recent years, fresh fruits and vegetables have attracted the attention of researchers not only because of the benefits that their consumption brings to the health of people, but also because of their association with various outbreaks of gastrointestinal diseases registered in the last decade (Xuan *et al.*, 2000). In recent years, the origin of foodborne diseases (ETA) has changed, new pathogens or more aggressive and antibiotic resistant strains have emerged (Rojas-Herrera and González-Flores, 2006).

SENASICA promotes the implementation of pollution risk reduction systems (SRRC), with the aim of reducing the risk of contamination in the production of fruits and vegetables (SAGARPA-SENASICA, 2018).

Thus, the implementation of good agricultural practices (GAP) in the production units of vegetables should reduce the risks of contamination and allow compliance with the microbiological limits and pesticide residues established in national and international regulations. The SRRC have been implemented in recent years mainly in export crops such as chili, tomato, onion and tomato (Callejas *et al.*, 2006). Only in 2016, the cultivated area of these 3 vegetables in the lagoon region was more than 9639.03 ha, generating a production value of \$16 535 million pesos (SIAP, 2016).

The Comarca Lagunera is a region with great agricultural and livestock activity and due to the great cattle and poultry production that exists, 1 million tons of bovine manure and chicken manure are produced each year with a great potential to be used as organic fertilizer in agricultural production (Ramírez *et al.*, 2016). However, these inputs represent a high risk of pathogen transmission (Peralta-Veran *et al.*, 2016).

According to Sandoval and Colli (2004), another of the risk factors of microbial contamination of horticultural products is the practice of irrigation with rounded water from natural tributaries, contaminated wells and the use of untreated wastewater. In the Lagunera Region 60% of the horticultural production is irrigated with water from the water waterwheel and 40% with river water (SIAP, 2016). Although the use of wastewater for irrigation provides a large number of nutrients, some studies indicate that the use of untreated water for agricultural irrigation is the practice that most influences the reduction of the sanitary quality of fruits and vegetables (Cazárez *et al.*, 2004; Allende *et al.*, 2008).

The ETA is a major problem worldwide because they cause high morbidity and mortality, according to the world health organization, it is estimated that each year around 600 million people in the world are ill, 1 in 10 people ingest food. contaminated and 420 000 die from this same cause (OMS, 2015). Food can be contaminated along the productive chain and transport (Fatem, 2011).

ETA-causing bacteria that are listed as dangerous include enterohemorrhagic *Escherichia coli* (particularly serovar O157: H7), *Campylobacter jejuni*, and *Salmonella typhimurium* (Blackburn and McClure, 2002). According to the CDC (2017), the bacterium *Salmonella* has been found in fruits of tomato, melon, pumpkin and mango, while *E. coli* O157:H7 in romaine lettuce and organic spinach.

The objective of this research was to identify the presence of enterobacteria in various sources of inputs used in agricultural production in La Comarca Lagunera and determine the risk involved in their use in agricultural activities.

Materials and methods

Sampling

The sampling of the various samples was made in the months of August to September during the years 2014-2016, in common and agricultural areas of the municipalities of San Pedro, Francisco I. Madero and Matamoros in the state of Coahuila, and of the municipalities of Lerdo and Tlahualilo in the state of Durango (Table 1).

	and a second sec	L
Type of sample	Location	Ν
Water from River Nazas	Common León Guzmán [*]	10
	Dam Francisco Zarco *	20
Water form well	Common Hidalgo **	15
	Fco. I. Madero ^{**}	20
	Common Porvenir **	15
	Common Venecia **	27

Table 1. Type of samples analyzed in La Comarca Lagunera and number of samples.

