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Diversity and Mechanisms of Alkali Tolerance in *Lactobacilli*[†]

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We determined the maximum pH that allows growth (pH_{max}) for 34 strains of *Lactobacilli*. High alkali tolerance was exhibited by strains of *Lactobacillus casei*, *L. paracasei* subsp. *tolerans*, *L. paracasei* subsp. *paracasei*, *L. curvatus*, *L. pentosus*, and *L. plantarum* that originated from plant material, with pH_{max} values between 8.5 and 8.9. Among these, *L. casei* NRIC 1917 and *L. paracasei* subsp. *tolerans* NRIC 1940 showed the highest pH_{max}, at 8.9. Digestive tract isolates of *L. gasseri*, *L. johnsonii*, *L. reuteri*, *L. salivarius* subsp. *salicinius*, and *L. salivarius* subsp. *salivarius* exhibited moderate alkali tolerance, with pH_{max} values between 8.1 and 8.5. Dairy isolates of *L. delbrueckii* subsp. *bulgaricus*, *L. delbrueckii* subsp. *lactis*, and *L. helveticus* exhibited no alkali tolerance, with pH_{max} values between 6.7 and 7.1. Measurement of the internal pH of representative strains revealed the formation of transmembrane proton gradients (Δ pH) in a reversed direction (i.e., acidic interior) at alkaline external-pH ranges, regardless of their degrees of alkali tolerance. Thus, the reversed Δ pH did not determine alkali tolerance diversity. However, the Δ pH contributed to alkali tolerance, as the pH_{max} values of several strains decreased with the addition of nigericin, which dissipates Δ pH. Although neutral external-pH values resulted in the highest glycolysis activity in the presence of nigericin regardless of alkali tolerance, substantial glucose utilization was still detected in the alkali-tolerant strains, even in a pH range of between 8.0 and 8.5, at which the remaining strains lost most activity. Therefore, the alkali tolerance of glycolysis reactions contributes greatly to the determination of alkali tolerance diversity.

Lactic acid bacteria (LAB) produce lactic acid as a main product of sugar fermentation. Many studies have examined the response of LAB to acid stress, including the mechanisms underlying acid tolerance (3, 4, 28). In contrast, very few studies have examined the stress response of LAB to alkaline environments. Several alkali-tolerant/alkaliphilic LAB, which are capable of growing at pH values of >9.5 as well as producing large amounts of lactic acid from glucose, have been reported to date: *Enterococcus hirae* (formerly *Streptococcus faecalis*) (8), *Pediococcus urinaeequi* (23), *Alkalibacterium olivapovlenticus* (12, 24), *Marinilactibacillus psychrotolerans* (12), *M. piezotolerans* (27), *Halolactibacillus halophilus*, and *H. miuensis* (13).

The mechanism underlying the alkali tolerance of LAB has been examined in *E. hirae* ATCC 9790^T. Biochemical experiments and mutant analyses suggest that *E. hirae* ATCC 9790^T maintains an internal pH lower than the external pH by the operation of an ATP-driven K⁺/H⁺ antiporter during growth at an external pH of around 9.5 (14–17). However, *E. hirae* ATCC 9790^T has no such system to maintain a near-neutral internal pH in an alkaline medium (22). Thus, the contribution of internal-pH maintenance to alkali tolerance in this bacterium is controversial. Adaptation to alkaline stress has been reported for *E. faecalis* ATCC 19433^T, in which the induction of two stress proteins, DnaK and GroEL, was detected (9).

Lactobacilli, although generally not considered alkali tolerant, have been used as a starter for the fermentation of alkaline

materials to prepare fermented foods. For example, *Lactobacillus pentosus* CECT 5138 has been studied as a starter to ferment lye-treated green olives (pH, >9.0) for the preparation of Spanish-style green olives (25). In our previous study (26), in which noodle dough with a pH of around 8.5 was fermented, *L. plantarum* NRIC 0380 was screened as a starter strain for fermented instant Chinese noodles.

Because there has been little examination of alkali tolerance and sensitivity in *Lactobacillus* strains, we investigated the effects of alkaline stress on the growth, physiology, and metabolism of *Lactobacilli* from various sources for more effective use of this group of bacteria in food industries.

