



D-Methionine and D-Phenylalanine Improve Lactococcus lactis F44 Acid Resistance and Nisin Yield by Governing Cell Wall Remodeling

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ABSTRACT Lactococcus lactis encounters various environmental challenges, especially acid stress, during its growth. The cell wall can maintain the integrity and shape of the cell under environmental stress, and D-amino acids play an important role in cell wall synthesis. Here, by analyzing the effects of 19 different D-amino acids on the physiology of L. lactis F44, we found that exogenously supplied D-methionine and D-phenylalanine increased the nisin yield by 93.22% and 101.29%, respectively, as well as significantly increasing the acid resistance of L. lactis F44. The composition of the cell wall in L. lactis F44 with exogenously supplied D-Met or D-Phe was further investigated via a vancomycin fluorescence experiment and a liquid chromatography-mass spectrometry assay, which demonstrated that D-Met could be incorporated into the fifth position of peptidoglycan (PG) muropeptides and D-Phe could be added to the fourth and fifth positions. Moreover, overexpression of the PG synthesis gene murF further enhanced the levels of D-Met and D-Phe involved in PG and increased the survival rate under acid stress and the nisin yield of the strain. This study reveals that the exogenous supply of D-Met or D-Phe can change the composition of the cell wall and influence acid tolerance as well as nisin yield in L. lactis.

IMPORTANCE As D-amino acids play an important role in cell wall synthesis, we analyzed the effects of 19 different D-amino acids on *L. lactis* F44, demonstrating that D-Met and D-Phe can participate in peptidoglycan (PG) synthesis and improve the acid resistance and nisin yield of this strain. *murF* overexpression further increased the levels of D-Met and D-Phe incorporated into PG and contributed to the acid resistance of the strain. These findings suggest that D-Met and D-Phe can be incorporated into PG to improve the acid resistance and nisin yield of the strain study provides new ideas for the enhancement of nisin production.

KEYWORDS D-amino acid, Lactococcus lactis, acid resistance, cell wall, nisin

L actococcus lactis, widely used in the food industry, is a Gram-positive bacterium. Nisin, an antimicrobial peptide produced by *L. lactis*, is used as a safe food preservative and antimicrobial (1, 2). Many studies have focused on methods to increase nisin yield, such as metabolic regulation, fermentation optimization, and genetic modifications (2–8). It is widely known that *L. lactis* encounters several environmental stresses, especially lactate, during fermentation, and the acidic medium inhibits its growth and nisin production (9), so we previously utilized a range of strategies to improve the acid resistance of *L. lactis*, which contributes to increases in nisin yield (10, 11). Citation Wu H, Xue E, Zhi N, Song Q, Tian K, Caiyin Q, Yuan L, Qiao J. 2020. p-Methionine and p-phenylalanine improve *Lactococcus lactis* F44 acid resistance and nisin yield by governing cell wall remodeling. Appl Environ Microbiol 86:e02981-19. https://doi.org/10 1128/AFM 02981-19.

Editor Maia Kivisaar, University of Tartu Copyright © 2020 American Society for Microbiology. All Rights Reserved.

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Received 22 December 2019 Accepted 22 February 2020

Accepted manuscript posted online 28 February 2020 Published 17 April 2020

The cell wall, a reticular polymer located outside the cell membrane, can maintain the integrity and shape of the cell. In Gram-positive bacteria, the peptide polysaccharide layer is thick and provides anchor points for other components of the cell wall, such as wall phosphoric acid and capsule polysaccharide (12, 13). Peptidoglycan (PG), an essential component of the cell wall, is a polymer of N-acetylmuramic acid (MurNAc) and N-acetylglucosamine (GlcNAc) cross-linked by peptide bridges. The D-Ala-D-Ala dipeptide (catalyzed by the D-alanine ligase Ddl) is incorporated into the monosaccharide tripeptide (generated by MurABCDE) through MurF (UDP-N-acetylmuramoyltripeptide-D-alanyl-D-alanine ligase). After the formation of monosaccharide pentapeptide, MraY catalyzes the transfer of phospho-MurNAc-pentapeptide to the lipid carrier undecaprenyl phosphate to form a lipid-linked N-acetylmuramic acid derivative (lipid I), and MurG couples GlcNAc to the C-4 hydroxyl of lipid I to generate the beta-linked disaccharide (lipid II). D-Asp is then attached to the third L-Lys to form a peptide bridge under the action of YxbA ligase (14). The membrane protein MurJ, Amj, or FtsW serves as the lipid II flippase that can transport lipid II across the membrane (15-17). The polymerization of lipid II to generate individual PG strands that are subsequently cross-linked to form PG layers is catalyzed by the penicillin-binding proteins and glycosyltransferases (18). The newly synthesized PG matures through the modification of synthetases and hydrolases (such as L,D-transpeptidase and carboxypeptidase) and is eventually hydrolyzed by PG hydrolases during cell growth and division. Partially hydrolyzed PG fragments are transported to the cytoplasm and reused in lipid II biosynthesis during PG recycling (19, 20).