Type of sample	Location	Ν
Residual water	Gómez Palacio *	20
Water form canal	Common Hidalgo **	10
	Common Escuadrón **	10
	Common Hormiguero **	12
	Common Coyote **	20
	Common 20 de noviembre **	12
	Common Jaboncillo **	8
	Common Santa María **	7
	Common Presa de Guadalupe **	8
	Common Finisterre **	10
	Common Florida **	10
	Common San Felipe *	12
	Common Viñedo [*]	10
	Common Venecia *	20
	Common Lucero *	20
	Tlahualilo *	20
	Matamoros **	12
Soil (onion growing)	Fco. I. Madero ^{**}	27
	Matamoros **	22
	Common Hidalgo **	13
Soil (chili cultivation)	Common Porvenir **	12
	San Pedro de las Col. **	10
Soil (tomato crop)	Common Hidalgo **	20
	Common Florida **	16
	Common Venecia *	10
Bovine manure	Fco. I. Madero ^{**}	20
	Common Venecia *	20
	Common Lucero *	20
Goat manure	Common Florida **	15
	Common Rosita **	15
	Common Venecia *	20
Chicken manure	Common Florida **	10
	Common Hidalgo **	10
	Common Esfuerzo **	12
Total		600

*= located in the state of Durango; **= located in the state of Coahuila. A total of 600 samples were collected (agricultural land 130, manure 142 and water 328).

The methodology for taking, handling and transporting samples was based on NOM-230-SSA1-2002 for water samples, and NOM-004-SEMARNAT-2002 for soil and manure samples.

The samples for water consisted of 500 ml, the samples for soil of 500 g and the samples for the manure of 300 g in fresh and 500 g in dry. After being collected, the samples were immediately stored on ice at a temperature below 10 $^{\circ}$ C and transported to the laboratory where they were stored at 4 $^{\circ}$ C. The microbiological analyzes were carried out within 24 h after the collection of the samples.

Microbiological analysis

The analysis for the presence of fecal coliforms was carried out following the NOM-004-SEMARNAT-2002. In the presumptive test, 1 ml of each dilution prepared in tubes containing lactosed broth was passed and incubated at 35 °C \pm 0.5 °C for 24 \pm 2 h, + 24 h if necessary. The positive tubes of the presumptive test were reseeded by triple roasting (sterilized to the lighter and cooled) in tubes containing EC broth and incubated at 44.5 \pm 0.2 °C in a water bath for 24 \pm 2 h. The result was positive when there was gas production from the fermentation of the lactose contained in the EC medium.

The positive tubes were transferred to Mac Conkey sorbitol agar with cefixime and tellurite, incubated from 18 to 24 h at 37 °C, then transferred to TSAYE agar and EMB agar, and incubated for 24 h at 35 °C. The colonies characteristic of *E. coli* was isolated and subjected to the biochemical tests of indole (+), TSI (A/A + Gas), Citrate of Simmons (-), sorbitol (-), being sorbitol negative. Since the Enterobacteria group also includes *Klebsiella*, it was isolated in the samples analyzed with this methodology.

The isolation and identification of *Salmonella* was carried out according to the protocol described by Andrews *et al.* (2007), which briefly describes, 25 mL or 25 g of sample were passed to 225 mL of lactose broth and incubated for 24 h at 35 °C. After the incubation period, the samples were transferred to tetrathionate broth (1 mL of sample plus 10 mL of tetrathionate broth) and allowed to incubate from 18 to 24 h at 42 °C, to be later sown on XLD agar (Becton Dickinson de México).

These plates were incubated from 22 to 50 h at 35 °C. Colonies with *Salmonella* characteristics were passed to biochemical tests on triple sugar and iron agar (TSI), iron-lysine agar (LIA) and Simmons citrate agar (Becton, Dickinson and Company EE UU). The identification of the genus was carried out through the biochemical tests of LIA, TSI and Simmons' Citrate.

The microbiological analyzes were performed in the microbiology laboratory of the Faculty of Agriculture and Zootechnics, located at km 28 of the Gómez Palacio-Tlahualilo highway, Durango, in the common Venecia, Municipality of Gómez Palacio, Durango, Mexico.