MATERIALS AND METHODS

Bacterial strains and chemicals. The bacterial strains used (Table 1) were obtained from the Japan Collection of Microorganisms (JCM, Wako, Japan), the Culture Collection Center, Tokyo University of Agriculture (NRIC, Tokyo, Japan); the Laboratory of Applied Microbiology, Research Faculty of Agriculture, Hokkaido University (AHU, Sapporo, Japan); and the Wageningen Center for Food Sciences (Wageningen, The Netherlands). [(2-Hydroxy-1,1-bis(hydroxymethyl)ethyl)amino]-1-propanesulfonic acid; N-[Tris(hydroxymethyl)methyl]-3-aminopropanesulfonic acid (TAPS); N-Tris(hydroxymethyl)methyl-4-aminobutanesulfonic acid (TABS); 2-(cyclohexylamino)ethanesulfonic acid (CHES); and nigericin were obtained from Sigma-Aldrich (St. Louis, MO). 2-(Morpholino)ethanesulfonic acid monohydrate (MES) and 2-[4-(2-hydroxyethyl)-1-piperazinyl]ethanesulfonic acid (HEPES) were obtained from Dojindo Laboratories (Mashikimachi, Kumamoto, Japan). 3-(N-morpholino)propanesulfonic acid (MOPS) was purchased from Nacalai Tesque (Kyoto, Japan). The 5 (and 6)-carboxyfluorescein diacetate succinimidyl ester (cFDA) was obtained from Molecular Probes (Eugene, OR). Other chemicals used were of the highest quality available commercially.

Determination of maximum alkaline pH allowing growth of *Lactobacilli*. For the preparation of seed cultures, frozen stocks of *Lactobacillus* strains and *E. hirae* JCM 8729^T kept under –80°C were transferred to 3 ml of half-strength MRS (1/2 MRS) medium (6) and incubated overnight at 30°C or 37°C, as indicated in the legend to Table 1. All cultures were grown in screw-cap test tubes tightly sealed with rubber septums. No other anaerobic treatment was

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TABLE 1. Determination of the maximum pH that allows the growth of various *Lactobacillus* strains and *E. hirae* JCM 8729^{Ta}

Species	Strain	Source	pH range allowing growth							
			6.5	6.9	7.3	7.7	8.1	8.5	8.9	9.3
<i>L. casei</i>	JCM 1134 ^T	Cheese								
	NRIC 1917	Sugar cane wine								
<i>L. paracasei</i> subsp. <i>tolerans</i> *	JCM 1171 ^T	Pasteurized milk								
	NRIC 1940	Coconut juice								
<i>L. paracasei</i> subsp. <i>paracasei</i> *	JCM 8130 ^T	Milk products								
	NRIC 1946	Coconut juice								
<i>L. curvatus</i>	JCM 1096 ^T	Milk								
	AHU GD12	“Ragi tape” (starter for sake)								
<i>L. pentosus</i>	JCM 1558 ^T	Corn silage								
	NRIC 0413	Fermented tea leaves								
<i>L. plantarum</i>	JCM 1149 ^T	Pickled cabbage								
	WCFS1	Human saliva								
	NRIC 0380	Pickles								
<i>L. salivarius</i> subsp. <i>salicinius</i>	JCM 1150 ^T	Saliva								
<i>L. salivarius</i> subsp. <i>salivarius</i>	JCM 1040	Human intestine								
	JCM 1044	Human intestine								
	JCM 1231 ^T	Human saliva								
<i>L. gasseri</i>	JCM 1131 ^T	Human intestine								
	JCM 5813	Infant feces								
<i>L. johnsonii</i>	JCM 1022	Human intestine								
	JCM 2012 ^T	Human blood								
<i>L. sakei</i> subsp. <i>sakei</i> *	JCM 1157 ^T	“Moto” (starter for sake)								
<i>L. reuteri</i>	JCM 1112 ^T	Adult intestine								
<i>L. brevis</i> *	NRIC 0137	Fermented bamboo shoots								
<i>L. delbrueckii</i> subsp. <i>delbrueckii</i>	JCM 1012 ^T	Sour grain mash								
<i>L. fermentum</i>	JCM 1173 ^T	Fermented beets								
	NRIC 0131	Pickled mustard								
<i>L. mali</i>	JCM 1116 ^T	Apple juice from cider press								
<i>L. delbrueckii</i> subsp. <i>bulgaricus</i>	JCM 1001	Not described								
	JCM 1002 ^T	Bulgarian yogurt								
<i>L. delbrueckii</i> subsp. <i>lactis</i>	JCM 1248 ^T	Emmental (Swiss) cheese								
<i>L. helveticus</i>	JCM 1062	Dairy products								
	JCM 1120 ^T	Emmental (Swiss) cheese								
<i>L. acidophilus</i>	JCM 1132 ^T	Human feces								
<i>E. hirae</i>	JCM 8729 ^T	Not described								