Our previous studies showed that the cell wall modification could improve the acid resistance of *L. lactis*. The overexpression of *asnH* increased the acylation level of the cell wall peptide bridge D-Asp, helped to maintain the rigidity of PG, and promoted the acid tolerance and nisin yield of *L. lactis* (10). Moreover, the increase in the O-acetylation of MurNAc and the N-acetylation of GlcNAc in the cell wall contributed to the integrity of the cell wall in *L. lactis* F44 and increased its acid resistance (11).

The D-amino acids involved in the PG synthesis are normally D-alanine and D-glutamate in bacteria. However, Lam and coworkers and Cava and coworkers found that Vibrio cholerae could produce D-methionine and D-leucine, called noncanonical D-amino acids (NCDAAs), which influenced the composition, amount, and flexibility of PG (21, 22). The presence of NCDAAs in PG was confirmed by adding NCDAAs to the medium (23-25). Other studies found that Gly was exchanged with D-Ala at the first, fourth, or fifth position of the peptide chain in bacterial PG (26) and that D-Met, D-Trp, or D-Phe replaced D-Ala at the fourth position of the peptide chain in the PG of Escherichia coli. L-Isomer addition to the external source affected neither the structure of the PG nor the synthesis (27). So far, NCDAAs have been found to participate in PG synthesis in four ways: (i) L,D-carboxypeptidase catalyzes exogenous NCDAAs to exchange with the D-Ala residue of a disaccharide tetrapeptide (27, 28); (ii) L,Dtranspeptidase, like LdtA and LdtB, directly replaces D-Ala with NCDAAs at the fourth or fifth position of the cross-linked peptide chain (22); (iii) D,D-transpeptidase has the potential to mediate NCDAAs to replace D-Ala at the fifth position of the stem peptide (24); and (iv) NCDAAs participate in PG synthesis through two steps catalyzed by the enzymes Ddl and MurF (22).

Although NCDAAs which can be incorporated into the PG have been extensively studied in several Gram-negative strains, little information is available about the effects of NCDAAs on the Gram-positive species *L. lactis*. This study gives insights into the pivotal role of *D*-amino acids in acid resistance and nisin yield and provides biological evidence that *D*-Met and *D*-Phe can participate in cell wall synthesis in *L. lactis* F44.

RESULTS

Effects of different D-amino acids on acid resistance of *L. lactis* **F44.** To investigate the effects of various D-amino acids on *L. lactis* **F44**, we added the 19 different D-amino acids to the medium separately and detected the acid resistance of F44 in each sample. The experimental groups were *L. lactis* **F44** with the separate addition of



FIG 1 (A) Survival rate of *L. lactis* F44 with the addition of various D-amino acids after 1.5 h acid shock. (B) Survival rate of *L. lactis* F44 with the addition of various D-amino acids after 2.0 h acid shock. Error bars indicate standard deviations (SD) for three independent experiments. *, P < 0.01; **, P < 0.001 (*t* test).

various D-amino acids (50 mM) in the medium, and the control group was F44 without any D-amino acid addition. In the mid-log phase, the cells were exposed to pH 3.0 for 1.5 or 2 h. As shown in Fig. 1, the survival rates of F44 added with D-Met, D-Phe, D-Asn, or D-Glu/Gln were significantly higher than that of the control (P < 0.01). Among them, the survival rate of the strain with added D-Phe was the highest—70.82% ± 2.26% and 41.33% ± 1.25% at 1.5 h and 2 h, respectively, which were 1.84 and 1.92 times higher than that of the control (38.47% ± 2.37% at 1.5 h; 21.52% ± 1.00% at 2.0 h). D-Met addition increased the survival rate by 58.33% and 79.53% after 1.5 h and 2 h of acid stress, respectively. Therefore, the separate addition of D-Met, D-Phe, D-Asn, and D-Glu in the medium could improve the survival of F44 under acid stress.