Results and discussion

Analysis of water samples

56.09% (184/328) of water samples (Table 2) presented contamination with some enterobacteria of the most commonly mentioned by the literature (Brisse *et al.*, 2009; Puerta-Garcia and Mateos-Rodríguez, 2010). The prevalence by microorganism isolated was as follows: *Klebsiella* spp. 46.95% (154/328), *Proteus* spp. 8.53% (28/328), *E. coli* 7.92% (26/328), *Salmonella* spp. 7.31% (24/328). In this study, *Klebsiella* was identified as the main risk factor for bacterial contamination in the water used to irrigate the different crops in the Comarca Lagunera. *Klebsiella* spp. it is a ubiquitous pathogen and together with *E. coli* and *Enterobacter* spp. they are bacteria with the capacity to cause gastroenteritis (by ingestion), respiratory infections, in the ears, skin and others (Rodríguez *et al.*, 2017).

Type of sample	Location	Ν	Microorganism	Prevalence
Water from river	Common Leon Guzmán [*]	10	Klebsiella	10/10
Nazas	Dam Francisco Zarco *	20	Klebsiella	10/20
Water form well	Common Hidalgo *	15	Proteus and	10/15
	Fco. I. Madero [*]	20	Klebsiella	15/20
	Common Porvenir *	15	Klebsiella	15/15
	Common Venecia *	27	Klebsiella Salmonella	3/27
Residual water	Gomez Palacio *	20	E. coli and Klebsiella	20/20
Water form canal	Common Hidalgo *	10	Proteus and	10/10
	Common Escuadron *	10	Klebsiella	8/10
	Common Hormiguero *	12	Proteus and	5/12
	Common Coyote *	20	Klebsiella	16/20
	Common 20 de noviembre *	12	Salmonella	7/12
	Common Jaboncillo *	8	Klebsiella	7/8
	Common Santa Maria *	7	Salmonella	2/7
	Common Presa de	8	Klebsiella	3/8
	Guadalupe *	10	Klebsiella	2/10
	Common Finisterre *	10	Klebsiella	2/10
	Common Florida *	12	E. coli	3/12
	Common San Felipe **	10	Salmonella	1/10
	Common Viñedo **	20	Salmonella	3/20
	Common Venecia **	20	Salmonella	15/20
	Common Lucero **	20	Salmonella	13/20
	Tlahualilo **	12	Klebsiella	4/12
	Matamoros *		Klebsiella	
			E. coli	
Total		328		184/328

 Table 2. Isolated microorganism and prevalence in four types of water samples used in the irrigation of crops in La Comarca Lagunera.

*= located in the state of Durango; **= located in the state of Coahuila.

The presence of *Klebsiella* in the water samples analyzed in this work represents a risk to human health and can become a big problem in the control of infections caused by this bacterium, since *Klebsiella* strains resistant to third generation cephalosporins have been isolated in fruits and vegetables that are consumed fresh (Mesbah *et al.*, 2017) and whose source of contamination has been irrigation water. Another important pathogen detected in the water samples was *Salmonella* spp. This bacterium is found on a daily basis in samples of soil, water and vegetation (Puerta-García and Mateos-Rodríguez, 2010).

Its presence in water samples from irrigation canals represents a risk, since it can pass to fruits and vegetables irrigated with this water and from there to the final consumer (Goodburn and Wallace, 2013; Nuesch-Inderbinen and Stephan, 2016). The water samples taken from the Nazas River

showed a prevalence of contamination with enterobacteria of 66.66% (20/30), the samples of well water a value of 55.84% (43/77), the samples of residual water a value 100% (20/20) and the channel water samples a value of 50.24% (101/201). These values are lower than that found by Luczkiewicz *et al.* (2011), who determined a prevalence of fecal coliforms of 97% in water samples; however, they are higher than that reported by Mesbah *et al.* (2017) who found a frequency of contamination by enterobacteria equal to 9% in the irrigation water.

The presence of enterobacteria in the four types of water samples analyzed in this work, and which are used as irrigation water in the crops of La Comarca Lagunera, constitutes a risk factor, since there is evidence of contamination of food of origin agricultural irrigation water and an association of foodborne outbreaks with vegetables, juices and other contaminated products, especially minimally processed or raw consumption (De Giglio *et al.*, 2017).

Analysis of soil samples

In the different types of soil (Table 3) and independently of the culture established at the time of sampling, only the bacterium *Klebsiella* spp. with a prevalence of 60% (78/130).

