Malaysian Journal of Microbiology, Vol 16(1) 2020, pp. 34-39 DOI: http://dx.doi.org/10.21161/mjm.180325



Malaysian Journal of Microbiology

Published by Malaysian Society for Microbiology (In SCOPUS since 2011)



Prevalence and antimicrobial resistance of *Staphylococcus aureus* isolated from vegetables sold in Malaysian wet markets

Mustapha Goni Abatcha^{1*}, Dauda Goni Mohammed², Adamu Abbas Muhammad³, Muhammad Jalo Ibrahim⁴ and Mohammed Goje⁵

¹Food Technology Division, School of Industrial Technology, Universiti Sains Malaysia, 11800 Minden, Penang, Malaysia.

²Department of Medical Microbiology and Parasitology, Universiti Sains Malaysia, Health Campus, 16150 Kubang Kerian Kelantan, Malaysia.

³Department of Medical Microbiology and Parasitology, Faculty of Clinical Sciences, College of Health Sciences, Bayero University Kano, Nigeria.

⁴Veterinary Service Department, Ministry of Agriculture and Environment, Damaturu Yobe State, Nigeria.

⁵Faculty of Veterinary Medicine, University of Maiduguri, Borno State Nigeria.

Email: mustygoni@gmail.com

Received 21 December 2018; Received in revised form 12 June 2019; Accepted 23 August 2019

ABSTRACT

Aims: *Staphylococcus aureus* is the most common pathogen found in humans, animals and foods worldwide. Vegetables contain essential vitamins, minerals, and fibres that may aid in protecting humans from chronic diseases and promote good health. This investigation aimed to find the prevalence of *S. aureus* contamination and antimicrobial resistance pattern of isolates from leafy vegetables in Peninsular Malaysia.

Methodology and results: A total of 397 samples, comprised of 16 different vegetables, were obtained from vendors in selected wet markets. Of the 397 samples, *S. aureus* was detected in 42 samples (10.6%) in which 9 (21.4%) were positive for methicillin-resistant *Staphylococcus aureus* (MRSA). The *S. aureus* isolates showed 52.3% resistance to tetracycline, followed by kanamycin (40.5%), penicillin G (35.7%), streptomycin (33.3%), oxacillin (21.4%) and cefoxitin (19.0%). The multiple antibiotic resistance indexes of *S. aureus* isolates varied from 0.21 to 0.50. The predominant antimicrobial resistance profiles of *S. aureus* were PDaOxSTeL (n=5), PDaSTeCe (n=4), TeKCnSxtC (n=3) and TeKCipQd, respectively.

Conclusion, significance and impact of study: The findings of this study revealed the consumption of raw or minimally prepared fresh leafy vegetables could be a possible source of infection with antimicrobial-resistant *S. aureus*.

Keywords: Staphylococcus aureus, antibiotic resistance, vegetables

INTRODUCTION

Leafy vegetables are well known as essential parts of a healthy and nutritious diet. Several countries, including Malaysia, have begun initiatives to inspire consumers to increase their consumption of these products (Denis *et al.*, 2016). However, fresh vegetables have been increasingly recognised to be among the leading food vehicles of foodborne microbial pathogens or hazards (Lynch *et al.*, 2009). Besides, fresh vegetables are also considered to harbour significant numbers of epiphytic microorganisms, mainly non-pathogenic (Alegbeleye *et al.*, 2018). In recent years, there have been a rising number of outbreaks of fresh produce-related foodborne illnesses in many parts of the world, and attempts are

being made to resolve these food safety harms (Jongman and Korsten, 2018).

Staphylococcus aureus is a major foodborne pathogen causing food intoxication worldwide (Kadariya *et al.*, 2014). This pathogen can contaminate numerous food products, including fresh vegetables and produce several kinds of enterotoxins, causing diseases with differing severity from superficial skin diseases to severe and potentially fatal, invasive ones (Bennett *et al.*, 2013). Studies published in Korea have shown that leafy vegetables such as Lettuce, sprouts, and perilla leaves, could be contaminated by *S. aureus* (Moon *et al.*, 2007; Seo *et al.*, 2010). In Malaysia, also, *S. aureus* is a wellknown pathogen associated with food poisoning (Fluit, 2012). Although some studies reported the bacterial contamination of raw leafy vegetables (Wadamori *et al.*,

*Corresponding author

2017), not much has been done in the area of *S. aureus* contamination (Puah *et al.*, 2016).

