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Changes in Membrane Fatty Acid Composition of *Pediococcus* sp. Strain NRRL B-2354 in Response to Growth Conditions and Its Effect on Thermal Resistance

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Membrane fatty acid composition and thermal resistance (*D* value) of *Pediococcus* sp. were determined for mid-exponential-phase (ME) and stationary-phase (ST) cells grown in tryptic soy broth (TSB) and tryptone-glucose-yeast extract (TGY) at 28 and 37°C. As the cells entered the stationary phase of growth, the unsaturated fatty acid, $C_{18:1\ n11c}$, produced during the exponential phase of growth was converted to its cyclic form, $C_{19:0\ \Delta 9c}$. This shift in membrane fatty acid composition was accompanied by an increase in the *D* values of this bacterium. Data from this study suggest that the membrane fatty acid composition of *Pediococcus* sp. is dependent on the growth conditions and that membrane fatty acid composition plays a critical role in thermal resistance. Thermal inactivation curves of *Pediococcus* sp. cells grown in TGY at 28°C indicated the presence of a cell population that is heterogeneous in thermal resistance. The growth of this bacterium in TGY at 37°C and in TSB at 28 and 37°C resulted in cell populations that were uniform in thermal resistance with a lag time for thermal inactivation. Thermal inactivation curves of ME and ST cultures were similar. The data presented here suggest that the cell population's uniformity of thermal inactivation is independent of the growth phase of the culture.

Bacterial cytoplasmic membrane, which consists mainly of lipids, has been shown to be a site for thermal injury (6, 8, 9, 32). This cytoplasmic membrane, the boundary between the cytoplasm and the external environment, regulates the flow of nutrients and metabolic products in and out of the cell, thereby permitting homeostasis of the cytoplasmic environment (3, 13). Growth conditions such as the composition of growth medium (1, 2, 12, 13, 17, 24), the growth phase (age) of the cells (13, 17, 18, 20, 32), the incubation temperature at which the bacteria were cultured (1, 6, 8, 13, 17, 18, 22–24, 26, 30), and the pH (3, 17, 23, 24) markedly affect the composition of the membrane lipid. These changes in the composition of the membrane lipid affect mainly the fluidity of the cellular membrane and are thought to occur in order to maintain both membrane integrity and functionality in the face of the external conditions (30). The major way in which bacteria maintain this ideal membrane fluidity is by changing their fatty acid composition (1, 22, 30). For example, as the growth temperature decreases, fatty acids with lower melting points are incorporated into the lipid bilayer, which lowers the order-disorder transition temperature of the membrane, thereby maintaining fluidity and compensating for the decreased temperature (1, 22, 30). This process has been described as a homeoviscous adaptation by Sinensky (26), a process in which the membrane fluidity is maintained (relatively) constant through lipid changes in response to changes in growth temperature.

Thermal inactivation of bacteria was shown to be dependent on environmental parameters such as the growth medium (2, 4, 9, 10, 27), the growth temperature (6, 8–10, 20, 21), the growth phase (10, 20), and the pH (2, 10). Also, changes in thermal resistance of bacteria due to changes in growth conditions were positively correlated with alterations in the membrane fluidity (6, 8, 9, 32, 33). These researchers reported that an increase in the fluidity of the bacterial membrane due to the changes in growth conditions corresponded to a decrease in thermal resistance. Furthermore, the use of procaine, a membrane fluidizing agent, in thermal inactivation studies resulted in an increase in the membrane fluidity, with a subsequent decrease in the thermal resistance of *Escherichia coli* (6, 33).

Pediococcus sp. (formerly, *Micrococcus freudenreichii* [2]) is a gram-positive, spherical, nonmotile, non-spore-forming, facultative anaerobe. This bacterium, originally isolated from milk and dairy utensils (28), is a heat-resistant, spoilage, nonpathogenic organism that has been used as a test organism in milk and milk byproduct pasteurization studies (14, 28, 29). These characteristics, i.e., nonpathogenicity and thermal resistance, have made this bacterium an attractive test organism for studying the destruction of bacteria by microwave energy in a food pilot plant (15, 16). Our data, obtained from a nonthermal, semicontinuous pilot plant process utilizing microwave energy, suggested that the lethal effect of electromagnetic energy on *Pediococcus* sp. was dependent on the growth medium and the processing fluid (15). Similarly, growth of this bacterium in different growth media resulted in different thermal resistance in all of the heating menstrua tested (2). This suggested that the thermal resistance of *Pediococcus* sp. was dependent on the growth medium. While this bacterium is considered to be a test organism in milk pasteurization studies, little is known about the influence of growth conditions on the membrane fatty acid composition and the thermal resistance of this organism.