^a Gray bars indicate the pH range allowing growth for each strain. Cultures were grown in buffered 1/2 MRS media at 37°C, except for species denoted with asterisks, which were cultured at 30°C. Buffers used in the buffered 1/2 MRS medium included MOPS (#), HEPES (%), and CHES (&); TAPS buffer was used for the rest of the cultures. The data are derived from at least two independent experiments that gave similar results. When different pH_{max} values were obtained for a given strain, the lower value was adopted.

applied. The 1/2 MRS medium contained 10 g/liter glucose, 5 g/liter Bacto Proteose Peptone No. 3 (Becton Dickinson, Franklin Lakes, NJ), 5 g/liter Lab-Lemco powder (Oxoid, Basingstoke, Hampshire, UK), 2.5 g/liter Bacto yeast extract (Becton Dickinson), 0.5 g/liter Tween 80, 1 g/liter K₂HPO₄, 2.5 g/liter NaCH₃CO₂ · 3H₂O, 1 g/liter triammonium citrate, 0.1 g/liter MgSO₄ · 7H₂O, and 0.025 g/liter MnSO₄ · 4H₂O; the pH was adjusted to 6.5, using HCl. The seed culture was inoculated into 3 ml of buffered 1/2 MRS medium at an optical density at 660 nm (OD₆₆₀) of around 0.03. The buffered 1/2 MRS medium was prepared by supplementing 1/2 MRS medium with an appropriate buffer at 100 mM to prevent acidification of the culture medium, using the following buffers: MOPS (pH 6.5 to 7.9), HEPES (pH 7.1 to 8.1), TAPS (pH 8.1 to 9.1), and CHES (pH 8.7 to 9.7). The pH of these media was adjusted using NaOH or HCl. Growth experiments using the lactobacilli strains were conducted at a pH range of 6.5 to 9.1; experiments using *E. hirae* JCM 8729^T were conducted at a pH

range of 6.5 to 9.7. Incubation was conducted at the same temperatures and same anaerobic conditions as used for the seed cultures for up to 24 h, and growth was monitored by the OD₆₆₀, using a spectrophotometer (Spectronic 20D+; Milton Roy Company). Strains that reached an OD₆₆₀ of >0.15 within 24 h of being cultured were considered to exhibit positive growth. At this point, acidification of the media appeared to be negligible (around 0.05 pH unit). Additionally, specific growth rates (μ ; expressed in h⁻¹) were calculated during the exponential growth phase, using the following equation: $\mu = (\ln x_{t2} - \ln x_{t1}) / (t2 - t1)$, where x_{t2} and x_{t1} are the OD₆₆₀ at times $t2$ and $t1$, respectively. Normally, calculation was conducted at the early exponential phase (OD₆₆₀, ~0.3), at which point acidification of the medium (around 0.15 pH unit) did not seem to affect μ values significantly.

Intracellular pH measurement. The internal pH of five representative strains of lactobacilli (see Fig. 2) and *E. hirae* JCM 8729^T was measured at external-pH

FIG. 1. Specific growth rates of representative *Lactobacillus* (*Lb.*) strains and *Enterococcus* (*Ec.*) *hirae* JCM 8729^T. The strains were cultured in buffered 1/2 MRS media. Each point represents the mean $\pm$ the standard deviation of the results of two independent experiments.

