

Kinetic Behavior of Pathogenic *Escherichia coli* and *Staphylococcus aureus* in Fresh Vegetables during Storage at Constant and Changing Temperature

Keywords: Fresh vegetables; *E. coli*; *S. aureus*; Kinetic model; Mathematical model

Abstract

This study used mathematical models to describe the kinetic behavior of pathogenic *Escherichia coli* and *Staphylococcus aureus* in lettuce, commercial sprout mix and water celery during storage at constant and changing temperature. μ_{max} values of *E. coli* in lettuce (0.002-0.294 log CFU/g/h) and water celery (0.002-0.325 log CFU/g/h) significantly increased ($P < 0.05$) as the temperature increased, but not in sprout mix. Regarding *S. aureus*, growth (0.003-0.024 log CFU/g/h) was observed on lettuce, but no growth in other vegetables. The square root model was appropriate to describe the temperature effect on the μ_{max} of *E. coli* in lettuce ($R^2 = 0.900$) and water celery ($R^2 = 0.951$), and μ_{max} of *S. aureus* in lettuce ($R^2 = 0.947$). RMSE (*E. coli*: lettuce = 0.691, water celery = 0.745; *S. aureus*: lettuce = 0.569) suggested that mathematical models were appropriate. The simulation with a dynamic model showed the gradual increase of *E. coli* cell counts in lettuce and water celery at changing temperature. The results indicate that pathogenic *E. coli* can be problematic in lettuce and water celery not in sprout mix, but *S. aureus* may not be risky in the vegetable examined in this study for both constant and changing temperature.

Introduction

The consumption of fresh vegetables has been increasing because of their health benefits; however, foodborne outbreaks are more often associated with such produce because fresh vegetables are usually consumed raw [1-4], and therefore, fresh vegetable can be a vector to transfer pathogenic bacteria to consumer [5]. Foodborne illness caused by fresh vegetables is often associated with pathogenic *Escherichia coli* such as enterotoxigenic *E. coli*, enterohemorrhagic *E. coli*, enteroaggregative *E. coli*, enteroinvasive *E. coli*, and enteropathogenic *E. coli* [6-8]. *Staphylococcus aureus* is also a common cause of foodborne illness because it can resist to various harmful environments [9,10]. Kim et al. also isolated *E. coli*, *Salmonella* spp., and *S. aureus* from various salads. Jo et al. recently found the evidence of *E. coli*, *S. aureus*, *Salmonella* spp., and *Listeria monocytogenes* contamination in fresh-cut produce and organic vegetables [11,12]. These studies then suggested that foodborne disease can be derived from contaminated fresh vegetables and fruits [13,14]. However, the growth patterns of foodborne pathogens can be different by storage temperature, especially for changing temperature during storage, and type of vegetables. Thus, it is necessary to describe



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the kinetic behavior of foodborne pathogens in various vegetables and at different temperature patterns.

To describe the kinetic behavior of foodborne pathogens, mathematical models can be used under specific conditions, in particular, the effects of storage temperatures on microbial growth [15-17]. Temperature is an important environmental factor for microbial growth, and changing temperature is a particularly important factor to consider when studying foodborne illnesses because circumstances change drastically during harvest, transport, and storage. A dynamic model developed by Baranyi and Roberts can predict bacterial growth under fluctuating temperature. With this mathematical concept, Koseki and Isobe developed mathematical models under changing temperatures during distribution to evaluate microbiological growth patterns on lettuce with pathogenic bacteria (*E. coli* O157:H7, *Salmonella* spp., and *Listeria monocytogenes*) [18,19]. Indeed, when Mckellar et al. simulated *E. coli* O157:H7 growth in fresh-cut lettuce under dynamic conditions, they were able to suggest the appropriate storage conditions that would effectively inhibit microbial growth under practical environment [20]. Also, Zeng et al. used the concept of mathematical model as an important method to compare bacterial growth (*Escherichia coli* O157:H7 and *Listeria monocytogenes*) in fresh-cut romaine mix at changing temperatures during transport, storage and display [14].