This study is a part of ongoing research to develop a cold pasteurization process utilizing electromagnetic energy (15, 16). The data from our pilot plant microwave process indicated that cells of *Pediococcus* sp. grown on tryptone-glucose-yeast extract (TGY) were more sensitive to microwave energy than cells grown in tryptic soy broth (TSB) (15) and that bacterial

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inactivation by microwave energy was dependent on the growth temperature (unpublished data). Also, we previously reported that the thermal resistance of *Pediococcus* sp. was dependent on the growth medium (2). It was the objective of the present study to determine the influence of growth medium, growth phase (age), and growth temperature on membrane fatty acid composition and the thermal resistance of *Pediococcus* sp.

MATERIALS AND METHODS

Microorganism, culture maintenance, and growth media. *Pediococcus* sp. strain NRRL B-2354 was supplied by L. K. Nakamura (U.S. Department of Agriculture, Peoria, Ill.). The culture was maintained on tryptose agar (TA; Difco Laboratories, Detroit, Mich.) at 4°C, with biweekly transfers to maintain strain viability. The three growth media used were TSB (Difco Laboratories) prepared in distilled water according to the manufacturer's guidelines and supplemented with glucose to a final concentration of 0.5% (wt/vol), TGY broth formulated in our laboratory (tryptone, 5 g; yeast extract, 5 g; glucose, 1 g; dibasic potassium phosphate, 1 g; double-distilled and deionized water, 1 liter; pH 7.00), and TGYG broth (TGY broth supplemented with glucose to a final concentration of 0.5% [wt/vol]). All ingredients were mixed prior to autoclaving, and the medium pH did not change after autoclaving.

Inoculum development, growth conditions, sample preparation, and thermal inactivation. A late-exponential-phase culture grown in the appropriate medium at either 28 or 37°C was used to inoculate 50 to 100 ml of the same medium at a 1% level (vol/vol). Growth was monitored by measuring the optical density at 600 nm with a Shimadzu UV-160 spectrophotometer (Shimadzu Scientific Instruments, Inc., Columbia, Md.). Cultures were grown to the mid-exponential (A_{600} = 0.5) or stationary (A_{600} = 1.0) phase of growth, harvested by centrifugation at 16,000 × *g* for 10 min at 4°C, and washed once with cold sterile distilled water. The cell pellet was suspended in tap water as the heating menstruum to a target level of 8 log CFU/ml. Culture samples (9.5 ml) were loaded onto a Techne submerged-coil heating apparatus model tempette TE-8D (Protocol Instruments Limited, West Byfleet, United Kingdom) and kept at 60°C. Heating time and sampling frequency was based on the culture growth conditions. After being heated, the samples were quickly stored on ice.

Assessment of bacterial viability. The bacterial suspensions were serially diluted in 0.1% peptone (Difco) and surface plated on TA plates by using the spiral plating system, model D (Spiral Systems Instruments, Inc., Bethesda, Md.). The plates were then incubated at 37°C for 18 to 24 h, and the survivors were enumerated by using a laser bacterial colony counter, model 500A (Spiral Systems Instruments, Inc.). Cell densities were reported as CFU per milliliter of sample.

***D* values.** *D* values (the times needed in order to inactivate 90% of the population) were calculated as the negative reciprocal slope of the linear portion of survivor curves (which were obtained by plotting logarithms of survival counts versus their corresponding heating times). Linear regression lines were fitted to the linear portion of two sets of independent data.

Fatty acid analysis. The total fatty acids were extracted and methyl esterified from 40 to 80 mg (wet weight) of cell pellets as previously described (1). A Hewlett-Packard 5890 gas-liquid chromatograph (Hewlett-Packard, Avondale, Pa.) equipped with a split-splitless injector, flame ionization detector, integrator, and a 30-m-by-0.25-mm SPB-1 (0.25-μm film thickness) fused silica capillary column (Supelco, Inc., Bellefonte, Pa.) was used for the separation and detection of the fatty acid methyl esters (FAME). The carrier gas flow (helium) was adjusted to 24 cm/s, and the injector and detector temperatures were maintained at 250 and 280°C, respectively. The sample (3 to 5 μl) was injected in the split mode (ratio of 100:1), and the column temperature was held at 150°C for 4 min before it was raised to 250°C at a rate of 4°C/min. FAME were identified by comparing the retention times of a qualitative standard bacterial FAME mixture (Matreya, Inc., Pleasant Gap, Pa.).