values of 6.5 to 9.0, using the pH-sensitive fluorescence probe 5 (and 6)-carboxyfluorescein succinimidyl ester (cFSE), based on the method originally described by Breeuwer et al. (2), with some modification. The fluorescence intensity of this probe is increased at alkaline pH and reduced at acidic pH. Seed culture was inoculated into an appropriate volume of 1/2 MRS medium in a screw-cap bottle at an OD₆₆₀ of about 0.05. The culture was conducted to an OD₆₆₀ of around 0.6 (i.e., exponential growth phase) under the same conditions as described in the preceding section. The cultured cells were harvested by centrifugation at $8,000 \times g$ for 5 min at 4°C and washed twice with buffer A (150 mM potassium phosphate buffer with 1 mM MgSO₄ · 7H₂O [pH 6.5]). The cells were resuspended in buffer A containing 4 μ M cFDASE, a precursor molecule of cFSE, at an OD₆₆₀ of around 0.5 and incubated for 30 min at the temperatures described in the footnote to Table 1. During this incubation, the membrane-permeating cFDASE was cleaved by intracellular esterases, and the resultant cFSE molecules were conjugated to the aliphatic amines of intracellular proteins. After being centrifuged and washed with buffer A, the cells were resuspended in the same volume of buffer A. The nonconjugated probe was eliminated by the addition of glucose at a final concentration of 10 mM and subsequent incubation for 1 h at the temperatures described in the footnote to Table 1. After being centrifuged and washed with buffer A, the cells were resuspended in buffer A, and their energy was depleted through incubation for 1 h at the temperatures described in the footnote to Table 1. After being centrifuged and washed with buffer A, the cells were resuspended in buffer A at an OD₆₆₀ of around 6. A 100- μ l aliquot of this cell suspension was diluted in 1.9 ml of buffer B (buffer A supplemented with 100 mM MOPS or TAPS [pH 6.5]) in a cuvette, and the cuvette was placed in a stirred and heated (same culture temperature) cuvette holder in an LS50B fluorimeter (PerkinElmer Life and Analytical Sciences, Wellesley, MA). The internal pH was determined by measuring the fluorescence intensities of the cell suspension with excitation and emission wavelengths of 490 and 520 nm, respectively (slit widths of 2.5 nm). In the experiments, while the fluorescence intensity was being monitored, the cells were energized with 10 mM glucose at pH 6.5 and incubated until the intensity stabilized. After that, the pH of the mixture was increased to the desired values between 7.0 and 9.0, using NaOH or KOH (see Fig. 2), and the mixture was incubated for 30 min more. Subsequently, the cells were treated with nigericin, which dissipates transmembrane proton gradient (Δ pH), at a final concentration of 2 μ M, and the fluorescence intensity was measured for calibration at appropriate external-pH values adjusted with the corresponding alkali (NaOH or KOH) and HCl.

Growth in the presence of nigericin. To evaluate the effect of Δ pH on alkali tolerance, growth experiments using four strains of lactobacilli (see Fig. 3A) and *E. hirae* JCM 8729^T were conducted in the presence or absence of nigericin at pH values between 6.5 and 9.0. Both seed cultures and main cultures were grown as described under "Determination of the maximum alkaline pH allowing the growth of lactobacilli," with the following modifications. Potassium-enriched 1/2 MRS (K-1/2 MRS) medium was used for the seed cultures and as the basal medium for the main cultures, and the main cultures were grown in the presence or absence of nigericin. Potassium was added to avoid possible growth inhibition caused by nigericin, as was previously observed in *E. hirae* ATCC 9790^T (JCM

8729^T) because of the loss of K⁺ from the cells (10). K-1/2 MRS medium was prepared by modifying 1/2 MRS medium. The concentration of K₂HPO₄ was increased to 10 g/liter, and NaCH₃CO₂ · 3H₂O was replaced with 1.8 g/liter of potassium acetate. The following buffers at 100 mM were added to the K-1/2 MRS medium for the main cultures, and the pH was adjusted to the desired values using KOH (buffered K-1/2 MRS medium): MES (pH 6.5), MOPS (pH 7.0 to 7.5), and TAPS (pH 8.0 to 9.0). Nigericin in ethanol solution was added at a final concentration of 7 μ M. Control cultures received 0.1% (vol/vol) ethanol. Specific growth rates were calculated during the exponential growth phase as described above. Acidification of the culture medium during the calculation appeared to be at negligible levels owing to the added buffer system.