Therefore, the objective of this study was to describe the kinetic behavior of pathogenic strains of *E. coli* and *S. aureus* in lettuce, commercial sprout mix, and water celery with mathematical models at constant and changing temperature.

Materials and Methods

Inoculum preparation

Staphylococcus aureus strains (ATCC 13565, ATCC 14458, ATCC 23235, ATCC 27664, and NCCP 10826) and pathogenic *E. coli* strains [enterohemorrhagic *E. coli* (NCCP 11142), enterotoxigenic *E. coli* (NCCP 14037), enteropathogenic *E. coli* (NCCP 14038), enteroaggregative *E. coli* (NCCP 14039), and enteropathogenic *E. coli* (NCCP 15661)] were cultured in 10 mL tryptic soy broth (TSB; Becton, Dickinson and Company, Sparks, MD, USA) at 35 °C for 24 h. A small portion of the culture (0.1 mL) was transferred into 10 mL TSB and subcultured at 35 °C for 24 h. Five *S. aureus* strains or five *E. coli* strains were mixed separately, centrifuged at 1,912 X g to: 1,912 xg at 4 °C for 15 min, and each pellet was washed twice with phosphate-buffered saline (PBS, pH 7.4; 0.2 g of KH₂PO₄, 1.5 g of Na₂HPO₄·7H₂O, 8.0 g of NaCl, and 0.2 g of KCl in 1 L of distilled water). Each suspension was then serially diluted with PBS to 5 log CFU/mL.

Sample preparation and inoculation

Lettuce, commercial sprout mix (broccoli, rapeseed, red kohlrabi, and alfalfa) which is consumed as mixed, and water celery were purchased from a grocery store, cut into small portions (lettuce: 10 g, sprout: 5 g, water celery: 6 g) and placed into a filter bag (Sample Bag, 3M, Korea). The 0.1 mL suspensions were used to inoculate the prepared vegetable samples at 3 log CFU/g. The samples were massaged thoroughly and then stored aerobically at 4 °C, 10 °C, 15 °C, 25 °C, and 30 °C. Each inoculated sample was analysed at specific intervals during their incubation [96 h (15 °C, 25 °C, and 30 °C), 288 h (10 °C), and 336 h (4 °C)].

Bacterial growth analysis

After incubation, 60 mL (lettuce) or 40 mL (sprout mix and water celery) of 0.1% buffered peptone water (BPW; Becton, Dickinson and Company) was added into the sample bags and pummelled in a pummeller (BagMixer; Interscience, St. Nom, France) for 1 min, and the homogenates were then serially diluted with BPW. Aliquots (0.1 mL) of the diluents were spread-plated on tryptic soy agar (TSA; Becton, Dickinson and Company, Sparks, MD, USA), mannitol salt agar (MSA; Becton, Dickinson and Company, Sparks, MD, USA), and *E. coli*/ Coliform Count Plate (Petrifilm™, 3M, St. Paul, MN, USA) to enumerate total bacterial, *S. aureus*, and *E. coli* cell counts, respectively. The plates were then incubated at 35 °C for 24 h (total bacteria) and 48 h (*S. aureus* and *E. coli*), and bacterial colonies were counted manually. Five presumptive *S. aureus* colonies on MSA were identified with an agglutination test (Microgen® Staph Latex Agglutination Kit; Microgen Bioproducts Ltd., Surrey, UK), and the ratio of the number of positive colonies to the number of total colonies was used to determine the number of *S. aureus* colonies.

Calculation of kinetic parameters

The Baranyi model (A) was fitted to the *S. aureus* and *E. coli* cell counts with DMFit (Institute of Food Research, Norwich, UK) to calculate kinetic parameters such as lag phase duration (*LPD*; h) and maximum specific growth rate (μ_{max} ; log CFU/g/h) as well as lower asymptote (N_0 ; log CFU/g) and upper asymptote (N_{max} ; log CFU/g) for each storage temperature.

$$N_t = N_0 + \mu_{max} \times A_t - \ln\left[1 + \frac{\exp(\mu_{max} \times A_t) - 1}{\exp(N_{max} - N_0)}\right]$$

$$A_t = t + \frac{1}{\mu_{max}} \ln\left(\frac{\exp(-\mu_{max} \times t) + q_0}{1 + q_0}\right) \tag{A}$$