All chemicals were analytical-grade reagents. Glassware was cleaned with Nochromix acid solution (Godax Laboratories, Inc., Takoma Park, Md.) and rinsed repeatedly with distilled water before use.

Statistical analysis. Standard errors were calculated from the regression analysis by using SAS software (SAS Institute, Inc., Cary, N.C.). Duncan's multiple-range test was used to determine the significant differences (unless otherwise mentioned, $P < 0.05$) among membrane lipid fatty acids and the *D* values of *Pediococcus* cells grown under different conditions.

RESULTS

Effect of growth conditions on membrane fatty acid composition. *Pediococcus* cells were grown to mid-exponential phase (ME) and stationary phase (ST) in TGY and TSB at 28 and 37°C, and their membrane fatty acid composition was determined. Total saturated fatty acids (SFA), total unsaturated fatty acids (USFA), and total cyclic fatty acids (CFA) were reported as the sum of the mean values of three independent replications ± the standard deviation (Tables 1 and 2) and were used in determining the significant differences among membrane fatty acids of *Pediococcus* cells grown under different conditions. The major fatty acids were, in order, USFA, SFA, and CFA, with USFA representing up to 72% of the total fatty acids (Tables 1 and 2).

The ST cells of *Pediococcus* sp. grown in TGY at 28°C showed a significant 4.3-fold increase in CFA compared to the ME cells grown in TGY at 28°C (Table 1). Growth of *Pediococcus* cells in TSB at 28°C to ST resulted in a 9.5-fold significant increase in CFA and a 36.5% significant decrease in USFA compared to ME cells grown in TSB at 28°C (Table 1). *Pediococcus* cells grown in TGYG at 28°C to ST had a membrane fatty acid profile similar to that of ST cells grown in TGY (Table 1).

Growth of *Pediococcus* cells at 37°C in TGY to ST resulted in a 1.4-fold significant (P = 0.07) increase in SFA (Table 2) compared to ST cells grown in TGY at 28°C (Table 1). When grown in TSB at 37°C, membrane fatty acid composition of ST cells (Table 2) was similar to that of ST cells grown at 28°C (Table 1). Also, the membrane fatty acid compositions of ME cells grown in TGY and TSB at 37°C (data not shown) was similar to those of ME cells grown in TGY and TSB at 28°C (Table 1).

Pediococcus cells grown in TSB at 28°C to ST exhibited a 1.9-fold significant increase in CFA and a 30.1% significant decrease in USFA compared to ST cells grown in TGY at 28°C (Table 1). When grown at 37°C, TSB-grown ST cells showed a 2.7-fold significant increase in CFA and a 24.8% significant decrease in USFA compared to TGY-grown ST cells (Table 2).

Effect of growth conditions on thermal resistance. ME and ST cultures grown in TGY and TSB at 28 and 37°C were used in studying the effect of growth conditions on the thermal resistance of *Pediococcus* sp. The thermal resistance (*D* value) at 60°C was determined in tap water since cells were shown to be stable in this heating menstruum (2). Logarithms of surviving *Pediococcus* cells (in CFU per milliliter) were plotted against heating time, and the *D* values were obtained by linear regression from the linear portion of the survivor curves (Fig. 1 and 2). The coefficient of correlation (r^2) range was 0.920 to 0.994. *D* values were reported as the mean of two independent replications ± the standard deviation (Table 3) and were used to determine the significance of growth conditions on the thermal resistance of *Pediococcus* sp.

A representative example of the effect of growth medium on survivor curves at 60°C is shown in Fig. 1 and 2, where the thermal inactivation curves demonstrated a biphasic inactivation characterized by a shoulder and/or tailing. While ME cells grown in TSB at 28°C exhibited a lag time for thermal inactivation (Fig. 1) of 0.25 min (Table 3), TGY-grown ME cells exhibited an initial period of higher thermal sensitivity (Fig. 2). Survivor curves of ST cells grown in TSB and TGY at 28°C (data not shown) were similar to those of ME cells grown in TSB (Fig. 1) and TGY (Fig. 2) at 28°C, with TSB-grown ST cells exhibiting a 1.00-min lag time for thermal inactivation (Table 3). Growth of *Pediococcus* cells in TGY and TSB at 37°C to ST resulted in survivor curves (data not shown) similar to that of ME cells grown in TSB at 28°C (Fig. 1), with lag times for thermal inactivation of 0.25 and 2.00 min, respectively (Table 3). When compared to ST cells grown in TGY at 28°C, the growth of *Pediococcus* cells in TGYG at 28°C to ST cells resulted in cell population with uniform thermal resistance (data not shown), a result similar to that seen with ME