The influence of p-amino acids on nisin yield is different from that of L-amino acids. The growth properties, optical density at 600 nm (OD₆₀₀), pH, and nisin production per cell were detected during fermentation. With the separate addition of 19 different p-amino acids, the final OD₆₀₀ value of *L. lactis* F44 was lower, and correspondingly the pH value was higher, than that of the control (see Fig. S1 and S2 in the supplemental material). Notably, p-Cys significantly inhibited strain growth. Figure 2 and Fig. S3 show that the addition of p-Met, p-Phe, p-Asn, p-Trp, or p-Leu to the medium significantly increased nisin production. Among them, p-Met addition enhanced the nisin yield per cell to $5.99 \times 10^{-7} \pm 1.08 \times 10^{-8}$ IU/CFU, which was 93.22% higher than that of the control ($6.24 \times 10^{-7} \pm 2.65 \times 10^{-8}$ IU/CFU) was 101.29% higher. Therefore, the addition of p-Met or p-Phe could improve the nisin yield of *L. lactis* F44.

L-Amino acids were added to the fermentation medium to investigate the effects of L-Met and L-Phe on nisin yield, which helped to illuminate the question of whether D-Met and D-Phe could be converted to L-Met and L-Phe to influence F44 nisin yield. As shown in Fig. 2, the nisin yield of *L. lactis* F44 with the 50 mM L-Met or L-Phe addition did not show an obvious change, which was different from the result with the D-amino acid addition.



FIG 2 Effects of D-amino acids and L-amino acids on the nisin titer per cell of *L. lactis* F44. Samples were taken every 2 h from 6 to 10 h. Error bars indicate the SD for three independent experiments. **, P < 0.01 (*t* test).

Preliminary verification of D-Met and D-Phe involved in PG synthesis by fluorescence detection. NCDAAs were found to be involved in the formation of PG in the fourth or fifth position of the peptide chain in *V. cholerae* (22). Here, to investigate whether D-Met and D-Phe could be incorporated into the PG muropeptides to influence the acid resistance and nisin yield of *L. lactis*, a fluorescent derivative of vancomycin (Van-FL) was used to detect the insertion of NCDAAs (29, 30). According to the previous study, Van-FL could interact with D-Ala–D-Ala at the fourth and fifth positions of the peptide chain through hydrogen bonding to form stable complexes (31), and its binding could be initially observed by the fluorescence of cell walls (21). Given that Van-FL could inhibit the growth of this strain, the tolerance of *L. lactis* F44 to Van-FL was determined, and 300 ng/ml was found to be a relatively high tolerance concentration (at which no visible inhibition of growth occurred) (see Fig. S4 in the supplemental material).

A fluorescence detection assay was performed with 300 ng/ml Van-FL. As shown in Fig. 3, the fluorescence intensities of the strains with either D-Met or D-Phe in the medium were weaker than that of the control, and the intensity of the sample with the D-Phe addition was the weakest. Therefore, we hypothesized that more D-Phe than D-Met was incorporated into the cell wall muropeptides, thus possibly leading to a more significant increase in acid resistance and nisin yield.

D-Met and D-Phe can be incorporated into the cell wall muropeptides, as verified by LC-MS. To test the above hypothesis, we extracted and detected the PG after *L. lactis* F44 had been grown in the fermentation medium containing D-Met or D-Phe by liquid chromatography-mass spectrometry (LC-MS). As shown in Fig. 4 and Table 1, the monomer Penta^{Met} (*N*-acetylglucosamine–*N*-acetylmuramyl pentapeptide with the asparagine peptide bridge) was found in the D-Met sample, which indicated that D-Met was incorporated at the fifth position of muropeptides in PG. According to the previous study, D-Met could be incorporated at two locations within PG, namely, the fourth and fifth positions of muropeptides (22). However, in the present study, D-Met was found only at the fifth position of muropeptides, and it accounted for 0.42% of the total PG in *L. lactis* F44 (Fig. 4).

The monomers Tetra^{Phe} (*N*-acetylglucosamine–*N*-acetylmuramyl tetrapeptide with the aspartate peptide bridge) and Penta^{Phe} (*N*-acetylglucosamine–*N*-acetylmuramyl pentapeptide with the asparagine peptide bridge) were found in the D-Phe sample, indicating that D-Phe replaced the D-Ala in the fourth and fifth positions of muropeptides, respectively. In our analysis, Tetra^{Phe} and Penta^{Phe} constituted 0.85% and 6.75% of total PG, respectively, demonstrating that the D-Phe sample showed a higher level of incorporation than the D-Met sample, which was consistent with the results of vancomycin fluorescence detection.





FIG 3 Verification of NCDAAs involved in PG synthesis through fluorescence detection. Fluorescence intensity of *L. lactis* F44 was measured after growth in the medium containing D-Met and D-Phe for 6 h. The control group did not have any added D-amino acid. Van-FL (300 ng/ml) was used to detect the insertion of NCDAAs. The fluorescence value is the fluorescence intensity of a single cell, and the fluorescence values of 50 cells in the field were taken to calculate the mean value.

