

Table S1. Growth of 31 strains of lactic acid bacteria isolated from traditional fermented products in MRS broth at 37°C for 18 h as shown with viable bacterial counts ($n = 3$, $\bar{x} \pm SD$).

Strain	Viable bacterial count(log cfu/ml)	Strain	Viable bacterial count(log cfu/ml)	Strain	Viable bacterial count(log cfu/ml)
<i>Lpb. pentosus</i> PC11	10.35±0.27	<i>Lpb. plantarum</i> P6	9.65±0.15	<i>Lpb. pentosus</i> PC1	6.93±0.10
<i>Lpb. plantarum</i> NMGL2	10.19±0.15	<i>Lcb. casei</i> JS2	9.61±0.09	<i>Lpb. plantarum</i> NMGL4	6.86±0.23
<i>Lpb. plantarum</i> P4	10.18±0.04	<i>Lcb. casei</i> JS1	9.57±0.20	<i>Lab. brevis</i> JS5	6.37±0.39
<i>Lpb. plantarum</i> NMGL1	10.14±0.15	<i>Lab. brevis</i> PC9	8.94±0.53	<i>Lpb. pentosus</i> PC13	6.21±0.27
<i>Lpb. plantarum</i> P3	10.10±0.03	<i>Lab. brevis</i> PC3	8.76±0.17	<i>Lpb. pentosus</i> PC4	6.15±0.51
<i>Lpb. plantarum</i> NMGL3	10.07±0.07	<i>Lab. brevis</i> PC2	8.70±0.15	<i>Lpb. plantarum</i> NMGL5	5.98±0.05
<i>Lpb. plantarum</i> PC10	10.06±0.06	<i>Lab. brevis</i> PC6	8.68±0.13	<i>Lab. brevis</i> JS3	5.84±0.31
<i>Lpb. plantarum</i> P2	9.88±0.07	<i>Lab. brevis</i> PC5	8.60±0.24	<i>Lab. brevis</i> JS4	5.19±0.52
<i>Lpb. plantarum</i> P5	9.82±0.07	<i>Lab. brevis</i> PC7	8.47±0.20	<i>Lpb. plantarum</i> PC12	5.18±0.43
<i>Lpb. plantarum</i> P7	9.79±0.20	<i>Lab. brevis</i> PC8	8.45±0.20		
<i>Lpb. plantarum</i> P1	9.71±0.30	<i>Lpb. pentosus</i> P7	7.02±0.20		

Table S2: The gene sequence of *Lpb. plantarum* NMGL2

GGGAGGGCGCGTGCTATAATGCAGTCGAACGAACTCTGGTATTGATTGGTGCTTGCATCATGATTACATTTG
AGTGAGTGGCGAACTGGTGAGTAACACGTGGGAAACCTGCCAGAAGCGGGGGATAACACCTGGAAACA
GATGCTAATACCGCATAACAACCTGGACCGCATGGTCCGAGTTTGAAAGATGGCTTCGGCTATCACTTTTGA
TGGTCCCGCGGCGTATTAGCTAGATGGTGGGGTAATGGCTCACCATGGCAATGATACGTAGCCGACCTGAGA
GGGTAATCGGCCACATTGGGACTGAGACACGGCCCAAACCTCTACGGGAGGCAGCAGTAGGGAATCTTCC
ACAATGGACGAAAGTCTGATGGAGCAACGCCGCGTGAGTGAAGAAGGGTTTCGGCTCGTAAACTCTGTT
GTTAAAGAAGAACATATCTGAGAGTAACTGTTTCAGGTATTGACGGTATTTAACCAGAAAGCCACGGCTAACT
ACGTGCCAGCAGCCGCGGTAATACGTAGGTGGCAAGCGTTGTCCGGATTTATTGGGCGTAAAGCGAGCGCA
GGCGGTTTTTTAAGTCTGATGTGAAAGCCTTCGGCTCAACCGAAGAAGTGCATCGGAAACTGGGAACTT
GAGTGCAGAAGAGGACAGTGGAACCTCCATGTGTAGCGGTGAAATGCGTAGATATATGGAAGAACACCAAGTG
GCGAAGGCGGCTGTCTGGTCTGTAACCTGACGCTGAGGCTCGAAAGTATGGGTAGCAAACAGGATTAGATAC
CCTGGTAGTCCATACCGTAAACGATGAATGCTAAGTGTTGGAGGGTTTCGCCCTTCAGTGCTGCAGCTAAC
GCATTAAGCATTCCGCCTGGGGAGTACGGCCGCAAGGCTGAAACTCAAAGGAATTGACGGGGGCCCGCAC
AAGCGGTGGAGCATGTGGTTTAATTCGAAGCTACGCGAAGAACCCTTACCAGTCTTGACATACTATGCAAAT
CTAAGAGATTAGACGTTCCCTTCGGGGACAATGGATACAGTGGTGCATGGTTGTGTCGTCAGCTCGTGTCGTGA
GATGTTGGATAAGTCCGCAACGAGCGCAACCTTATTATCAGTGCAGCATAAGTGGCACTCTGGTGAGACTGC
CGTGACAACCGAGAGTGGGATGACGTCAATCATCATGCCTATGACTGCTAACACGTGCTACATGATGGTACA
CGAGTTGCGAACCTCGCGAAG

Table S3. The expression levels of low temperature and acid-resistant proteins in *Lpb. plantarum* NMGL2 in fermented milk stored at low temperature (4°C) for days 1, 7, 14, and 21.