Measurement of glycolysis activity in the presence of nigericin. To evaluate the pH dependency of lactobacilli glycolysis activity, the activities of the four strains used in the preceding section together with that of *E. hirae* JCM 8729^T were measured in the presence of nigericin at various pHs between 4.0 and 9.5. The cells were cultured until the exponential phase (OD₆₆₀ = 0.6) in 1/2 MRS medium, harvested, and washed with buffer A, as described under "Intracellular pH measurement." The washed cell pellet was resuspended in the same buffer and incubated for 1 h at the same temperatures in which the cells were cultured to deplete their metabolic energy. The cells were washed once with buffer A and resuspended in the same buffer at an OD₆₆₀ of about 80. The basal composition of the reaction mixture for the glycolysis reaction contained 150 mM potassium phosphate buffer, 1 mM MgSO₄ · 7H₂O, and one of the following buffers at 100 mM with the pH adjusted to the desired values, using KOH: MES (pH 4.0 to 7.0), MOPS (pH 7.5), TAPS (pH 8.0 to 9.0), or TABS (pH 9.5). Subsequently, the energy-depleted cells were added at an OD₆₆₀ of about 4, and then nigericin in ethanol solution was added at a final concentration of 7 μ M. The control series received 0.1% (vol/vol) ethanol. The reaction was started by the addition of glucose at a final concentration of 10 mM. The mixtures were incubated at the temperature in which they were cultured, for appropriate time periods (see Fig. 3). After being incubated, the mixtures were centrifuged at $18,120 \times g$ for 5 min at 4°C, and the supernatant was filtered through a 0.2- μ m-pore-size membrane. The glucose concentrations of the filtered supernatants were measured, using Glucose C2 (Wako Pure Chemical Industries, Osaka, Japan).

RESULTS

Maximum pH allowing the growth of lactobacilli. We determined the maximum pH that allows growth (pH_{max}) for 34 strains of lactobacilli from various sources. We used *E. hirae* JCM 8729^T, a well-known alkali-tolerant strain (8), as a reference. As expected, *E. hirae* JCM 8729^T showed growth up to a pH of 9.7 (Table 1). Relatively high-alkali-tolerant strains in the lactobacilli included the plant isolates of *L. casei*, *L. paracasei* subsp. *tolerans*, *L. paracasei* subsp. *paracasei*, *L. curvatus*,

FIG. 2. Internal-pH measurements at various external pHs. MOPS buffer (○) or TAPS buffer (△) was used in buffer B (see Materials and Methods). The pH of buffer B was adjusted using either KOH (open symbols) or NaOH (filled symbols). The dashed lines represent the intracellular pH (pHin) which is equal to the extracellular pH (pHout). *Lb.*, *Lactobacillus*; *Ec.*, *Enterococcus*.

L. pentosus, and *L. plantarum*, with pHmax values between 8.5 and 8.9. Of these, *L. casei* NRIC 1917 and *L. paracasei* subsp. *tolerans* NRIC 1940 exhibited the highest alkali tolerance, with a pHmax of 8.9. Within the same species of lactobacilli, i.e., *L. casei*, *L. paracasei* subsp. *tolerans*, and *L. paracasei* subsp. *paracasei*, isolates from plant materials showed higher alkali tolerance than those from other sources. In contrast, strains of *L. brevis*, *L. delbrueckii* subsp. *delbrueckii*, *L. fermentum*, and *L. mali* that were isolated from plant materials showed lower pHmax values of between 7.5 and 7.7. Strains of *L. gasseri*, *L. johnsonii*, *L. reuteri*, *L. salivarius* subsp. *salicinius*, and *L. salivarius* subsp. *salivarius*, which were isolated from digestive tracts, exhibited moderate alkali tolerance, with pHmax values between 8.1 and 8.5. The dairy isolates of *L. delbrueckii* subsp. *bulgaricus*, *L. delbrueckii* subsp. *lactis*, and *L. helveticus* showed no alkali tolerance, with pHmax values between 6.7 and 7.1. The least-alkali-tolerant strains among those tested were *L. helveticus* JCM 1120^T and *L. acidophilus* JCM 1132^T, which had pHmax values as low as 6.7.

Among six representative strains of lactobacilli and *E. hirae* JCM 8729^T, *E. hirae* JCM 8729^T showed the highest μ as a function of the initial pH of the culture medium under the tested pH range of between 6.5 and 9.7 (Fig. 1). However, in all strains, μ decreased with the increasing initial pH of the medium.