Fluorescence assay of recombinant strains *L. lactis* **F44-***ddl* **and F44-***murF* **with the addition of D**-**amino acids.** According to a previous study, two cell wall synthesis enzymes, Ddl and MurF, are related to the addition of NCDAAs into the fifth position of muropeptides (22). In the present work, the effects of the two enzymes on the incorporation of D-Phe and D-Met in PG synthesis were further explored. The recombinants *L. lactis* F44-*ddl* and F44-*murF*, which overexpress the *ddl* and *murF* genes, respectively, were constructed, and then a Van-FL fluorescence assay was performed.

As shown in Fig. 5, *ddl* or *murF* overexpression enhanced the binding of Van-FL to the tripeptides, leading to an increase in fluorescence intensity. Compared with F44, the increase in F44-*murF* was more significant than that in F44-*ddl*. Furthermore, the fluorescence intensity decreased when D-Met or D-Phe was added to the medium of F44-*murF*, and the influence of D-Phe was greater than that of D-Met. Therefore, it was suggested that *murF* or *ddl* overexpression could promote the conjugation of D-Ala–D-Ala with monosaccharide tripeptides to form monosaccharide pentapeptides. Notably, *murF* influenced the PG synthesis more significantly than *ddl*, and we further explored the changes in the composition of the PG of F44-*murF* by LC-MS.

MurF catalyzes the formation of the precursors replaced by D-Met or D-Phe, as verified by LC-MS. By using LC-MS analysis, the monomer Penta^{Met} was identified (Fig. 6), and it accounted for 0.72% of its total PG extracted by F44-*murF* with the addition of D-Met. As shown in Table 2, with the D-Met addition, F44-*murF* had 71.42% more PG than F44. Meanwhile, the PG of F44-*murF* with the D-Phe addition was extracted, and the monomers Penta^{Phe}, Penta^{Phe(X)} (*N*-acetylglucosamine–*N*-acetylmuramyl pentapeptide with the aspartate peptide bridge and a molecule of water lost), and Tetra^{Phe} accounted for 10.48%, 1.18%, and 9.35% of their total PGs, respectively. Compared with *L. lactis* F44, Penta^{Phe} and Tetra^{Phe} were increased by 55.25% and approximately 10-fold, respectively, revealing that the D-Phe sample had a higher level of binding than the D-Met sample, and the amount of D-Ala–D-Phe participating in the syn-



FIG 4 Analysis of *L. lactis* F44 PG by LC-MS. (A) LC-MS analysis of *L. lactis* F44 without the addition of any D-amino acid. (B) LC-MS analysis of *L. lactis* F44 PG after growth in medium containing D-Met for 6 h. Peak 1 represents D-Met binding to a disaccharide pentapeptide with the asparagine-peptide bridge, namely, Penta^{Met} (*N*-acetylglucosamine-*N*-acetylmuramyl pentapeptide with the asparagine peptide bridge). The structures and parameters are shown in Table 1. (C) LC-MS analysis of *L. lactis* F44 PG after growth in medium containing D-Phe for 6 h. Peak 2 represents D-Phe binding to Penta^{Phe}, a disaccharide (*N*-acetylglucosamine-*N*-acetylmuramyl) pentapeptide with the asparagine peptide bridge. Peak 3 represents D-Phe binding to a Tetra^{Phe}, a disaccharide (*N*-acetylglucosamine-*N*-acetylmuramyl) tetrapeptide with the asparagine peptide bridge. The structures and parameters are shown in Table 1.

thesis of pentapeptides was higher than that of D-Ala–D-Met. From these results, we came to the conclusion that *murF* plays an important role in the incorporation of D-Met or D-Phe into PG.

Overexpressing *murF* enhances the effects of p-Met or p-Phe on acid resistance and nisin yield. The effect of *murF* on the incorporation of p-Met or p-Phe to PG was further determined by the acid resistance and nisin yield assays, through which we found that, without any p-amino acid addition, the survival rate under acid stress of F44-*murF* was not obviously different from that of F44 (Fig. 7A). Compared with that of F44 without any p-amino acid addition, the survival rate of F44-*murF* with the p-Met addition was enhanced 2.10-fold or 2.91-fold after 1.5 h or 2.0 h acid shock, respectively, and that of F44 with the p-Phe addition increased 1.92-fold (1.5 h) and 2.44-fold (2.0 h).

TABLE 1 Analysis of L. lactis F44 PG by LC-MS

Additive	Structure ^a	Mol wt	Retention time (min)	Area (%)
D-Met	Penta ^{Met}	1,140.6	2.5	0.42
D-Phe	PentaPhe	1,156.6	8.7	6.75
D-Phe	Tetra ^{Phe}	1,074.5	28.3	0.85

^aPenta^{Met}, G–M–L-Ala–D-Gln–L-Lys (D-Asn)–D-Ala–D-Met; Penta^{Phe}, G–M–L-Ala–D-Gln–L-Lys (D-Asn)–D-Ala–D-Phe; Tetra^{Phe}, G–M–L-Ala–D-Gln–L-Lys (D-Asp)–D-Phe (D-Asn or D-Asp is the peptide bridge bound to L-Lys, G is *N*-acetylglucosamine [GlcNAc], and M is *N*-acetylmuramic acid [MurNAc]).