	1 8		8	
Type of sample	Localidad	Ν	Microorganism	Prevalence
Ground	Fco. I. Madero **	27	Klebsiella	15/27
(Onion growing)	Matamoros ^{**}	22	Klebsiella	10/22
	Common Hidalgo **	13	Klebsiella	10/13
Ground	Common Porvenir **	12	Klebsiella	8/12
(chili cultivation)	San Pedro de las Col **	10	Klebsiella	6/10
Ground	Common Hidalgo **	20	Klebsiella	10/20
(tomato crop)	Common Florida **	16	Klebsiella	11/16
	Common Venecia *	10	Klebsiella	8/10
Total		130		78/130

Table 3. Isolated microorganism and prevalence in three different soils according to the crop
established at the time of sampling in La Comarca Lagunera.

*= located in the state of Durango; **= located in the state of Coahuila.

In the soil where onion cultivation was established, a prevalence of 56.45% (35/62) was observed, for the soil where chili culture was established, a prevalence of 63.63% (14/22) and for the soil where there was tomato cultivation a prevalence of 63% was observed (29/46). *Klebsiella* spp. it is found in the environment on a daily basis, and the soil is not the exception. Rossmann *et al.* (2012) described the incidence of *Klebsiella* spp. in 11.4% in the soil rhizosphere and also reported the presence of enterobacteria with human pathogenic potential with less prevalence such as *Salmonella* and *E. coli*. In this study, neither *Salmonella* nor *E. coli* was detected in the soil samples.

Analysis of manure samples

Regarding the enterobacteria isolated in the three types of manure (Table 4), the prevalence was 41.54% (59/142). The prevalence by type of manure was as follows: cattle 33.3% (20/60), goat 14.0% (7/50) and 100% chicken manure (32/32).

Type of sample	Location	Ν	Microorganism	Prevalence
Bovine manure	Fco. I. Madero **	20	E. coli	6/20
	Common Venecia *	20	E. coli	6/20
	Common Lucero *	20	Klebsiella	8/20
Goat manure	Common Florida **	15	Salmonella spp.	3/15
	Common Rosita **	15	Salmonella spp.	4/15
	Common Venecia *	20	Salmonella spp.	0/20
Chicken manure	Common Florida **	10	Salmonella spp.	10/10
	Common Hidalgo **	10	Salmonella spp.	10/10
	Common Esfuerzo **	12	Salmonella spp.	12/12
Total		142		59/142

 Table 4. Isolated microorganism and prevalence in three different types of manure in La Comarca Lagunera.

*= located in the state of Durango; **= located in the state of Coahuila.

In bovine manure, *E. coli* and *Klebsiella* spp. were isolated, while *Salmonella* spp. was isolated only in goat and chicken manure. The above is a strong risk factor for human health, because the use of antibiotics to treat humans and animals in agriculture can lead to the selection of bacteria resistant to antibiotics that escape into the environment in manures (Durso and Cook, 2014).

Environmental isolates of enterobacteria that have acquired resistance to third generation cephalosporins (3GC) constitute a crucial threat to public health as sources of resistance for pathogenic bacterial strains that could lead to a failure in antibiotic therapy (Blaak *et al.*, 2014).

The presence of residues in the housing of pigs, cows, chickens, goats has been identified as an important risk factor for *Salmonella* infection (Beloeil *et al.*, 2004). *Salmonella* can survive for months in remains of dust and organic matter present in pens, equipment and ventilation systems (Rajic *et al.*, 2005), being this survival marked by some factors such as serotype and climatic conditions. Therefore, it is very important to use effective cleaning and disinfection protocols in the stables.

Several studies have proven that the application of an adequate cleaning and disinfection protocol is an important measure to prevent infection by *Salmonella* within the stables (Schmidt *et al.*, 2004; Hautekiet *et al.*, 2008).

In the case of the three types of manure analyzed (Table 3), it is observed that *Salmonella* contamination is found in the chicken manure and not cow manure, possibly due to the fact that the region is an important dairy basin, which may indicate that in bovine stables there are good excreta management practices. The development of diagnostics for enteropathogenic bacteria, associated with horticultural products, will allow to establish more adequate control measures in the production chain by identifying the sources of contamination such as irrigation water, manure, soil with a tendency to horticultural crops, towards which directed the investigation.