	Day 1 Quantity			Day 7 Quantity			Day 14 Quantity			Day 21 Quantity		
				1218.94			31887.7 32372.9 34887.9 132057.			127528. 132707.		
dnaA	0	0	0	0	2	1233.82	4	5	3	1	6	8
	3805.29 3795.26 4291.25 4256.37 6119.11						92424.6 92574.8 99434.8 359405.			346913. 367442.		
dnaJ	9	4	5	4	4	3727.16	1	1	7	6	9	2
	34755.2 36828.5 33994.8 38748.5 42968.2 40247.7			755230. 768227. 828651.								
dnaK	2	5	2	7	9	6	3	7	6312466230474593179973			
	7625.47 8183.75 7353.17 7529.26 5828.14 163949.						192376. 703937. 726612. 753248.					
dnaN	9623.74	7	8	1	9	5	5	162473	7	8	1	8
							20836.3 22556.7 23097.5 76416.5 81724.6 81986.8					
dnaX	0	0	0	0	0	0	8	8	1	1	3	3
							68010.1			260744. 281193.		
grpE	0	0	0	0	0	0	6	68599.8 72019.2	5 276313 4			
	95403.9 99527.3 99294.3 95845.4 97333.3 99768.4											
groEL	7	6	1	7	6	3230864422405592507343832035081154938557151						
	3972.36 4539.80 4723.00 1517.15 122.257						70985.2 67682.2 81461.5 289681. 288781. 296821.					
rbfA	6	2	1	4	6	0	9	6	3	4	7	1
							7679.88 8889.86 154467. 147795. 165678. 343556. 346412.					
hsp1	0	0	0	8906.93	2	8	9	5	6	7	6	368582
	1762.36 1339.54 2306.95 1915.67 1497.80 1270.72			44610.6 43775.5 45799.5 160636. 176769. 166639.								
hsp2	9	3	7	7	6	5	4	3	5	4	6	1
	4972.72						156180. 150562. 169457. 533801. 522632. 559783.					
hsp3	0	0	7	0	0	0	1	7	5	8	7	6
murA							486.231 35503.4 31435.0 35458.4 171943. 180826. 173624.					
1	0	0	0	0	0	8	9	5	5	8	2	3
							12577.4 14110.9 15368.5			54802.6 56418.3		
murB	0	0	0	0	0	0	2	1	6	55851.3	8	1
							860.301 2584.85 33104.5 34521.5 37088.1 102248. 111729. 113298.					
murC	0	0	0	0	1	7	1	6	9	6	4	4
	2000.11 2035.17 1883.12 2404.64			1785.34 36012.5 38024.6 42767.0 119724. 127224. 132681.								
murD	5	3	6	6	2664.79	6	1	1	7	6	4	1
	1169.10			1942.33			63393.4 63811.3 71858.8 232787. 216100. 233090.					
murE1	5	0	0	0	7	1676.99	4	6	1	8	2	6
	1471.11 1465.11			1414.16 43052.0 44647.6 47934.7 162139. 164235.								
murF	3	1	0	0	0	8	4	2	9	7	7	172024
	2900.81			3875.99 5337.40 5686.64			228692. 224307. 258873. 288471. 297987. 298654.					
dacA1	7	3415.7	4	4	5	6326.56	5	6	1	6	6	1
							29276.5 27690.8			35962.2 34832.1		
dacB	0	0	0	0	0	0	9	6	32534.3	5	2	39989.3
	40200.5 36166.3 27443.0 52015.4 52309.0 61281.3											
cspC	4	9	2	4	2	7169643018222982056945284743827294813183768						
	4372.31 4114.33 4376.35 3086.68			3683.66 97415.9 97131.8 110254. 327683. 333298. 335478.								
gaLE1	8	9	6	5	3654.22	9	6	2	9	7	2	4

	gaLE2	12973.3	13678.5	8181.83	3962.85	8698.77	301711.	298306.	324952.											
		13581.7	4	9	4	7	8	7	4	81095517	10626971	145028								
		25288.1	25348.0	24169.0	12638.3	12480.8	11070.1	487946.	464839.	531454.										
	gaLK	6	2	4	1	1	4	2	8	421266852	1945262	260797								
	gaLM							9993.96	9949.47	15476.9	42983.9	47165.3								
	1	0	0	0	0	0	0	7	2	746253.9	6	1								
	gaLM	7341.62		7873.89	3466.31	3954.18	4624.28	237730.	235703.	264347.	736947.									
	3	5	6477.87	1	3	1	1	3	3	5	735577	699845	8							
		25027.1	25950.5	25558.9	23531.4	24250.9	24761.4	507787.	565960.											
	gaLU	9	9	6	5	8	1	9	490675	12073073	1907866	2013499								