FIG 5 Fluorescence detection of *ddl*- and *murF*-overexpressing strain. The fluorescence intensities of F44-*ddl* and F44-*murF* after growth in medium containing D-Met and D-Phe for 6 h are shown. The control group was *L. lactis* F44. Van-FL (300 ng/ml) was used to detect the insertion of NCDAAs. The fluorescence value was the fluorescence intensity of a single cell, and the fluorescence values of 50 cells in the field were taken to calculate the mean value.

As shown in Fig. 7B, the addition of D-Met and D-Phe in the medium further increased the nisin yield of F44-*murF* 2.02-fold and 2.34-fold, respectively, compared with that of F44 without any D-amino acid addition. These results suggested that overexpressing *murF* could further enhance the effects of D-Met or D-Phe on the acid resistance and nisin yield.

DISCUSSION

Microorganisms communicate with the environment by producing diverse metabolites (32). For instance, many bacteria have been found to produce NCDAAs during growth: *V. cholerae* produces D-Met, D-Leu, and D-Arg; *Bacillus subtilis* generates D-Tyr and D-Phe (21, 29, 33); and a variety of lactic acid bacteria species produce D-branchedchain amino acids (34). Bacteria utilize NCDAAs to support their growth, regulate spore germination, and reshape their cell walls to be resistant to several environmental stresses (21, 25, 35, 36).

This study demonstrated that two types of NCDAAs, p-Met and p-Phe, significantly increased the survival rate and nisin yield of *L. lactis* F44 (Fig. 1 and 2). NCDAAs were identified as being involved in PG composition and made the strain highly resistant to environmental stresses in *V. cholerae* (21, 22). Our previous studies showed that the changes in cell wall modification could increase acid resistance and indirectly enhance nisin yield (10, 11).

In *V. cholerae* and *B. subtilis*, NCDAAs can bind to the fourth or fifth position of the cell wall peptide chain, which affects the function of the cell wall (22). Vancomycin is a clinically important antibiotic that can specifically bind to D-Ala–D-Ala at the fourth and fifth positions of the newly synthesized PG (31). The fluorescence derivatives of vancomycin can provide technical support for the exploration of the topological structure of PG synthesis (30). In this study, the fluorescence derivative Van-FL was used to investigate the incorporation of D-amino acids, and the fluorescence intensity of strains with the D-Met or D-Phe addition was weaker than that of the control (Fig. 3). We



FIG 6 Analysis of *L. lactis* F44 PG by LC-MS. (A) Result for *L. lactis* F44-*murF* without addition of any D-amino acids. (B) LC-MS analysis of *L. lactis* F44-*murF* PG after growth in medium containing D-Met for 6 h. Peak 1 represents D-Met binding to Penta^{Met}, a disaccharide pentapeptide with the asparagine-peptide bridge. The structures and parameters are shown in Table 2. (C) LC-MS analysis of *L. lactis* F44-*murF* PG after growth in medium containing D-Phe for 6 h. Peaks 2 and 3 represent D-Phe binding to Penta^{Phe}, a disaccharide pentapeptide with the asparagine peptide bridge. Peak 4 represents D-Phe binding to Penta^{Phe}(X), a disaccharide (*N*-acetylglucosamine-*N*-acetylmuramyl) tetrapeptide with the asparate peptide bridge and a molecule of water loss. Peaks 5, 6, and 7 represent D-Phe binding to Tetra^{Phe}, a disaccharide tetrapeptide with an aspartate peptide bridge. The structures and parameters are shown in Table 2.

speculated that this was because D-Met or D-Phe was incorporated into the fourth or fifth position of the peptide chain, and this consequently hindered the binding of the Van-FL to the peptide chain (31). Moreover, an LC-MS assay showed that D-Met was incorporated at the fifth position of the PG peptide chain and D-Phe at the fourth and fifth positions of (Fig. 4 and Table 1). Our results are consistent with previous studies in *V. cholerae* and *B. subtilis* (21, 22), which indicates that the incorporation of NCDAAs in PG is a common phenomenon in bacteria.