The results found in this research, highlight the importance of giving manure a treatment aimed at minimizing the burden of pathogenic microorganisms before being used as fertilizer in agricultural production, since it is shown that in addition to improving soil fertility, they are also carriers of zoonotic pathogens that have the potential to cause disease in humans (Hutchison *et al.*, 2005).

Conclusions

The inputs evaluated in this work and that are applied in agricultural production in La Comarca Lagunera represent a risk factor for the consumer due to the presence of enterobacteria. This, in addition to being a source of diseases transmitted by food, makes producers less competitive to sell their agricultural products. In addition, the final consumer should be informed about these microbiological health hazards and promote good housekeeping practices. It also follows that more research is needed along the fruit and vegetable production chain in order to involve more professionals in agricultural safety in actions to reduce this contamination.

Acknowledgments

To the Juárez University of the State of Durango for the financial support to carry out the research and to the National Council for Science and Technology for the scholarship granted to conclude my doctoral studies.

Cited literature

- Allende, A.; Selma, M. V.; López-Golvez, F.; Villaescusa, R. and Gil, M. I. 2008. Role of commercial sanitizers and washing systems on epiphytic microorganisms and sensory quality of fresh-cut escarole and lettuce. Postharvest Biol. Technol. 49(1):155-163.
- Andrews, W.; Wang, H.; Jacobson, A. and Hammack, T. 2007. Bacteriological analytical manual. Chapter 5. Salmonella. https://www.fda.gov/food/foodscienceresearch/laboratorymethods/ ucm070149.htm.
- Beloeil, P. A.; Fravalo, P.; Fablet C.; Jolly, J. P.; Eveno, E.; Hascoet, Y.; Chauvin, C.; Salvat, G. and Madec, F. 2004. Risk factors for *Salmonella enterica* subsp. enterica shedding by market-age pigs in French farrow-to-finish herds. Prev. Vet. Med. 63(1-2):103-120.
- Blaak, H.; van Hoek, A. H.; Veenman, C.; van Leeuwen, A. E.; Lynch, G.; van Overbeek, W. M.; de Roda Husman, A. M. 2014. Extended spectrum ß-lactamase- and constitutively AmpCproducing Enterobacteriaceae on fresh produce and in the agricultural environment. Int. J. Food Microbiol. (168-169):8-16. doi: 10.1016/j.ijfoodmicro.2013.10.006.
- Blackburn, C. and McClure, P. J. 2002. Foodborne pathogens: hazards, risk analysis and control. CRC Press. Boca Raton, FL. USA. 521 p.
- Brandl, M. T.; Cox, C. E.; Teplitski, M. 2013. *Salmonella* interactions with plants and their associated microbiota. Phytopathology. 103(4):316-325.
- Brisse, S.; Fevre, C.; Passet, V.; Issenhuth-Jeanjean, S.; Tournebize, R.; Diancourt, L. and Grimont, P. 2009. Virulent clones of Klebsiella pneumoniae: identification and evolutionary scenario based on genomic and phenotypic characterization. PLoS ONE. 4(3):1-13.
- Callejas, J. N.; Matus-Gardea, J. A.; García-Salazar, J. A.; Martínez-Damián, M. Á. y Salas-González, J. Ma. 2006. Situación actual y perspectivas de mercado para la tuna, el nopalito y derivados en el Estado de México. Agrociencia. 43(1):73-82.