Intracellular pH measurements. Next, we clarified the mechanism(s) determining the diversity of alkali tolerance. Previous studies suggested that antiporters acidify cytoplasmic pH under alkaline conditions in several alkali-tolerant species, including *E. hirae* ATCC 9790^T (15–17), and some alkaliphiles (20, 21). To verify the contribution of this mechanism in alkali tolerance, internal-pH measurements were performed, using five strains of lactobacilli with different pHmax values and *E. hirae* JCM 8729^T under a range of external-pH values from 6.5

to 9.0 (Fig. 2). An ATP-driven K⁺/H⁺ antiporter has been suggested for *E. hirae* ATCC 9790^T (JCM 8729^T) (15–17), whereas Na⁺/H⁺ antiporter activity has been implied for the alkaliphilic bacteria *Bacillus firmus* RAB (20) and *Exiguobacterium aurantiacum* NCIB 11798 (21). Thus, both NaOH and KOH were used to adjust the assay mixtures for pH measurements. At an external pH of 6.5, an Δ pH (i.e., an alkaline interior) was generated in all tested strains by energization with glucose. When the external pH was increased using KOH, the internal pH also rose, but at a slower rate, thereby decreasing the Δ pH. Above a specific external pH, Δ pH was reversed (i.e., the interior became acidic), and under highly alkaline external pH, the internal pH increased while maintaining the magnitude of the reversed Δ pH at about 0.5 in all tested strains, including *E. hirae* JCM 8729^T. We did not observe the remarkable cytoplasmic pH regulation previously reported at alkaline external pH in *E. hirae* ATCC 9790^T (14). The formation of the reversed Δ pH was observed not only in the highly alkali-tolerant strains *E. hirae* JCM 8729^T and *L. paracasei* subsp. *tolerans* NRIC 1940, but also in the least-alkali-tolerant strain, *L. acidophilus* JCM 1132^T (Fig. 2). Similar results were obtained when NaOH, rather than KOH, was used to adjust the pH, except that a significantly larger reversed Δ pH was observed in *L. acidophilus* JCM 1132^T than that observed when KOH was used for pH adjustment. Internal-pH regulation does not influence the diversity of lactobacilli alkali tolerance.

Growth in the presence of nigericin. The reversed Δ pH observed in the preceding experiments was expected to contribute to growth in alkaline conditions. To verify this, we examined growth in the presence or absence of nigericin, which dissipates Δ pH, using four strains of lactobacilli and *E. hirae* JCM 8729^T in buffered K-1/2 MRS. We observed dramatic decreases in the μ values of all the *Lactobacillus* strains and *E.*

FIG. 3. (A) Specific growth rates and (B) glycolysis activities of representative *Lactobacillus* (*Lb.*) strains and *Enterococcus* (*Ec.*) *hirae* JCM 8729^T. The experiments were carried out in the presence (filled symbols) or absence (open symbols) of nigericin. Glycolysis reactions were conducted for the following time periods: *L. acidophilus* JCM 1132^T and *L. paracasei* subsp. *tolerans* NRIC 1940, 2 h; *L. johnsonii*

hirae JCM 8729^T in the presence of nigericin under a range of acidic pH from 6.5 to 7.0 (Fig. 3A). This was expected from previous studies showing that maintenance of intracellular pH by H⁺-ATPase is important for growth and viability in acidic conditions (1, 18, 29). Under a range of alkaline pH, the μ values of *L. johnsonii* JCM 1022, *L. plantarum* NRIC 0380, and *L. paracasei* subsp. *tolerans* NRIC 1940 were decreased in the presence of nigericin and the pH_{max} of each strain was decreased by 0.5. Thus, the reversed Δ pH formed in alkaline conditions contributes to the alkali tolerance of the tested *Lactobacillus* strains. Furthermore, we observed that the pH_{max} of a given strain in the presence of nigericin was positively dependent on the pH_{max} in the absence of nigericin (Fig. 3A), suggesting that the degree of resistance of the cellular metabolism, including catabolism and anabolism to an alkaline environment, may determine the alkali tolerance of a given strain.