- N_t : Bacterial cell counts at time *t*
- N_0 : Initial bacterial counts
- μ_{max} : Maximum specific growth rate
- A_t : Adjustment function related to *LPD*
- N_{max} : Final bacterial counts
- q_0 : a Parameter defining the initial physiological state of the cells
- t*: Time

The square root model (B) was used to analyse the temperature effect on *E. coli* and *S. aureus* μ_{max} in lettuce to develop secondary models as a function of storage temperature (Baranyi and Roberts) [18].

$$\sqrt{\mu_{max}} = \alpha \times (T - T_{min}) \tag{B}$$

- μ_{max} : Maximum specific growth rate
- α : Slope of linear equation
- T*: Storage temperature
- T_{min} : Theoretical minimum growth temperature

Validation

To evaluate the model performance, additional experiments were conducted for each bacteria/vegetable at 12 °C and 20 °C which were not used in developing mathematical models, and the observed bacterial cell counts were compared to the predicted bacterial cell counts of developed models. The root mean square error (*RMSE*) (C), *A* factor (D) and *B* factor (E) were then calculated by comparing the observed values and predicted values as follows:

$$RMSE = \sqrt{\frac{\sum (\text{observed values} - \text{predicted values})^2}{n}} \tag{C}$$

$$A \text{ factor} = 10^{\sum |\log(\text{Predicted values}/\text{Observed values})|/n} \tag{D}$$

$$B \text{ factor} = 10^{\sum \log(\text{Predicted values}/\text{Observed values})/n} \tag{E}$$

Kinetic behavior at changing temperature

To predict *S. aureus* and *E. coli* cell counts under changing temperatures, the dynamic model was developed using the equation developed by Baranyi and Roberts [18]. The bacterial cell counts from lettuce, sprout mix and water celery were then simulated under changing temperature profiles, which were collected by measuring every 30 min from 10 am to 7:30 pm with an infrared light thermometer (HS33CT, Hansung, Seoul, Korea) in six grocery stores.

Statistical analysis

μ_{max} values were analysed using the general linear model

procedure of SAS® version 9.3 (SAS Institute, Cary, NC, USA). All mean comparisons were performed using a pairwise *t*-test at $\alpha = 0.05$.

Results

LPD values were observed for *E. coli* strains that were grown in lettuce but only at 4 °C (119.91 h), 25 °C (1.55 h) and 30 °C (1.30 h), and μ_{max} values for the pathogen increased ($P < 0.05$) as the storage temperature increased from 4 °C (0.002 log CFU/g/h) to 30 °C (0.289 log CFU/g/h) (Table 1). For *E. coli* in water celery, no growth was observed at 4 °C (Table 1). μ_{max} values for *E. coli* in water celery then increased ($P < 0.05$) when the storage temperature increased from 10 °C (0.002 log CFU/g/h) to 30 °C (0.325 log CFU/g/h), and very short LPD (0.64 h) were observed only at 30 °C (Table 1). However, no *E. coli* growth in the sprout mix was observed at all temperatures (Figure 1). Instead, the minimal death of *E. coli* was observed in sprout mix at 4 °C (Figure 1A).

To describe the changes to the kinetic growth parameters of the *E. coli* strains in response to increases in temperature, the square root model was fitted to μ_{max} values as a function of temperature, and R^2 values were 0.900-0.951, indicating that the square root model was appropriate to evaluate the temperature effect on μ_{max} (lettuce: = $0.0215 \times (T - 3.944)$, $R^2 = 0.900$; water celery: = $0.0224 \times (T - 6.790)$, $R^2 = 0.951$). From the square root model, T_{min} values, reflecting the theoretical minimum growth temperature for *E. coli* strains, can be calculated, and the values were 3.944 °C and 6.790 °C for lettuce and water celery, respectively.