- Cazárez, G.; Gortares, P.; Rubio, W.; Martínez, C.; Meza, P. y Chaidez, C. 2004. Presencia y supervivencia de coliformes fecales, *Salmonella* spp. y *Listeria* spp. en agua de uso agrícola en el valle de Culiacán. *In*: XIV Congreso Nacional. Federación Mexicana de Ingeniería Sanitaria y Ciencias Ambientales. Mazatlán, Sinaloa. 12 al 14 de mayo.
- CDC. Center of Disease Control. 2017. Foodborne Illness and Outbreaks. https://www.cdc.gov/ foodsafety/outbreaks/multistate-outbreaks/outbreaks-list.html.
- De Giglio, O.; Caggiano, G.; Bagordo, F.; Barbuti, G.; Brigida, S.; Lugoli, F.; Grassi, T.; La Rosa, G.; Lucentini, L.; Uricchio, V. F.; De Donno, A. and Montagna, M. T. 2017. Enteric viruses and fecal bacteria indicators to assess groundwater quality and suitability for irrigation. Int. J. Environ. Res. Public Health. 14(6):2-13.
- Durso, L. M. and Cook, K. L. 2014. Impacts of antibiotic use in agriculture: what are the benefits and risks? Curr. Opin. Microbiol. 19(1):37-44.
- Fatem, M. 2011. The incidence of Enterobacteriaceae causing food poisoning in some meat products. Adv. J. Food Sci. Technol. 3(2):116-121.
- Ferens, W. A. and Hovde, C. J. 2011. Escherichia coli O157:H7: animal reservoir and sources of human infection. Foodborne Pathog. Dis. 8(4):465-87.
- Francis, G. A.; Gallone, A.; Nychas, G. J.; Sofos, J. N.; Colelli, G.; Amodio, M. L. and Spano, G. 2012. Factors affecting quality and safety of fresh-cut produce. Crit. Rev. Food Sci. Nutr. 52(7):595-610.
- Franz, E.; van Diepeningen, A. D.; De Vos, O. J. and van Bruggen, A. H. C. 2005. Effects of cattle feeding regimen and soil management type on the fate of *Escherichia coli* O157:H7 and *Salmonella enterica* serovar typhimurium in manure, manure-amended soil, and lettuce. Appl. Environ. Microbiol. 71(10):6165-6174.
- Goodburn, C. and Wallace, C. A. 2013. The microbiological efficacy of decontamination methodologies for fresh produce: A review. Food Control. 32(2):418-427.
- Hautekiet, V.; Geert, V.; Marc, V. and Rony, G. 2008. Development of a sanitary risk index for *Salmonella* seroprevalence in Belgian pig farms. Prev. Vet. Med. 86(1-2):75-92.
- Holvoet, K.; Sampers, I.; Seynnaeve, M.; Jacxsens, L. and Uyttendaele, M. 2014. Agricultural and management practices and bacterial contamination in greenhouse versus open field lettuce production. Int. J. Environ. Res. Public. Health. 12(1):32-63.
- Hutchison, M.; Walters, L.; Moore, T.; Thomas, D.; and Avery, S. 2005. Fate of pathogens present in livestock wastes spread onto fescue plots. Appl. Environ. Microbiol. 71(2):691-696.
- Luczkiewicz, A.; Jankowska, K.; Kurlenda, J.; Olanczuk-Neyman, K. 2011. Identification and antimicrobial susceptibility of fecal coliforms isolated from surface water. Polish J. Environ. Stud. 20(4):941-950.
- Mesbah, Z. F.; Granier, S. A.; Marault, M.; Yaici, L.; Gassilloud, B.; Manceau, C.; Touati, A. and Millemann, Y. 2017. From farms to markets: gram-negative bacteria resistant to third generation Cephalosporins in fruits and vegetables in a region of North Africa. Front. Microbiol. 8:1569. doi:10.3389/fmicb.2017.01569.
- NOM-004-SEMARNAT-2002. Protección ambiental. -lodos y biosólidos. -especificaciones y límites máximos permisibles de contaminantes para su aprovechamiento y disposición final. http://dof.gob.mx/nota-detalle.php?codigo=691939&fecha=15/08/2003.
- NOM-230-SSA1-2002. Salud ambiental. Agua para uso y consumo humano, requisitos sanitarios que se deben cumplir en los sistemas de abastecimiento públicos y privados durante el manejo del agua. Procedimientos sanitarios para el muestreo. http://www.salud.gob.mx/unidades/cdi/nom/230ssa102.html.