TABLE 1. Effect of growth medium[a] on the membrane fatty acid[b] composition of *Pediococcus* cells grown at 28°C until ST or ME

Acyl chain[c]	% Total fatty acid on:				
	TGY		TSB		TGYG
	ST	ME	ST	ME	ST
SFA					
$C_{13:0}$	0.09	0.12	ND	0.14	0.08
$C_{14:0}$	5.75	5.41	8.59	4.41	2.54
$C_{15:0}$	0.62	0.99	0.12	0.36	1.03
$C_{16:0}$	12.93	16.30	16.15	17.56	11.81
$C_{17:0}$	ND[d]	ND	ND	0.11	0.30
$C_{18:0}$	1.06	1.62	1.11	2.34	2.33
$C_{20:0}$	0.31	0.29	0.45	0.51	1.04
Total[e]	20.76 ± 3.64	24.73 ± 2.28	26.41 ± 1.47	25.42 ± 4.44	19.13 ± 4.38
USFA					
$C_{16:1\ n9c}$	18.71	16.20	16.95	11.47	11.31
$C_{18:2\ n9c,\ n12c}$	0.21	ND	0.92	0.19	ND
$C_{18:1\ n9c}$	ND	0.25	ND	0.39	ND
$C_{18:1\ n11c}$	46.24	55.51	27.70	59.72	57.78
Total[e]	65.16 ± 0.24	71.97 ± 1.21	45.56 ± 3.00	71.76 ± 3.74	69.09 ± 14.10
CFA					
$C_{17:0\ \Delta 9c}$	0.40	0.07	2.25	0.09	0.11
$C_{19:0\ \Delta 9c}$	13.69	3.24	24.69	2.73	11.15
Total[e]	14.10 ± 3.42	3.31 ± 1.09	26.93 ± 2.73	2.83 ± 0.74	11.26 ± 8.93
Others	ND	ND	1.09	ND	0.57

[a] Cells were grown in 100 ml of TGY, TSB, or TGYG medium at 28°C until ST (A_{600} = 1.0) or ME (A_{600} = 0.5), centrifuged, and washed once with sterile distilled water. Wet cells (40 to 80 mg) were used in the preparation and analysis of FAME as described in Materials and Methods.
[b] Data are presented as percent total fatty acid and represent the mean of three replications.
[c] *n*, location of double bond; Δ, location of cyclopropane ring; c, *cis*.
[d] ND, not detected (species absent or below detection limit).
[e] Data are the total of the mean values ± the standard deviation of replicate values.

cells grown in TSB at 28°C (Fig. 1), but no lag time for inactivation was detected (Table 3).

Growth of *Pediococcus* cells at 28°C to ST in TGY and TSB resulted in 2.1- and 6.3-fold significant increases in *D* values, respectively, compared to ME cells grown in the same medium (Table 3). ST cells grown at 37°C in TGY showed a 2.6-fold significant increase in *D* value compared to ST cells grown in TGY at 28°C (Table 3).

The growth of *Pediococcus* cells in TSB at 28 and 37°C to ST resulted in 4.1- and 1.9-fold significant increases in *D* values, respectively, compared to ST cells grown in TGY at 28 and 37°C (Table 3).

DISCUSSION

Fatty acids play an important role in determining the physiochemical properties of cellular and membrane lipids. Microorganisms possess species-specific fatty acid profiles (30), and the membrane fatty acid composition of *Pediococcus* cultures (Tables 1 and 2) was consistent with that of the genus *Pediococcus* (19). Fatty acids derived from ST cultures grown in TGY and TSB at different temperatures revealed a marked shift in membrane fatty acid composition as cells entered the ST of growth, where a significant proportion of the $C_{18:1\ n11c}$ in ME cells was converted to $C_{19:0\ \Delta 9c}$ in ST cells (Table 1). The formation of CFA is considered to be a postsynthetic modification of the phospholipid bilayer that occurs predominantly as cultures enter the ST (3, 5, 24, 31). The CFA are formed by CFA synthase through the addition of a methylene group from *S*-adenosyl-L-methionine to the *cis* double bond of the USFA moiety of the phospholipid. The conversion of USFA to CFA as the cells enter the ST is believed to serve as a protective measure against lipid oxidation (18), low pH (3), and thermal inactivation (32). On the other hand, bacterial mutant strains that completely lacked the ability to synthesize CFA were able to grow and survive normally under virtually all conditions (31). The survival of these mutant strains was somewhat reduced after repeated cycles of freezing and thawing, indicating that this fatty acid modification is not essential but may be beneficial under certain conditions (31).