There has been an alarming surge in antibioticresistant S. aureus in both developed and developing countries over the past few decades (Fair and Tor, 2014; Kashani et al., 2018). The excessive use of antimicrobial agents as prophylactic, therapeutic and growth promoters in food animals has steered to the emergence of resistant strains and remains a significant public health challenge (Medina and Pieper, 2016). In many countries, Penicillin has remained the antibiotic of choice to treat S. aureus infections (Kali, 2015). However, the production of an enzyme called penicillinase or beta-lactamase causes resistance in several strains of S. aureus to a penicillin (Jamali et al., 2015). Most isolated S. aureus strains from foods and animals are increasingly resistant to many frequently used antibiotics including methicillin, penicillin, streptomycin, macrolides, gentamicin and tetracycline (Alenizi, 2014; Jamali et al., 2015). Nevertheless, the resistance patterns of isolated S. aureus from fresh leafy vegetables in recent years show the emergence of multidrug resistance (MDR) S. aureus strains which may cause a substantial threat to human health in Malaysia (Seo et al., 2010; Dhama et al., 2014). Therefore, this study aimed to examine the prevalence and antimicrobial resistance of S. aureus isolated from raw leafy vegetables obtained from wet markets in Malaysia.

MATERIALS AND METHODS

Sampling

In this study, a total of 397 samples comprised of 16 different raw leafy vegetables were purchased from selected wet markets located in four cities of Malaysia from May 2015 to March 2016. The selected vegetables were Indian pennywort (Pegaga), amaranth red (Bayam), sweet basil (Selasi), bean sprout (Kecambah), coriander (Ketumber), lettuce salad (Daun salad), spring onion (Daun bawang), Japanese parsley (Selom), winged bean (Kacang botol), laksa leaves (Kesom), amaranth green (Bayam), iceberg lettuce (Lettuce ais), mint (Daun pudina), wild parsley (Ulam raja), water spinach (Kankong) and Chinese flowering cabbage (Sawi bunga). The samples were collected in sterile plastic bags and transported instantly to the laboratory for analysis.

Isolation and detection of S. aureus

For the detection and isolation of *S. aureus*, 25 g (raw leafy vegetable) of each sample was added 225 mL of buffered peptone water (Merck, Germany) and homogenised. The samples were streaked onto Baird-Parker Agar (Merck, Germany) with an added 5% egg yolk and tellurite (Merck, Germany). The plates were then incubated at 37°C for 24-48 h. The suspected colonies with a typical black display in white circles on the Baird-Parker Agar were enumerated as *S. aureus* (Loncarevic *et al.*, 2005; Jamali *et al.*, 2015). The API Staph

identification system was used to confirm suspected colonies (bioM-erieux, Marcy-I'-Etoile, France).

Antimicrobial susceptibility assay

The resistance to antimicrobial agents was performed by the Kirby-Bauer disc diffusion method using ready antibiotic disks (Oxoid, UK) on Mueller-Hinton agar (Merck, Germany) according to the guidelines of the Clinical and Laboratory Standards Institute (CLSI, 2006). The following fourteen panels of antimicrobial agents (Oxoid, UK) were used: tetracycline (30 mg), kanamycin (30 mg), gentamicin (10 mg), penicillin G (10 U), oxacillin (1 mg), chloramphenicol (30 mg), erythromycin (15 mg), tobramycin (10 mg), lincomycin (15 mg), ciprofloxacin (5 mg), quinupristin-dalfopristin (15 mg), clindamycin (2 mg), trimethoprim-sulfamethoxazole (1.25/23.75 mg) and streptomycin (10 mg). Escherichia coli ATCC 35218 were used for quality control.

PCR for the amplification of the mecA gene

DNA templates were prepared using crude cell lysates as specified by Zhang et al. (2012). A PCR using the primers *mecA1* (5 AAAATCGATGGTAAAGGTTGGC) and *mecA2* (5-AGTTCTGCAGTACCGGATTTGC), was applied for the determination of *mecA* gene using conditions as reported by Murakami *et al.* (1991).

Table	1:	Prevalenc	e of	· S.	aureus	in	various	raw	leafy
vegeta	ble	samples of	obtai	ned	from we	et n	narkets.		