To evaluate the performance of the mathematical models used in this study, an additional experiment was performed to obtain observed *E. coli* growth data at 12 °C and 20 °C. To obtain predicted *E. coli* cell counts, the kinetic parameters (μ_{max} and LPD) were

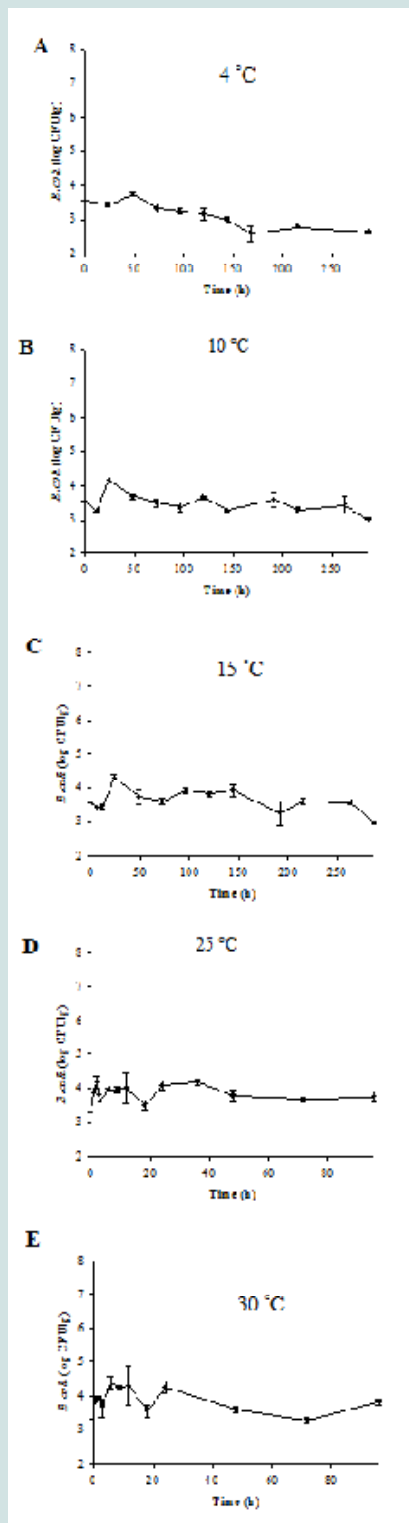


Figure 1: Bacterial populations of pathogenic *Escherichia coli* in sprout mix during storage at 4 °C (A), 10 °C (B), 15 °C (C), 25 °C (D) and 30 °C (E) for 288 h, 288 h, 288 h, 96 h and 96 h, respectively.

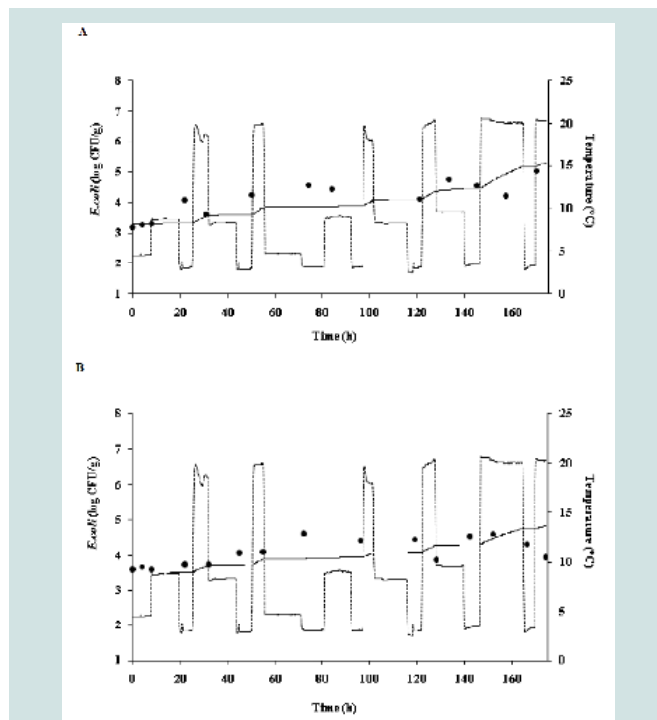


Figure 2: Predicted growth of pathogenic *Escherichia coli* in lettuce (A) and water celery (B) under dynamic temperature condition. Symbol: observed bacterial cell counts, line: predicted bacterial cell counts, dotted line: storage temperature.