The most commonly found CFA in the bacterial membrane are $C_{19:0\ \Delta 11c}$, which is derived from $C_{18:1\ n11c}$; $C_{17:0\ \Delta 9c}$, which is derived from $C_{16:1\ n9c}$; and $C_{19:0\ \Delta 9c}$, which is derived from $C_{18:1\ n9c}$ (24). While the conversion of $C_{18:1\ n11c}$ to $C_{19:0\ \Delta 11c}$ was reported in ST cells of *E. coli* (18, 24, 31), Brown et al. (3) and Yatvin et al. (32) did not indicate the location of the cyclopropane ring on the cyclic fatty acid $C_{19:0}$ that was derived from $C_{18:1\ n11c}$. The data reported here indicate that the $C_{19:0\ \Delta 9c}$ in the *Pediococcus* cytoplasmic membrane was derived from $C_{18:1\ n11c}$ (Tables 1 and 2) rather than from $C_{18:1\ n9c}$. To check the validity of this data, we added the known FAME $C_{18:1\ n11c}$, $C_{18:1\ n9c}$, $C_{19:0\ \Delta 11c}$, or $C_{19:0\ \Delta 9c}$ to the membrane FAME samples and analyzed them by gas-liquid chromatography. The results indicated that $C_{19:0\ \Delta 9c}$ fatty acid was indeed derived from $C_{18:1\ n11c}$. This indicates that unlike other microorganisms, *Pediococcus* sp. might contain an enzyme system that is capable of carrying on the conversion of $C_{18:1\ n11c}$ to $C_{19:0\ \Delta 9c}$ instead of $C_{19:0\ \Delta 11c}$.

The experiments described here focus attention on $C_{19:0\ \Delta 9c}$ as playing a critical role in thermal resistance of *Pediococcus* sp. This fatty acid becomes one of the major fatty acids as cultures enter ST irrespective of the growth medium and the growth temperature (Tables 1 and 2). The phase transition

TABLE 2. Membrane fatty acid composition[a] of *Pediococcus* cells[b] grown at 37°C in TGY or TSB medium until ST

Acyl chain[c]	% Total fatty acid on:		T_c[d] (°C)
	TGY	TSB	
SFA			
$C_{13:0}$	ND[e]	0.05	13.5
$C_{14:0}$	7.14	8.99	23.0
$C_{15:0}$	0.56	0.13	34.2
$C_{16:0}$	18.98	17.50	41.0
$C_{17:0}$	ND	ND	47.8
$C_{18:0}$	2.37	1.49	58.0
$C_{20:0}$	0.33	0.19	66.0
Total[f]	29.38 ± 8.70	28.35 ± 7.79	
USFA			
$C_{16:1\ n9c}$	15.56	18.02	−36.0
$C_{18:2\ n9c,\ n12c}$	0.39	0.16	−17.6
$C_{18:1\ n9c}$	4.35	2.90	−22.0
$C_{18:1\ n11c}$	40.91	24.97	−19.0
Total[f]	61.22 ± 12.45	46.05 ± 7.37	
CFA			
$C_{17:0\ \Delta 9c}$	0.26	1.88	−19.9
$C_{19:0\ \Delta 9c}$	9.15	23.60	−0.5
Total[f]	9.41 ± 5.17	25.48 ± 4.43	
Others	ND	0.13	

[a] Data represent the mean of three replications.

[b] Cells were grown in 100 ml of TGY or TSB medium at 37°C until ST (A_{600} = 1.0), centrifuged, and washed once with sterile distilled water. Wet cells (40 to 80 mg) were used in the preparation and analysis of FAME as described in Materials and Methods.

[c] *n*, location of double bond; Δ, location of cyclopropane ring; c, *cis*.

[d] T_c, phospholipid gel to liquid-crystalline-phase transition temperature (25).

[e] ND, not detected (species absent or below detection limit).