Glycolysis activity in the presence of nigericin. To evaluate the preceding hypothesis, we measured the glycolysis activity of the strains in the presence or absence of nigericin at various external pH values. As expected, we observed dramatic decreases in glycolysis activity at a range of acidic pH values for all strains tested (Fig. 3B). The glycolysis activities of the highly alkali-tolerant strains NRIC 1940 and *E. hirae* JCM 8729^T were also greatly decreased at a range of alkaline pH above 9.0 in the presence of nigericin (Fig. 3B). However, the optimum pH for glycolysis activity in the presence of nigericin of all strains tested was neutral, regardless of the strain's alkali tolerance. Interestingly, substantial glucose utilization in the highly alkali-tolerant strains NRIC 1940 and *E. hirae* JCM 8729^T was still detected up to a pH range of between 8.0 and 8.5 in the presence of nigericin, whereas the remaining strains lost most glycolysis activity. Therefore, the diversity of alkali tolerance is strongly dependent on the alkali tolerance of cellular metabolism, especially glycolysis reactions involved in the generation of energy.

DISCUSSION

We determined the pH_{max} of many lactobacilli for the first time. The pH_{max} ranged between 6.7 and 8.9, demonstrating the diversity of alkali tolerance in lactobacilli (Table 1). Strains of *L. casei*, *L. paracasei* subsp. *tolerans*, *L. paracasei* subsp. *paracasei*, *L. curvatus*, *L. pentosus*, and *L. plantarum* originating from plant material exhibited high alkali tolerance, with pH_{max} values of >8.5. It is not clear if there is a correlation between the plant environment and the strain's alkali tolerance. Most of these strains belong to two phylogenetic groups of lactobacilli: the *L. casei* group (strains tested, *L. casei*, *L. paracasei* subsp. *paracasei*, and *L. paracasei* subsp. *tolerans*) and the *L. plantarum* group (strains tested, *L. plantarum* and *L. pentosus*) (5). Therefore, we speculated that strains belonging to these groups may have a tendency for alkali tolerance from an evolutionary viewpoint. The next-highest alkali-tolerant

JCM 1022, 1.5 h; *L. plantarum* NRIC 0380 and *E. hirae* JCM 8729^T, 1 h. Each point represents the mean $\pm$ the standard deviation of the results of two independent experiments.

group contained strains of *L. gasseri*, *L. johnsonii*, *L. reuteri*, *L. salivarius* subsp. *salicinius*, and *L. salivarius* subsp. *salivarius* isolated from digestive tracts; these ranged in pH_{max} from 8.1 to 8.5. The small intestine, which is colonized by various intestinal bacteria, including lactobacilli and *E. hirae*, has a weakly alkaline environment of around pH 7.5 (7). Thus, we assumed that *Lactobacillus* strains originating in human intestine would exhibit relatively high pH_{max} as a result of adaptation to the alkaline environment of the small intestine. *L. acidophilus* JCM 1132^T seems an exceptional strain, with the pH_{max} as low as 6.7. However, as discussed later in the Discussion section, this strain possibly has certain system(s) to create significantly larger reversed ΔpH under alkaline conditions than any other strain tested (Fig. 2), suggesting survival under an alkaline environment (without growth). We found extremely low alkali tolerance in the dairy strains of *L. delbrueckii* subsp. *bulgaricus*, *L. delbrueckii* subsp. *lactis*, and *L. helveticus*, with pH_{max} values between 6.7 and 7.1. Dairy starter fermentations generally proceed under severe acidic conditions of around pH 4. Thus, dairy strains may have little chance of encountering an alkaline environment and, therefore, have not evolved high tolerance.

We found that the diversity of alkali tolerance depended strongly on the alkali tolerance of the cellular metabolism, including catabolism and anabolism, as observed in terms of growth (Fig. 3A), especially on the glycolysis reactions of a given strain (Fig. 3B). In fact, the alkali-tolerant reference strain, *E. hirae* JCM 8729^T, exhibited the optimum pH for growth of 8.5 in the presence of nigericin, which is significantly higher than those of the other tested strains, which showed the pH optima to be around 7.5 (Fig. 3A). The pH/glycolysis activity profile of *E. hirae* JCM 8729^T also shifted to the alkaline side compared to those of the other tested strains in the presence of nigericin (Fig. 3B). A dramatic decrease in glycolysis activity, which was not observed in the other strains, occurred at pH 6.0 (Fig. 3B). This shift in the pH/glycolysis activity profile was also observed to a lesser extent in *L. paracasei* subsp. *tolerans* NRIC 1940, which showed the highest alkali tolerance of the tested lactobacilli (Fig. 3B). Nevertheless, the optimal pH for glycolysis activity was neutral in all strains, regardless of their alkali tolerance (Fig. 3B), which is consistent with a previous finding of neutral optimal pH in many enzymes of LAB involved in carbohydrate and amino acid metabolism (11). It is interesting that in *L. acidophilus* JCM 1132^T, substantial glycolysis activity in the absence of nigericin was still observed at pH 7.5 or even at pH 8.5 (Fig. 3B), while the pH_{max} was 6.7 (Table 1). These results suggest that in this strain anabolic reactions may be more sensitive to an alkaline environment than catabolic reactions (glycolysis reactions).