- Nuesch-Inderbinen, M. and Stephan, R. 2016. Fresh fruit and vegetables as vehicles of bacterial foodborne disease: a review and analysis of outbreaks registered by proMED-mail associated with fresh produce. J. Food Saf. Food Qual. 67(2):32-39.
- Olaimat, A. N. and Holley, R. A. 2012. Factors influencing the microbial safety of fresh produce: a review. Food Microbiol. 32(1):1-19.
- Peralta-Veran, L.; Juscamaita-Morales, J. y Meza-Contreras, V. 2016. Obtención y caracterización de abono orgánico líquido a través del tratamiento de excretas del ganado vacuno de un establo lechero usando un consorcio microbiano ácido láctico. Ecol. Apl. 15(1):1-10.
- Puerta-García, A. y Mateos-Rodríguez, F. 2010. Enterobacterias. Medicine. 10(51):3426-3431.
- Rajic, A.; Keenliside, J.; McFall, M.; Deckert, A.; Muckle, A.; O'Connor, B.P.; Manninen, K.; Dewey, C. and McEwen, S. 2005. Longitudinal study of *Salmonella* species in 90 Alberta swine finishing Farms. Vet. Microbiol. 105(1):47-56.
- Ramírez, J. A.; Figueroa, U.; Núñez, G.; Reta, D. G. y García, J. L. 2016. Evaluation of tillage methods and manure incorporation into corn silage production. RChSZA. 15(2):67-76.
- Rodríguez, R.; Retamozo-Chavez, R.; Aponte, H. y Valdivia, E. 2017. Evaluación microbiológica de un cuerpo de agua del ACR humedales de ventanilla (Callao, Perú) y su importancia para la salud pública local. Ecología Aplicada. 16(1):15-21.
- Rojas-Herrera, R. A. y González-Flores, T. 2006. Detección e identificación de bacterias causantes de enfermedades transmitidas por alimentos mediante la reacción en cadena de la polimerasa. Bioquimia. 31(2):69-76.
- Rossmann, B.; Müller, H.; Smalla, K.; Mpiira, S.; Baptist, J.; Staver, Ch. and Berg, G. 2012. Banana-associated microbial communities in uganda are highly diverse but dominated by Enterobacteriaceae. Appl. Environ. Microbiol. 78(14):4933-4941.
- SAGARPA-SENASICA. 2018. Secretaría de Agricultura, Ganadería, Desarrollo Rural, Pesca y Alimentación-Servicio Nacional de Sanidad-Inocuidad y Calidad Agroalimentaria. Guía de Evaluación de los SRRC. https://www.gob.mx/senasica/documentos/guias-para-laevaluacion-de-los-srrc.
- Sandoval, L. y Collí, J. 2004. Tratamiento integral de agua residual municipal, su desinfección y reuso en la agricultura. *In*: XXIX Congreso Interamericano de Ingeniería Sanitaria y Ambiental. San Juan, Puerto Rico.
- Schmidt, P. L.; O'Connor, A. M.; McKean, J. D. and Hurd, H. S. 2004. The association between cleaning and desinfection of lairage pens and the prevalence of *Salmonella* enterica in swine harvest. J. Food Prot. 67(7):1384-1388.
- SIAP. 2016. Servicio de Información Agroalimentaria y Pesquera. Cierre de la producción agrícola. Nopalitos. http://www.siap.gob.mx.
- Teplitski, M.; Warriner, K.; Bartz, J. and Schneider, K. R. 2011. Untangling metabolic and communication networks: interactions of enterics with phytobacteria and their implications in produce safety. Trends. Microbiol. 19(3):121-127.
- Tyler, H. L. and Triplett, E. 2008. Plants as a habitat for beneficial and human pathogenic bacteria. Annu. Rev. Phytopathol. 46:53-73.
- Warriner, K.; Huber, A.; Namvar, A.; Fan, W. and Dunfield, K. 2009. Recent advances in the microbial safety of fresh fruits and vegetables. Adv. Food Nutr. Res. 57:155-208.
- Xuan, G.; Chen, J.; Beuchat, L. R. and Brackett, R. E. 2000. PCR detection of *Salmonella* enterica serotype Montevideo in and on raw tomatoes using primers derived from *hilA*. Appl. Environ. Microbiol. 66(12):5248-5252.