In *V. cholerae*, the incorporation of D-Met into the fourth position in muropeptides is generated by two enzymes involved in the synthesis of PG precursors, Ddl and MurF (22). Ddl connects two D-Ala to form D-Ala–D-Ala, and MurF connects the monosaccharide tripeptide with the dipeptide D-Ala–D-Ala (or D-Ala–D-Phe or D-Ala–D-Met) to form

TABLE 2 Analysis of L. lactis F44-murF PG by LC-MS

Additive	Structure ^a	Mol wt	Retention time (min)	Area (%)
D-Met	Penta ^{Met}	1,140.6	2.03	0.72
D-Phe	Penta ^{Phe}	1,156.6	5.6/10.4	10.48
D-Phe	Tetra ^{Phe}	1,074.5	24.5/24.7/29.4	9.35
D-Phe	Penta ^{Phe(X)}	1,139.5	21.7	1.18

^aPenta^{Met}, G–M–L-Ala–D-Gln–L-Lys (D-Asn)–D-Ala–D-Met; Penta^{Phe}, G–M–L-Ala–D-Gln–L-Lys (D-Asn)–D-Ala–D-Phe; Tetra^{Phe}, G–M–L-Ala–D-Gln–L-Lys (D-Asp)–D-Phe; Penta^{Phe(X)}, G–M–L-Ala–D-Gln–L-Lys (D-Asp)–D-Ala–D-Phe (D-Asn) or D-Asp is the peptide bridge bound to L-Lys, G is *N*-acetylglucosamine [GlcNAc], M is *N*acetylmuramic acid [MurNAc], and X is the loss of an H₂O molecule (18 Da).



FIG 7 (A) Survival rate of *L. lactis* F44 and F44-*murF* with added D-Met or D-Phe after 1.5 h and 2.0 h acid shock. **, P < 0.001 (*t* test). (B) Nisin titer of *L. lactis* F44 and F44-*murF* with added D-Met or D-Phe. Samples were taken every 2 h from 6 to 10 h. Error bars indicate the SD for three independent experiments.

a conventional (or unconventional) monosaccharide pentapeptide (12, 13). In this study, the overexpression of *ddl* and *murF* increased the fluorescence intensity (Fig. 5), because the enhanced synthesis of D-Ala–D-Ala provided more substrates binding to the monosaccharide tripeptides. However, the fluorescence intensities of the recombinant strains were weakened when D-Met or D-Phe was added, and that seen with D-Phe was lower, indicating a higher binding level of D-Ala–D-Phe. This result was further confirmed by the LC-MS assay for the PG of F44-*murF* (Fig. 6 and Table 2). Moreover, the addition of D-Met or D-Phe further increased the acid resistance of F44-*murF* (Fig. 7). A previous study found that strains producing NCDAAs were more resistant to osmotic stress, because NCDAAs could cause alterations in cell wall structure, such as changing the length of glycan chains and the degree of cross-linking, leading to a stronger PG in the strain (21). Hence, we suggest that overexpressing *murF* in *L. lactis* F44 could contribute to the incorporation of D-Met and D-Phe into the PG and, therefore, increase the acid resistance and indirectly enhance the nisin titer of *L. lactis* F44.

In addition, two Ldts (LdtA and LdtB), inner membrane-anchored periplasmic proteins, contributed to the incorporation of D-Met in the fourth position of the PG peptide chain (22). In *L. lactis* F44, only an Ldt (ATY88872.1) containing an ErfK (*E. coli* LdtA) conserved domain was annotated, but it showed a low similarity (25.81% identity), and the function was unknown. *E. coli* is a non-NCDAA producer, but it can incorporate D-Met into the PG muropeptides by exogenous addition, and similar results were also obtained with some strains of the Gram-positive organisms *B. subtilis, Enterococcus faecalis*, and *S. aureus* (22). Whether the Gram-positive strain F44 can produce NCDAAs is still unclear and remains to be further verified. However, in this study, we determined that F44 could also incorporate exogenously supplied D-Met and D-Phe into muropeptides. These results suggest that NCDAAs may serve as signals for microbial communication in microbial communities under the changing environmental threats (22).

In conclusion, this study found that the NCDAAs D-Met and D-Phe participate in the

Strain or plasmid	Description ^a	Reference or source
Strains		
L. lactis F44	Evolved from L. lactis YF11 (accession no. PRJNA419050)	6
F44-ddl	Overexpression of <i>ddl</i> in WT	This study
F44-murF	Overexpression of <i>murF</i> in WT	This study
F44-pLEB124	Introduction of expression vectors in WT	Laboratory stock
M. flavus	Indicator strains for nisin yield diffusion	Laboratory stock
Plasmids		
pLEB124·P _{P45}	E. coli-L. lactis shuttle and expression vector, Em ^r ; P45 promoter, low copy no.	Laboratory stock
pLEB124·P _{P45} ddl	ddl cloned in pLEB124 P_{P45}	This study
pLEB124·P _{P45} murF	murF cloned in pLEB124 P _{P45}	This study

TABLE	3	Strains	and	pl	asmid	s	used	in	this	stud	y
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^aEm^r, erythromycin resistant.

synthesis of PG and affected the survival rate and nisin yield of *L. lactis* F44. Moreover, overexpressing *ddl* or *murF* enhanced the level of p-Met and p-Phe participating in cell wall synthesis, and *murF* played an important role in the incorporation process. This study gives insight into the pivotal role of NCDAAs in the acid resistance and provides biological evidence that p-Met and p-Phe can participate in *L. lactis* F44 cell wall synthesis.