Table 1: Kinetic growth parameters (mean ± SD) calculated by the Baranyi model for pathogenic *Escherichia coli* growth in lettuce and water celery.

	Storage temperature (°C)	LPD ¹⁾ (h)	$\mu_{max}^{2)}$ (log CFU/g/h)	$N_0^{3)}$ (log CFU/g)	$N_{max}^{4)}$ (log CFU/g)
Lettuce	4	119.91±169.58	0.002±0.003 ^B	3.3±0.2	3.8±0.2
	10	0.00±0.00	0.018±0.003 ^B	2.8±0.1	5.9±0.1
	15	0.00±0.00	0.016±0.003 ^B	3.6±0.0	6.0±0.1
	25	1.55±0.68	0.294±0.079 ^A	3.5±0.1	5.7±0.1
	30	1.30±1.11	0.289±0.081 ^A	3.5±0.1	6.1±0.0
Water celery	4	- ⁵⁾	-	-	-
	10	0.00±0.00	0.002±0.000 ^C	3.5±0.1	4.4±0.1
	15	0.00±0.00	0.014±0.005 ^C	3.5±0.1	5.2±0.2
	25	0.00±0.00	0.151±0.023 ^B	3.5±0.1	5.1±0.1
	30	0.64±0.90	0.325±0.026 ^A	3.5±0.0	5.4±0.1

- 1) lag phase duration
 - 2) maximum specific growth rate
 - 3) initial cell concentration
 - 4) maximum cell concentration
 - 5) no growth
- A-C:** different letters in a same column mean significantly different at P<0.05.

Table 2: Kinetic growth parameters (mean ± SD) calculated by the Baranyi model for *Staphylococcus aureus* growth in lettuce.

Storage temperature (°C)	LPD ¹⁾ (h)	$\mu_{max}^{2)}$ (log CFU/g/h)	$N_0^{3)}$ (log CFU/g)	$N_{max}^{4)}$ (log CFU/g)
4	64.68±25.81	0.003±0.000	4.6±0.0	5.6±0.0
10	20.11±0.27	0.005±0.003	4.8±0.1	6.4±0.6
15	0.00±0.00	0.006±0.001	5.1±0.0	6.7±0.0
25	0.00±0.00	0.023±0.015	4.2±0.4	6.8±0.7
30	0.00±0.00	0.024±0.001	4.4±0.1	8.9±0.0

- 1) lag phase duration
- 2) maximum specific growth rate
- 3) initial cell concentration
- 4) maximum cell concentration

calculated by substituting the specific temperature to the developed model with the square root model, and the kinetic parameters were substituted into the Baranyi model at given condition. The observed values were then compared to the predicted values, and *A* factor, *B* factor and *RMSE* values were then calculated. *A* factor values were 1.114 (lettuce) and 1.148 (water celery) and *B* factor values were 1.071 (lettuce) and 0.939 (water celery). *RMSE* values (lettuce: 0.691; water celery: 0.745) were interpreted that the differences between predicted values and observed values were less than 0.691 log CFU/g in lettuce and 0.745 log CFU/g in water celery.

To describe the kinetic behavior of *E. coli* during storage, which has changing temperature rather than constant temperature, a simulation of *E. coli* growth was conducted under changing temperature. The temperature was set to 4.2 °C, 21.2 °C, and 10.5 °C at an interval of 6 h to describe the changes of *E. coli* cell counts as the temperature changed. The temperature settings were chosen by calculating the minimum, maximum, and mean values for *E. coli* growth on lettuce (4.1 °C, 21.0 °C, and 10.5 °C) and water celery (4.2 °C, 24.4 °C, and 12.3 °C), which were measured at market displays from six separate locations. Simulated *E. coli* cell counts at changing temperature are presented in Figure 2. Under the changing storage temperatures, *E. coli* grew gradually in both vegetables, and *E. coli* growth was greater

in lettuce samples than when grown in water celery samples, which were also proven by N_{max} values in (Table 1 and Figure 2).