Reversed ΔpH formation was observed at alkaline external pH in all of the tested lactobacilli, as well as in *E. hirae* JCM 8729^T, with a magnitude of around 0.5 (Fig. 2). The dissipation of this ΔpH by the addition of nigericin lowered the pH_{max} of each tested strain by 0.5 (Fig. 3A). These consistent results clearly indicate the contribution of the reversed ΔpH to alkali tolerance. The formation of the reversed ΔpH under alkaline external pH has already been reported for *E. hirae* ATCC 9790^T (14, 18, 19), and such ATP-dependent K⁺/H⁺ antiporters (15–17) and/or physicochemical forces as the Donnan potential (19) have been suggested to contribute to these phenomena. In the case of *L. acidophilus* JCM 1132^T, *L. plantarum*

NRIC 0380, and *L. paracasei* subsp. *tolerans* NRIC 1940, glucose utilization was still observed under the alkaline pH range in which the reversed ΔpH was observed (Fig. 2 and 3B), thereby ensuring an energy supply under these conditions. Therefore, the observed reversed ΔpH formation in these *Lactobacillus* strains (Fig. 2) may possibly be mediated not only by some physicochemical force but also by certain energy-dependent Na⁺(K⁺)/H⁺ antiporter(s). The larger reversed ΔpH observed in *L. acidophilus* JCM 1132^T when NaOH was used for pH adjustment than that observed when KOH was used (Fig. 2) suggested the possible operation of an Na⁺/H⁺ antiporter. In this bacterium, since the pH_{max} is 6.7 (Table 1), the observed reversed ΔpH (Fig. 2) may not contribute to the growth but may contribute to the survival of this bacterium under alkaline stress conditions.

We did not detect the remarkable regulation of cytoplasmic pH in *E. hirae* JCM 8729^T (ATCC 9790^T) in external alkaline pH (Fig. 2) by such transporters as the ATP-driven K⁺/H⁺ antiporter reported previously for this bacterium (14–17). Similar results have been reported by Mugikura et al. (22). These discrepancies may have been caused by differences in the methods used for intracellular-pH measurements. It has been argued that the amines used for the measurement of internal pH by Kakinuma (14) tend to overestimate ΔpH because of the binding of amines to intracellular materials such as RNA as well as to the cell surface (22). In contrast, Mugikura et al. (22) used weak acid for internal pH measurements, and its binding to intracellular materials is negligible. We used a pH-sensitive fluorescent probe, cFSE, which allows the direct monitoring of intracellular pH.

We investigated the physiological response of many *Lactobacillus* strains to an alkaline environment. The most important factor determining the diversity of alkali tolerance in lactobacilli was the alkali tolerance of the strains' glycolysis reactions. In addition, the reversed ΔpH formed in alkaline environments, probably because of the Donnan potential and/or certain energy-dependent Na⁺(K⁺)/H⁺ antiporter(s), contributed to the alkali tolerance. Although a variety of mechanisms aid LAB survival in acidic environments, lactobacilli seem to have poor evolutionary preparation for growth in alkaline environments. It has been reported that prior exposure to weak environmental stresses dramatically increases the tolerance to the corresponding stress in various LAB strains (3, 4, 28). Potential alkaline stress adaptation, as has been reported for *E. faecalis* ATCC 19433^T (9), needs to be checked in lactobacilli for a more-comprehensive understanding of the mechanism of alkali tolerance. We have illustrated previously unexplored aspects of lactobacilli in alkaline conditions. We believe that these findings will help to understand the basic biology of lactobacilli and expand the application range of this group of bacteria.

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