MATERIALS AND METHODS

Bacterial strains and growth conditions. Bacterial strains are listed in Table 3. *L. lactis* F44 strain (accession number PRJNA419050) was cultured in seed medium or fermentation medium separately supplemented with different D-amino acids when needed. The seed medium was prepared as follows (wt/vol): yeast extract (1.5%), peptone (1.5%), KH₂PO₄ (2.0%), sucrose (2.0%), NaCl (0.15%), and MgSO₄·7H₂O (0.015%), pH 7.2. The fermentation medium was similar to the seed medium except for the addition of corn steep liquor (0.3%) and cysteine (0.26%). The constructed vectors were transformed into *E. coli* for enrichment and then electrotransformed into *L. lactis* F44 (9). All the *L. lactis* strains were incubated at 30°C, and *E. coli* strains were grown in liquid Luria-Bertani (LB) medium at 37°C, with shaking at 180 rpm. *Micrococcus flavus*, an indicator in the nisin activity assay, was incubated in LB solid medium at 37°C. The media for agar diffusion in the nisin activity assay were as follows (wt/vol): tryptone (0.8%), yeast extract (0.25%), glucose (0.5%), Na₂HPO₄ (0.2%), NaCl (0.5%), and agar powder (1.5%). All chemicals were purchased from Sangon Biotech, Shanghai, China.

Construction of overexpressing strains. Primers used in this study are listed in Table 4. *L. lactis* F44 was used for the transformation of pLEB124 with the P45 promoter upstream of the multiple cloning sites. The genes *ddl* and *murF* were amplified from the *L. lactis* F44 genome. PCR was performed using TransStart FastPfu DNA polymerase (TransGen, Beijing, China) using the protocol recommended by the manufacturer. The PCR products were digested with HindIII plus BamHI or with BamHI plus Ncol (NEB, Beijing, China) and were ligated to pLEB124 at the same restriction sites. The recombinant plasmids were transformed into *E. coli* and electrically transformed into *L. lactis* F44, generating the overexpressing strains F44-*ddl* and F44-*murF*. The sequences were confirmed by DNA sequencing (Genewiz, Suzhou, China).

Acid resistance assay. The acid resistance assay was performed as described in a previous study (8). The F44 strains were incubated in the seed medium (control) and the seed medium with the p-amino acid addition (experimental samples) for three generations before being used in acid tolerance assays. After *L. lactis* F44 grew to the mid-log phase, cells were centrifuged (2,604 × g, 5 min), washed twice with normal saline, and then exposed to the pH 3.0 seed medium for 1.5 h or 2.0 h. The number of viable cells was detected via plating on agar plates after proper dilution. Plates were incubated at 30°C for 24 h. The number of viable cells before the acid shock was defined as the T_0 (0 h) prestress cell number (CFU). The number after the acid shock was defined as the $T_{1.5}$ or T_2 poststress cell number [CFU]/ T_0 prestress cell number [CFU]) × 100. The relative survival rate is the survival rate of the experimental sample divided by the survival rate of the control.

TABLE 4	Primers	used in	this	study	/
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Primer	Sequence
ddl-F	CCCAAGCTTATGCTGTCAGTAAAAAGAAGTG
ddl-R	CGCGGATCCAAATCTGCCAGTTGCGAC
murF-F	CGCGGATCCACGATTGCTGACAGATTTTTT
<i>murF-</i> R	CATGCCATGGTCTGCTTTCTCCAAATCGTAA
p124-F	AGGGAACCTAGAATAGTGAA
p124-R	TTCATTCTGCTAACCAGTAAGGC

Fermentation and nisin titer assay. Samples were taken every 2 h to measure the extracellular pH and the optical density at 600 nm (OD_{600}). Nisin titers were determined using the agar diffusion method (9). The stock solution of nisin ($10^6 IU/mI$) was prepared by adding standard nisin (2.5%; balance, sodium chloride; Sigma, St. Louis, MO) to 0.02 M HCl and boiling for 5 min. The stock solution was diluted by adding 0.02 M HCl to standard nisin solutions (50, 100, 200, and 500 IU/mI). Every 2 h, $500 \,\mu$ l of fermentation liquid was added to the same volume of 0.02 M HCl. After boiling for 5 min and centrifugation ($8,228 \times g$ for 5 min, the supernatant was serially diluted with 0.02 M HCl. The nisin titer assay was performed as described previously (9). Meanwhile, $100-\mu$ l fermentation samples were serially diluted in the normal saline and spread on the seed medium plate. After cultivation at 30° C, the number of viable cells was counted. Each sample was tested in triplicate. The nisin yield per cell was calculated as the nisin titer divided by the number of viable cells.