Increase in the cell counts of *S. aureus* that was grown on the sprout mix and water celery samples were minimal at all tested temperatures (Figures 3 and 4). Thus, kinetic parameters were calculated only for lettuce samples. Slow *S. aureus* growth in lettuce were observed at 4 °C (0.003 log CFU/g/h) - 30 °C (0.024 log CFU/g/h) (Table 2). *LPD* values were estimated for *S. aureus* in lettuce only at 4 °C (64.68 h) and 10 °C (20.11 h) (Table 2). The square root model was fitted to the μ_{max} values and appropriate fitting was obtained to describe temperature effect on the μ_{max} of *S. aureus* in lettuce (0.022×(T+6.077), R²= 0.947). T_{min} value of the model was -6.077°C, which is not realistic temperature for *S. aureus* growth.

The observed *S. aureus* cell counts obtained from lettuce stored at 12 °C and 20 °C were compared to the predicted *S. aureus* cell counts from the developed model. *A* factor and *B* factor values were 1.096 and 0.981, respectively. The *RMSE* value of *S. aureus* in lettuce was 0.569, indicating that the developed model is appropriate to describe the kinetic behaviour of *S. aureus* in lettuce. The growth pattern of *S. aureus* in lettuce was also simulated under changing temperature condition (4.2 °C, 21.2 °C, and 10.5 °C at 6 h intervals) to describe the fate of the pathogen, and the simulation showed only minimal increases of *S. aureus* cell counts in lettuce (Figure 5).

Discussion

T_{min} values of developed models were that 3.944 °C (*E. coli* in lettuce), -6.077°C (*E. coli* in water celery) and -6.077°C (*S. aureus* in lettuce). These results indicate that the T_{min} for the pathogenic *E. coli* is dependent on vegetable. Ratkowsky et al. previously suggested that the T_{min} values for *E. coli* O157:H7 grown on mechanically tenderized

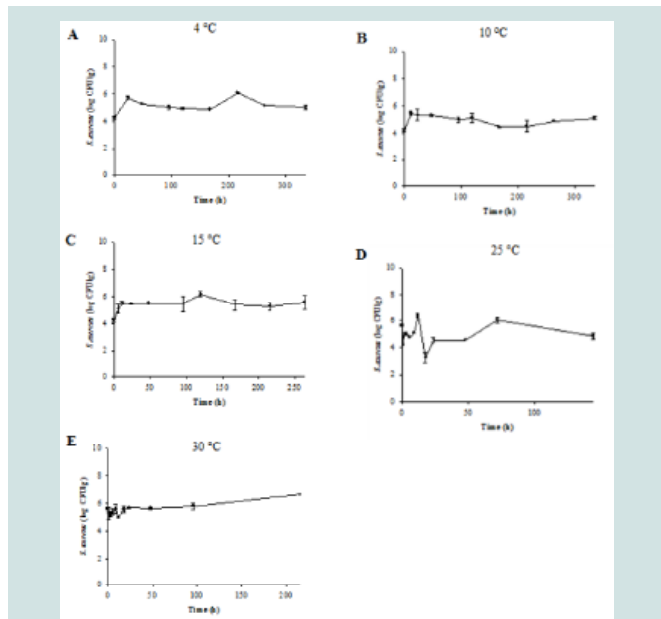


Figure 3: Bacterial populations of *Staphylococcus aureus* in sprout mix (broccoli, rapeseed, red kohlrabi, and alfalfa) during storage at 4 °C (A), 10 °C (B), 15 °C (C), 25 °C (D) and 30 °C (E) for 336 h, 336 h, 288 h, 144 h and 216 h, respectively.