Samples	No. of Samples	Samples positive (%)	Prevalence (%)
Amaranth green	25	6	24.0
Amaranth red	25	2	8.0
Bean sprouts	25	3	12.0
Coriander	25	4	16.0
Water spinach	25	3	12.0
Winged bean	25	2	8.0
Laksa leaves	25	1	4.0
Iceberg lettuce	25	2	8.0
Mint	25	3	12.0
Spring onion	25	1	4.0
Indian pennywort	25	5	20.0
Wild parsley	25	2	8.0
Lettuce salad	23	1	4.3
Chinese flowering cabbage	25	3	12.0
Sweet basil	24	3	12.5
Japanese parsley	25	1	4.0
Total	397	42	10.6

Statistical analysis

Statistical analysis was performed using the Chi-square test to find the relationship between contaminated

samples and several types of vegetable. The variations were regarded as significant at $p \le 0.05$.

Table 2: Antimicrobial susceptibil	ty profile of S. aureu	s (n=42) isolated from v	arious raw leafy	vegetable samples
------------------------------------	------------------------	--------------------------	------------------	-------------------

Susceptible N (%)	Intermediate N (%)	Resistant N (%)
19 (45.2)	8 (19.0)	15 (35.7)
34 (80.9)	0 (0)	8 (19.0)
35 (83.3)	5 (11.9)	2 (4.7)
37 (88.1)	0 (0)	5 (11.9)
25 (59.5)	5 (11.9)	12 (28.6)
30 (71.4)	3 (7.1)	9 (21.4)
20 (47.6)	0 (0)	22 (52.3)
25 (59.5)	0 (0)	17 (40.5)
32 (76.1)	0 (0)	10 (23.8)
36 (85.7)	0 (0)	6 (14.3)
25 (59.5)	5 (11.9)	12 (28.60
28 (66.6)	0 (0)	14 (33.3)
35 (83.3)	2 (4.7)	5 (11.9)
37 (88.1)	0 (0)	5 (11.9)
36 (85.7)	0 (0)	6 (14.3)
	Susceptible N (%) 19 (45.2) 34 (80.9) 35 (83.3) 37 (88.1) 25 (59.5) 30 (71.4) 20 (47.6) 25 (59.5) 32 (76.1) 36 (85.7) 25 (59.5) 28 (66.6) 35 (83.3) 37 (88.1) 36 (85.7)	Susceptible N (%)Intermediate N (%)19 (45.2) 8 (19.0)34 (80.9)0 (0)35 (83.3) 5 (11.9)37 (88.1)0 (0)25 (59.5) 5 (11.9)30 (71.4) 3 (7.1)20 (47.6)0 (0)25 (59.5)0 (0)32 (76.1)0 (0)36 (85.7)0 (0)25 (59.5) 5 (11.9)28 (66.6)0 (0)35 (83.3) 2 (4.7)37 (88.1)0 (0)36 (85.7)0 (0)

obtained from wet markets.

 Table 3 Antibiogram of S. aureus isolated from various raw leafy vegetable samples obtained from wet markets.

Antibiogram	No. of isolates	Source	MAR index
PDaOxSTeL	5	Amaranth green (2), Mint (2), Laksa leaves	0.42
PDaSTeCe	4	Coriander, Japanese parsley, Water spinach, Indian pennywort	0.35
PDaEOxS	2	Bean sprouts, Chinese flower Cabbage	0.35
KPDaEOxSTe	1	Amaranth green	0.5
KPEOxTeL	1	Coriander	0.42
PSLCeK	2	Indian pennywort	0.42
KCeTor	2	Amaranth green, Indian pennywort	0.35
TeKCipQd	3	Amaranth red, Spring Onion, Sweet basil	0.28
TeKCnSxtC	3	Amaranth green (2), Lettuce salad	0.35
TeKCnSxtCQd	2	Sweet basil, Wild parsley	0.42
TeKCnC	1	Water spinach	0.28
TeKCip	2	Winged bean, Bean sprouts	0.21

RESULTS AND DISCUSSION

A total of 397 samples of raw leafy vegetables obtained from wet markets were investigated in this study. The overall prevalence of *S. aureus* isolated from the vegetables was 10.6% (42/397) as shown in Table 1, an indication that vegetables can be a possible cause of *S. aureus* infection in Malaysia. *S. aureus* was isolated from all the various types of vegetables examined.