[f] Data are the total of the mean values ± the standard deviation of replicate values.

temperature (T_c) of phosphatidylcholine containing $C_{19:0\ \Delta 9c}$ (−0.5°C) is significantly higher than that of phosphatidylcholine containing $C_{18:1\ n11c}$ (−19°C) (Table 2). Also, $C_{19:0\ \Delta 9c}$, which has less rotational freedom than $C_{18:1\ n11c}$, imparts increased rigidity to the cytoplasmic membrane (7, 32). Thus, the increase in CFA and the decrease in USFA would cause a decrease in membrane fluidity, thereby increasing the thermal resistance (6, 8, 32).

Pediococcus cells grown in TGY and TSB at 28°C to ST showed a significant increase in CFA, a significant decrease in USFA (Table 2), and a significant increase in *D* values (Table 3) compared to the ME cells. These data suggest that ST cells are more thermotolerant than ME cells (11, 20, 32). The increase in CFA within membrane fatty acids increases the membrane rigidity and hence decreases the fluidity of the cell membrane (3, 7, 18, 20, 32). This decrease in membrane fluidity could explain the increase in the thermal resistance of these ST cultures (Table 3). A similar correlation between thermal resistance (6, 8, 32) or acid resistance (3) and the CFA content of cytoplasmic membrane fluidity was reported for *E. coli*.

Growth of *Pediococcus* cells in TSB to ST resulted in a significant increase in CFA and a significant decrease in USFA compared to ST cultures grown in TGY at 28°C (Table 1) and 37°C (Table 2). The ST cells of *Pediococcus* sp. grown in TSB at 28 and 37°C had significantly higher *D* values and exhibited up to a 1.75-min increase in lag time for thermal inactivation compared to TGY-grown ST cells (Table 3). The increase in membrane rigidity as a result of producing more CFA and less USFA could explain the increase in the *D* values of TSB-grown

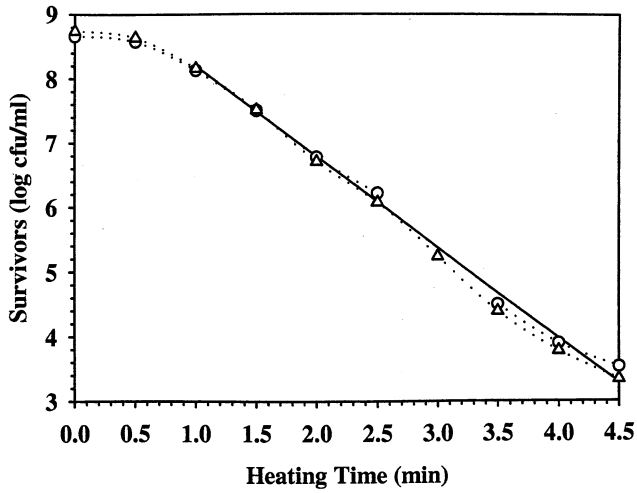

FIG. 1. Thermal inactivation curve of TSB-grown *Pediococcus* cells in tap water at 60°C. Circles and triangles represent the data of two independent studies. The solid straight line is the average regression plot of the straight portion of the two survivor curves (dotted lines). Cells were grown to ME at 28°C, washed once with sterile distilled water, and suspended in tap water to ca. 8 log CFU/ml.

ST cells compared to the TGY-grown ST cells. While the total levels of SFA, USFA, and CFA of ME cells grown in TGY and TSB at 28°C were similar (Table 1), TSB-grown ME cells seemed to contain higher concentrations of the individual fatty acids with higher melting points (see Table 2 for T_c values) compared to TGY-grown ME cells. Also, TSB-grown ME cells possessed a 1.4-fold higher *D* value and a 0.25-min increase in lag time for thermal inactivation compared to ME cells grown in TGY at 28°C (Table 3). Thus, the change in membrane fluidity due to higher concentrations of fatty acids with higher melting points could explain the change in the thermal resistance of this bacterium. These data indicate that the TSB-grown cells were more resistant to thermal inactivation than cells grown in TGY. These differences in *D* values (Table 3)