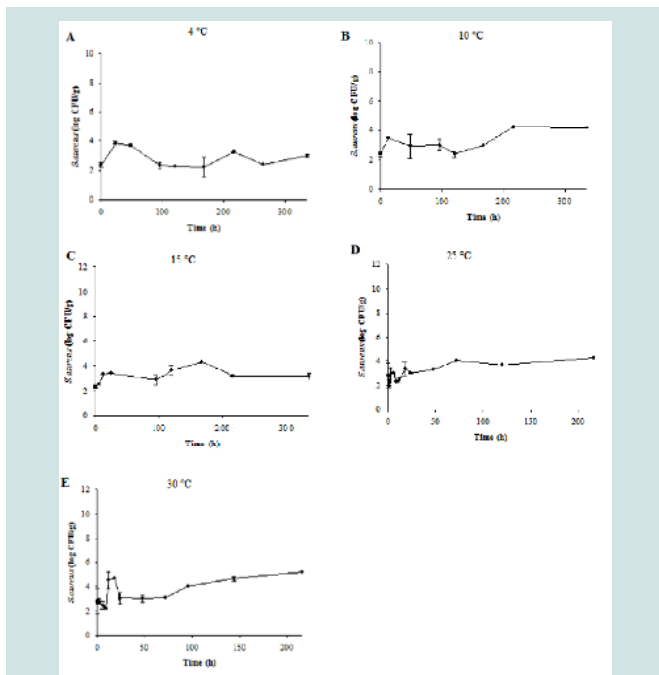


Figure 4: Bacterial populations of *Staphylococcus aureus* in water celery during storage at 4°C (A), 10 °C (B), 15 °C (C), 25 °C (D) and 30 °C (E) for 336 h, 336 h, 336 h, 216 h and 216 h, respectively.

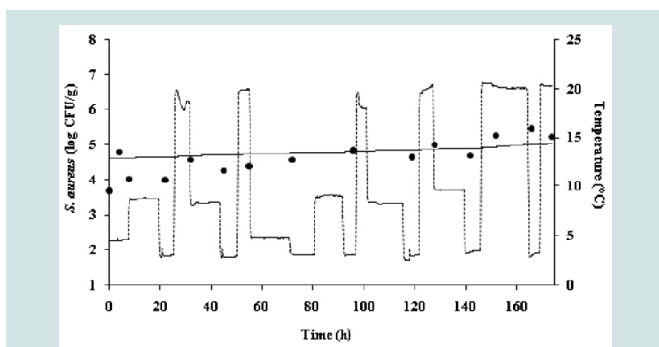


Figure 5: Predicted *Staphylococcus aureus* growth in lettuce under dynamic temperature condition. Symbol: observed bacterial cell counts, line: predicted bacterial cell counts, dotted line: storage temperature.

beef was 1.5 °C -4.7 °C, but Huang suggested that the T_{min} values were 6.64 °C -8.76 °C [21,22]. Taken together, T_{min} values for *E. coli* can be influenced by food matrix. Since T_{min} is the lowest temperature limit theoretically below which the calculated growth rate is close to zero [23], T_{min} and minimum growth temperature observed in food can be different such as T_{min} value of *S. aureus* in lettuce -6.077°C.

RMSE values were that 0.691 (*E. coli* in lettuce), 0.745 (*E. coli* in water celery) and 0.569 (*S. aureus* in lettuce). Because NACMCF suggested that more than 1 log growth could be generally considered as a significant change, a mathematical model with less than 1.0 of RMSE could be considered appropriate to describe the kinetic behavior of the pathogenic *E. coli* in lettuce and water celery at constant temperatures [24]. In previous studies, Perez-Rodriguez et al. and Lee et al. suggested that the performance of developed models

were appropriate to assay the temperature requirements for bacterial growth with similar RMSE values [0.300-0.450 for Perez-Rodriguez et al.; 0.326-0.361 for Lee et al. as those of our study [25,26]. Taken together, predictive models should be useful in describing kinetic behavior of foodborne pathogens [27,28].

In the dynamic model, although the *S. aureus* showed minimal growth in lettuce at changing temperature, the pathogen had very low risk of intoxication because *S. aureus* produce enterotoxin as they had growth up to 5-6 log CFU/g [29].

In conclusion, *E. coli* may not grow in commercial sprout mix, but they may grow in lettuce and water celery above at 10 °C as well as changing temperature found in grocery stores. Therefore, storage temperatures for lettuce and water celery should be below 10 °C to prevent *E. coli* growth on them. Regarding *S. aureus*, the pathogen may not grow in water celery and sprout mix, and only minimal growth in lettuce for both constant temperature and practical storage temperature, indicating that *S. aureus* may not be risky pathogen in the vegetables examined in this study.

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