The prevalence of *S. aureus* significantly varied between samples (P < 0.05). Incidentally, Amaranth green had the highest incidence of 24.0% (6/25), followed by Indian pennywort 20.0% (5/24) and coriander 16.0 % (4/25). Other studies also found a prevalence of *S. aureus*

in fresh produce and raw leafy vegetables. In Korea, a survey of fresh fruits and vegetables such as lettuce,

mixed salads and green onions reported contamination with *S. aureus* (Seo *et al.*, 2010) while another study published by Nguz *et al.* (2005) isolated it from vegetables such as radish, lettuce, and seed sprouts. According to Olaimat *et al.* (2016), vegetables can become contaminated with such pathogens while growing, during harvest, during post-harvest handling, or distribution. This could be due to improper human handling and storage, contaminated irrigation water, containers, animal waste fertilisers and post-harvest washings (Mritunjay *et al.*, 2015).

S. aureus contamination can also be found at the dining point. *S. aureus* is known to be carried by food handlers (Faour-Klingbeil *et al.*, 2016). High levels of *S. aureus* were detected in Russian salad (25.0%), and vegetable salad (12.0%) served in Turkish dining facilities (Aycicek *et al.*, 2005).

Because S. aureus is well known for the tendency to become resistant to some antimicrobials, this study also looked into the resistance pattern of S. aureus to some antimicrobial agents, as shown in Table 2. There was a high prevalence of resistance to tetracycline, kanamycin, penicillin G, and to a minor extent, quinupristindalfopristin, trimethoprim-sulfamethoxazole, ciprofloxacin and The antimicrobial tobramvcin. resistances demonstrated by the isolates were as follows; tetracycline 22/42 (52.3%), kanamycin 17/42 (40.5%), penicillin G 15/42 (35.7%), streptomycin 14/42 (33.3%), oxacillin 9/42 (21.4%) and cefoxitin 8/42 (19.0%). Furthermore, nine isolates (21.4%) were identified as Methicillin-resistant S. aureus (MRSA) by antibiotic susceptibility test, Oxacillinresistant and confirmed by detection of the mecA gene. This result was like a study by Hong et al., (2015) in which eleven S. aureus isolates from leafy vegetables were MRSA, which showed the presence of the mecA gene and phenotypic resistance to oxacillin.

The high prevalence of tetracycline, kanamycin and penicillin Ğ resistant S. aureus noticed in the present study, was similar to earlier studies reported by Jamali et al. (2015); Seo et al. (2010). Hong et al. (2015) also found almost 15 of 19 (78.95%) of S. aureus isolates isolated from perilla leaves being multidrug-resistant. This high prevalence of S. aureus resistance could be due to the wide use of animal manures as fertiliser in vegetable farming. Tetracycline, sulfonamides, and quinolone antibiotics have been detected in animal manures (chicken, cattle and swine) from China (Gao et al., 2012). Moreover, MRSA has increasingly been found in farm animal population and food of animal origin (Vicca et al., 2008). Other factors play a significant role in increasing the burden of environmental exposure to resistant bacteria. These come from several human activities, including emissions of antibiotic residues, fish production, animal production, wastewater treatment, and antibiotic manufacturing. Furthermore, antibiotics could be taken up by many vegetables if they were planted in soil contaminated by antibiotics (Hu et al., 2010). The results we obtained in this study showed that the most predominant antibiogram of *S. aureus* isolated from raw leafy vegetables were PDaOxSTeL (n = 5), PDaSTeCe (n = 4), TeKCnSxtC (n = 3) and TeKCipQd, respectively (Table 3). The MAR index of S. aureus isolated from vegetables ranged from 0.21 to 0.50 and was resistant to 2-7 various types of antibiotics. When isolates have a MAR index of 0.2 and above it indicates that strains were exposed to several antibiotics or originated from the environment where antibiotics were frequently used (Krumperman, 1983).

CONCLUSION

In summary, the presence of S. aureus from raw leafy vegetables retailed in wet markets is worrying. It showed that the consumption of fresh or minimally cooked vegetables might be a possible risk of foodborne bacterial pathogens. Increased antibiotic resistance amongst the S. aureus isolated from wet markets vegetables is also of great concern to the society. More importantly, the isolation of MRSA from vegetables obtained from wet markets is a public health concern for the consumers. The findings here highlighted the need for strict hygienic standards in wet markets to reduce the prevalence of S. aureus. The use of treated animal manures in agricultural practice also needs strict monitoring to curtail the emerging antimicrobial resistance bacteria in vegetables. The findings of this study highlighted the likely problems of the spread of S. aureus through raw vegetables and the need to curtail them.