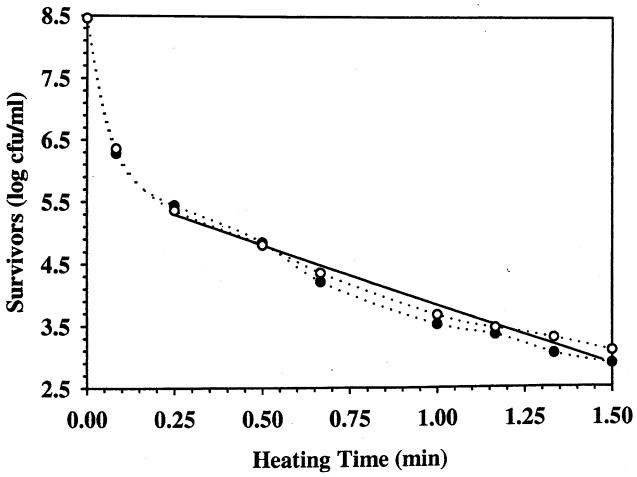

FIG. 2. Thermal inactivation curve of TGY-grown *Pediococcus* cells in tap water at 60°C. Circles and triangles represent the data of two independent studies. The solid straight line is the average regression plot of the straight portion of the two survivor curves (dotted lines). Cells were grown to ME at 28°C, washed once with sterile distilled water, and suspended in tap water to ca. 8 log CFU/ml.

TABLE 3. Effect of growth conditions on *D* values of *Pediococcus* sp. at 60°C

Culture growth conditions[a]	*D* value (min)[b]	Lag time (min)[c]	pH at end of growth[d]
TGY, ME, 28°C	0.51 ± 0.04	ND[e]	6.34
TGY, ST, 28°C	1.09 ± 0.06	ND	5.15
TGY, ST, 37°C	2.81 ± 0.38	0.25	4.86
TSB, ME, 28°C	0.71 ± 0.02	0.25	6.17
TSB, ST, 28°C	4.46 ± 0.06	1.00	5.28
TSB, ST, 37°C	5.28 ± 1.75	2.00	4.68
TGYG, ST, 28°C	0.70 ± 0.02	ND	4.28

[a] Cells were grown in the appropriate medium (TGY, TSB, or TGYG) until ME or ST at the appropriate temperature (28 or 37°C), centrifuged, washed once with sterile distilled water, and resuspended in tap water to a final cell concentration of ca. 8 log CFU/ml.

[b] Data represent the mean values of two replications ± the standard deviation of replicate values.

[c] Initial period for thermal resistance.

[d] Medium pH was measured prior to harvesting of the cells.

[e] ND, not detected.

and initial thermal protection (Fig. 1) or sensitivity (Fig. 2) appear to result from effects of the growth medium regardless of the growth phase used. A similar report (2) suggested that the changes in *D* values and the initial thermal protection or sensitivity of *Pediococcus* ST cells seemed to be an effect of the growth medium regardless of the heating menstruum used. This effect could be due to the higher concentrations of glucose and/or the nitrogen source in TSB medium compared to that of TGY medium. Concentrations of glucose and/or the nitrogen source in the growth medium have been known to influence the bacterial cell membrane fluidity (1, 2, 12, 17), where an increase in membrane fluidity was reported to result in a decrease in the thermal resistance of the bacterial cell (3, 6, 8, 9, 32, 33).

The *D* values of TSB (containing 0.5% glucose)-grown cells of *Pediococcus* sp. were significantly higher than those of TGY (containing 0.1% glucose)-grown cells in all heating menstrua tested (2). These results, in conjunction with the data presented here (see above), suggest that the increase in thermal resistance of *Pediococcus* cells is a result of an increasing glucose concentration in the growth medium. To check the validity of this hypothesis, we studied the effect of glucose on thermal inactivation of *Pediococcus* sp. Cells were grown in TGYG (TGY broth containing 0.5% glucose) to ST at 28°C, and the *D* values were determined in tap water at 60°C. There was a 36% decrease in the *D* value of ST cells grown in TGYG compared to TGY-grown ST cells (Table 3). These data suggest that the increase in glucose concentration in the growth medium had a negative effect on the thermal resistance of this bacterium. Smith et al. (27) reported that the presence of glucose in the growth medium potentiated the heat injury of *Staphylococcus aureus*. When ST *Pediococcus* cells were grown in TGY or TGYG at 28°C, the pH values of the growth media at the end of growth were 5.15 and 4.28, respectively. Thus, the decrease in thermal resistance as a result of increasing the glucose concentration in the growth medium might be related to the pH of the medium at the end of growth. The data presented in Table 3 show that the increase in *D* values of cells grown in TGY and TSB at 37°C was associated with a decrease in medium pH at the end of growth compared to growth at 28°C. However, data presented in Table 3 indicated that ST cells of *Pediococcus* sp. grown in TGYG at 28°C had a low pH (4.28), yet the cells were more sensitive to thermal inactivation than the TGY-grown ST cells (Table 3). Thus, the decrease in pH was merely indicative of glucose metabolism (27).