Fluorescent detection. A fluorescent vancomycin derivative (Van-FL) was prepared as in a previous study (37). The 500- μ l vancomycin solution (Sigma; 10 mg/ml in water) was mixed with 50 μ l 5(6)-carboxyfluorescein-*N*-hydroxy succinimide ester (Fluos [Sigma]; 5 mg/ml in dimethyl sulfoxide) and kept at 4°C overnight. After this solution was mixed with 450 μ l 0.1 M Tris (pH 8.0) and reheated for 1 h at 22°C, the resulting solution was stored at –20°C away from light.

The *L. lactis* culture was diluted to an OD₆₀₀ of about 0.8 and inoculated into the seed medium. Then Van-FL was added to the seed medium, with the concentrations ranging from 1 ng/ml to 500 μ g/ml. The OD₆₀₀ was measured every 2 h with a microplate reader (Thermo Fisher Scientific, Vantaa, Finland). The highest concentration tolerated (at which no visible inhibition of growth occurred) was determined by the growth curve (see Fig. S4 in the supplemental material).

Van-FL was added to the seed medium at a final concentration of 0.3 μ g/ml (30°C for 15 to 20 min) when *L. lactis* F44 grew to stationary phase. After centrifugation to remove the supernatant (8,228 × *g*, 5 min), the pellet was washed with precooled phosphate buffer solution (pH 7.0) three times. A 20- μ l sample was placed on the microscope slide with 30 μ l of anti-fluorescence quencher. Images were taken with a confocal laser scanning microscope (LSM 880; Carl Zeiss AG, Oberkochen, Germany). The digital images were analyzed with Carl Zeiss software version 2.3.

HPLC coupled with quadrupole time of flight MS analysis of PG. PG was extracted as previously described (10). A 500-ml portion of *L. lactis* F44 cells at stationary phase were harvested by centrifugation (7,500 × g, 10 min, 4°C). The pellets were resuspended in sterile water and boiled for 10 min. Samples were centrifuged at 40,000 × g for 10 min at 4°C, resuspended in 5% sodium dodecyl sulfate (SDS) solution, and boiled and stirred for 25 min. The previous steps were repeated with 4% SDS solution, and then the pellets were washed with ultrapure water to remove SDS. The pellets were treated with 2 mg/ml pronase for 90 min at 40°C and then with 200 μ g/ml trypsin for 16 h at 37°C. After centrifugation, they were resuspended in 40% hydrofluoric acid (16 h, 4°C). Then they were washed with Tris-HCI (0.25 M, pH 7.0) and ultrapure water several times. The final pellets were freeze-dried and stored at -20° C.

After muramidase digestion (18 h) and filtration, the solutions were determined by high-performance LC-MS (HPLC-MS; Agilent 1200 HPLC system; Agilent, Santa Clara, CA) using a C_{18} column (3.5 μ m; 100 \times 2.1 mm; Waters, Ireland) at a flow rate of 0.2 ml/min. For the flow phase, buffer A was 0.1% formic acid and buffer B was 100% acetonitrile. Parameters were as follows: 0 to 63 min, 95% to 80% A; 63 to 83 min, 80% to 10% A; 83 to 100 min, 10% to 95% B. Muropeptide peaks were collected by electrospray ionization–tandem mass spectrometry (ESI–MS/MS) by using a mass spectrometer (micrOTOF-QII; Bruker Daltonics, Hamburg, Germany). The parameters were set as previously described (10).

Statistical analysis. Three independent experiments, each containing three replicates, were performed in the acid resistance experiment. To evaluate the statistical significance of the survival rate under acid stress, a *t* test was carried out.

SUPPLEMENTAL MATERIAL

Supplemental material is available online only. **SUPPLEMENTAL FILE 1**, PDF file, 0.7 MB.

ACKNOWLEDGMENTS

This study was supported by the Funds for Creative Research Groups of China (21621004), the National Key R&D Program of China (2017YFD0201405), and the National Natural Science Foundation of China (31770076). Jianjun Qiao was supported by the 131 innovative personnel training project of Tianjin (China) and the New Century Outstanding Talent Support Program, Education Ministry of China.

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