ACKNOWLEDGEMENTS

The authors are thankful to USM-IPS for providing a USM Global fellowship.

REFERENCES

- Alegbeleye, O. O., Singleton, I. and Sant'Ana, A. S. (2018). Sources and contamination routes of microbial pathogens to fresh produce during field cultivation: A review. *Food Microbiology* **73**, **177-208**.
- Alenizi, D. A. (2014). Prevalence of Staphylococcus aureus and antibiotic resistance in children with atopic dermatitis in Arar, Saudi Arabia. Journal of Dermatology and Dermatologic Surgery 18(1-2), 22-26.
- Aycicek, H., Cakiroglu, S. and Stevenson, T. H. (2005). Prevalence of *Staphylococcus aureus* in ready-to-eat meals from military cafeterias in Ankara, Turkey. *Food Control* **16(6)**, **531-534**.
- Bennett, S. D., Walsh, K. A. and Gould, L. H. (2013). Foodborne disease outbreaks caused by *Bacillus cereus*, *Clostridium perfringens*, and *Staphylococcus aureus*-the United States, 1998-2008. *Clinical infectious diseases* 57(3), 425-433.
- Clinical and Laboratory Standards Institute (CLSI) (2006). Methods for antimicrobial dilution and disk susceptibility testing of infrequently isolated or fastidious bacteria; approved guideline. *Clinical and Laboratory Standards Institute* 26.
- Denis, N., Zhang, H., Leroux, A., Trudel, R. and Bietlot, H. (2016). Prevalence and trends of bacterial contamination in fresh fruits and vegetables sold at retail in Canada. *Food Control* 67, 225-234.
- Dhama, K., Tiwari, R., Chakraborty, S., Saminathan, M., Kumar, A., Karthik, K. and Rahal, A. (2014). Evidence-based antibacterial potentials of medicinal plants and herbs countering bacterial pathogens,

especially in the era of emerging drug resistance: An integrated update. *International Journal of Pharmacology* **10(1), 1-43.**

- Fair, R. J. and Tor, Y. (2014). Antibiotics and bacterial resistance in the 21st century. *Perspectives in Medicinal Chemistry* 6, 25-64.
- Faour-Klingbeil, D., Murtada, M., Kuri, V. and Todd, E.
 C. (2016). Understanding the routes of contamination of ready-to-eat vegetables in the Middle East. *Food Control* 62, 125-133.
- Fluit, A. C. (2012). Livestock-associated Staphylococcus aureus. Clinical Microbiology and Infection 18(8), 735-744.
- Gao, J., Ferreri, M., Yu, F., Liu, X., Chen, L., Su, J. and Han, B. (2012). Molecular types and antibiotic resistance of *Staphylococcus aureus* isolates from bovine mastitis in a single herd in China. *The Veterinary Journal* 192(3), 550-552.
- Hong, J., Kim, Y., Kim, J., Heu, S., Kim, S. R., Kim, K. P. and Roh, E. (2015). Genetic diversity and antibiotic resistance patterns of *Staphylococcus aureus* isolated from leaf vegetables in Korea. *Journal of food science* 80(7), 1526-1531.
- Hu, X., Zhou, Q. and Luo, Y. (2010). Occurrence and source analysis of typical veterinary antibiotics in manure, soil, vegetables and groundwater from organic vegetable bases, northern China. *Environmental Pollution* 158(9), 2992-2998.
- Jamali, H., Paydar, M., Radmehr, B., Ismail, S. and Dadrasnia, A. (2015). Prevalence and antimicrobial resistance of *Staphylococcus aureus* isolated from raw milk and dairy products. *Food Control* 54, 383-388.
- Jongman, M. and Korsten, L. (2018). Irrigation water quality and microbial safety of leafy greens in different vegetable production systems: A review. Food *Reviews International* 34(4), 308-328.
- Kadariya, J., Smith, T. C. and Thapaliya, D. (2014). Staphylococcus aureus and staphylococcal foodborne disease: An ongoing challenge in public health. BioMed Research International 2014, Article ID 827965.
- Kali, A. (2015). Antibiotics and bioactive natural products in the treatment of methicillin-resistant *Staphylococcus aureus*: A brief review. *Pharmacognosy Reviews* 9(17), 29.
- Kashani, H. H., Schmelcher, M., Sabzalipoor, H., Hosseini, E. S. and Moniri, R. (2018). Recombinant endolysins as potential therapeutics against antibioticresistant *Staphylococcus aureus*: current status of research and novel delivery strategies. *Clinical Microbiology Reviews* 31(1), 1-26.
- Krumperman, P. H. (1983). Multiple antibiotic resistance indexing *Escherichia coli* to identify risk sources of faecal contamination of foods. *Applied and Environmental Microbiology* 46, 165-170.
- Loncarevic, S., Jørgensen, H. J., Løvseth, A., Mathisen, T. and Rørvik, L. M. (2005). Diversity of Staphylococcus aureus enterotoxin types within single