ST cells of *Pediococcus* sp. grown in TGYG medium at 28°C showed a 31% decrease in CFA (Table 1) and a 36% decrease in the *D* value (Table 3) compared to ST cells grown in TGY. This decrease in the CFA could increase the membrane fluidity and thus explain the decrease in the thermal resistance of this bacterium (6, 8, 32). However, the presence of 0.5% glucose in the TSB medium resulted in a decreased membrane fluidity in ST cells grown at 28°C (Table 1), with a concomitant increase in the *D* value (Table 3) compared to TGYG-grown ST cells. This suggests that nutrient sources other than glucose present in TSB compared to TGY probably affected the membrane fluidity. Since the composition of growth medium was reported to affect the bacterial membrane fluidity through the alteration of fatty acid content (see above), determination of thermal resistance would ultimately depend on growth medium.

We sought to investigate the effect of growth temperature on membrane fatty acid composition and thermal resistance of this bacterium. ME cultures of *Pediococcus* sp. grown in TGY and TSB at 28°C (Table 1) had similar fatty acid profiles compared to ME cultures grown at 37°C (data not shown). Also, ST cells grown in TSB at 37°C (Table 2) showed a similar fatty acid profile (slight increase in SFA) compared to those of ST cells grown at 28°C (Table 1), a ca. 1.2-fold increase in *D* value, and a 1-min increase in lag time for thermal inactivation (Table 3). When grown in TGY at 37°C (Table 2), ST cells showed a significant increase in SFA compared to ST cells grown at 28°C (Table 1), a significant increase in *D* value (ca. 2.6-fold), and a 0.25-min increase in lag time for thermal inactivation (Table 3). This suggests that cyclization of fatty acid does not play a role in temperature adaptation (18, 22). On the other hand, the increase in SFA, which has a significantly higher T_c (Table 2) than USFA, can result in lower membrane fluidity and hence can explain the increase in the *D* values of *Pediococcus* sp. in response to the increase in growth temperature.

The increase in growth temperature of *Pediococcus* sp. revealed that the mode of adaptation of fatty acid composition was dependent on the growth medium (Tables 1 and 2). The activity of CFA synthase responsible for synthesis of $C_{19:0\ \Delta 9c}$ fatty acid from the $C_{18:1\ n11c}$ intermediate was possibly produced at lower levels when cells were grown in TGY compared to TSB-grown cells (Tables 1 and 2).

The biphasic nature of the survivor curves suggests that two discrete populations were present (2, 10, 11). The survivor curves of *Pediococcus* sp. grown in TSB at 28°C to ME exhibit a lag time for thermal inactivation and a linear decrease in cell concentration during thermal inactivation (Fig. 1). This suggests that the cell population is uniform in its thermal resistance (2, 10, 11). On the other hand, ME cells grown in TGY exhibited an initial period of higher thermal sensitivity (Fig. 2), suggesting the presence of a cell population heterogeneous in thermal resistance (2, 10, 11). Similar survivor curves were reported for *Pediococcus* cultures grown in TGY and TSB at 28°C to ST (2), suggesting that the cell population's uniformity in thermal resistance is independent of the culture growth phase. Similar biphasic thermal inactivation kinetics in *Salmonella enteritidis* PT4 have been reported to be independent of culture age (11). When *Pediococcus* cultures were grown in TGY and TSB at 37°C to ST, cell populations were uniform in thermal resistance (data not shown), as previously seen with ME cells grown in TSB at 28°C (Fig. 1), with a lag time for thermal inactivation (Table 3). When compared to ST cells grown in TGY at 28°C, the ST cell population of *Pediococcus* sp. grown in TGYG at 28°C was uniform in thermal resistance

(data not shown), with no lag time for thermal inactivation (Table 3). Thus, the data suggest that the thermal inactivation curves of *Pediococcus* cell populations are dependent on the growth medium and growth temperature and not on the culture growth phase.

In conclusion, thermal resistance of *Pediococcus* sp. was shown here to be dependent on many factors, including growth medium, growth temperature, and growth phase (see above), as well as the heating menstruum (2). These growth conditions, as well as the methodology used for bacterial recovery, could make it difficult to compare *D* values obtained in different laboratories. This study is important because it shows that until a universal growth medium suitable for *D* value determination is developed, the interpretation and prediction of the bacterial thermal resistance in foods from data obtained under one growth condition and/or one experimental procedure are not advisable.

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