samples of raw milk and raw milk products. *Journal of Applied Microbiology* **98(2), 344-350.**

- Lynch, M. F., Tauxe, R. V. and Hedberg, C. W. (2009). The growing burden of foodborne outbreaks due to contaminated fresh produce: Risks and opportunities. *Epidemiology and Infection* 137(3), 307-315.
- Medina, E. and Pieper, D. H. (2016). Tackling threats and future problems of multidrug-resistant bacteria. *In:* How to Overcome the Antibiotic Crisis. Current Topics in Microbiology and Immunology. Stadler M., Dersch P. (eds.). vol. 398 Springer, Cham, Switzerland. pp. 3-33.
- Moon, J. S., Lee, A. R., Jaw, S. H., Kang, H. M., Joo, Y. S., Park, Y. H. and Koo, H. C. (2007). Comparison of antibiogram, staphylococcal enterotoxin productivity, and coagulase genotypes among *Staphylococcus aureus* isolated from animal and vegetable sources in Korea. *Journal of Food Protection* 70(11), 2541-2548.
- Mritunjay, S. K. and Kumar, V. (2015). The fresh farm produces as a source of pathogens: A review. Research Journal of Environmental Toxicology 9(2), 59-70.
- Murakami, K., Minamide, W., Wada, K., Nakamura, E., Teraoka, H. and Watanabe, S. (1991). Identification of methicillin-resistant strains of staphylococci by a polymerase chain reaction. *Journal of Clinical Microbiology* 29(10), 2240-2244.
- Nguz, K., Shindano, J., Samapundo, S. and Huyghebaert, A. (2005). Microbiological evaluation of fresh-cut organic vegetables produced in Zambia. *Food Control* 16(7), 623-628.
- Olaimat, A. N., Delaquis, P. J. and Holley, R. A. (2016). Impact of materials handling at pre-and post-harvest operations on the microbial ecology of foods of vegetable origin. *Quantitative Microbiology in Food Processing: Modeling the Microbial Ecology* 2016, 85-116.
- Puah, S. M., Chua, K. H. and Tan, J. A. M. A. (2016). Virulence factors and antibiotic susceptibility of *Staphylococcus aureus* isolates in ready-to-eat foods: Detection of *S. aureus* contamination and a high prevalence of virulence genes. *International Journal of Environmental Research and Public Health* 13(2), 199.
- Seo, Y. H., Jang, J. H. and Moon, K. D. (2010). Occurrence and characterisation of enterotoxigenic *Staphylococcus aureus* isolated from minimally processed vegetables and sprouts in Korea. *Food Science and Biotechnology* 19(2), 313-319.
- Strommenger, B., Kettlitz, C., Werner, G. and Witte, W. (2003). The multiplex PCR assay for simultaneous detection of nine clinically relevant antibiotic resistance genes in *Staphylococcus aureus*. *Journal* of *Clinical Microbiology* **41(9)**, **4089-4094**.
- Vicca, J., Vanderhaegehen, W., Cerpentier, T. and Butaye, P. (2008). Prevalence at herd-level of methicillin-resistant *Staphylococcus aureus* in milk samples of dairy herds. *Mastitis Control-from Science* to Practice 2008, 71-75.

- Wadamori, Y., Gooneratne, R. and Hussain, M. A. (2017). Outbreaks and factors are influencing microbiological contamination of fresh produce. *Journal of the Science of Food and Agriculture* 97(5), 1396-1403.
- Zhang, Q. Y., Zhou, W. W., Zhou, Y., Wang, X. F. and Xu, J. F. (2012). Response surface methodology to design a selective co-enrichment broth of *Escherichia coli*, *Salmonella* spp. and *Staphylococcus aureus* for simultaneous detection by multiplex PCR. *Microbiological Research* 167(7), 405-412.