

Supplementary tables

Supplementary Table S1. The PCR conditions for the identification of suspected pathogens

Pathogens	Target gene	Size	Sequence (5' to 3')	Reaction Materials Final Volume: 25 ul	PCR Condition	Reference
<i>Acinetobacter bau- mannii</i>	ITS region	208	p-Ab-ITSF: 5'-CATTATCACGGTAATTAGTG-3' p-AbI-TSB: 5'-AGAGCACTGTGCACTTAAG-3'	DNA: 300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 94°C 5 min Denaturation: 94°C 30s Annealing: 55°C 30s Extension: 72°C 30s D.A.E. Cycles: 30 cycles Final extension: 72°C 7 min	[1]
<i>Bacillus cereus sensu lato s.l</i>	<i>Bal</i>	533	BalF: 5'-TGCAACTGTATTAGCACAAGCT-3' BalR: 5'-TACCACGAAGTTTGTTCACTACT-3'	DNA: 100-300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 95°C 5 min Denaturation: 94°C 45s Annealing: 55°C 45s Extension: 72°C 45s D.A.E. Cycles: 30 cycles Final extension: 72°C 5 min	[2]
<i>Clostridium difficile</i>	<i>tcdA</i> <i>tcdB</i> <i>cdtA</i> <i>cdtB</i>	632, 441, 260, 179	tcdA-F 5'-GTATGGATAGGTGGAGAAGTCAGTG-3' tcdA-R 5'-CGGTCTAGTCCAATAGAGCTAGGTC-3' tcdB-F 5'-GAAGATTTAGGAAATGAAGAAGGTGA-3' tcdB-R 5'-AACCACTATATTCAACTGCTTGTCC-3' cdtA-F 5'-ATGCACAAGACTTACAAAGCTATAGTG-3' cdtA-R 5'-CGAGAATTTGCTTCTATTTGATAATC-3' cdtB-F 5'-ATTGGCAATAATCTATCTCCTGGA-3' cdtB-R 5'-CCAAAATTTCCACTTACTTGTGTG-3'	DNA: 100 ng Primer: 25 nM tcdA FR, 10 nM tcdB FR, 200 nM,cdtA FR 500 nM,cdtB FR Master mix: 5 ul	Pre-denaturation: 93°C 2 min Denaturation: 93°C 20s Annealing: 60°C 65s Extension: 68°C 70s D.A.E. Cycles: 30 cycles Final extension: 68°C 5 min	[3]

<i>Klebsie pneumoniae</i>	<i>tyrB</i>	931	tyrB-F: 5'-GGCTGTACTACAACGATGAC-3' tyrB-R: 3'-TTGAGCAGGTAATCCACTTTG-3'	DNA: 300 ng Primer: 500 nM Master mix: 5 ul	Pre-denaturation: 95°C 5 min Denaturation: 95°C 60s Annealing: 55°C 60s Extension: 72°C 60s D.A.E. Cycles: 35 cycles Final extension: 72°C 10 min	[4]
<i>Staphylococcus aureus</i>	<i>nuc</i>	270	nuc-F 5'-GCGATTGATGGTGATACGGTT-3' nuc-R 5'-AGCCAAGCCTTGACGAACTAAAGC-3'	DNA: 300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 95°C 5 min Denaturation: 94°C 60s Annealing: 55°C 60s Extension: 72°C 60s D.A.E. Cycles: 30 cycles Final extension: 72°C 10 min	[5]
<i>Pseudomonas aeru- ginosa</i>	16S rRNA	956	PA-SS-F: 5'-GGGGGATCTTCGGACCTCA-3' PA-SS-R: 5'-TCCTTAGAGTGCCCCACCCG-3'	DNA: 300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 95°C 2 min Denaturation: 94°C 20s Annealing: 58°C 20s Extension: 72°C 40s D.A.E. Cycles: 25 cycles Final extension: 72°C 4 min	[6]

<i>Streptococcus pneumoniae</i>	<i>pneumolysin</i>	75	F 5'-AGCGATAGCTTTCTCCAAGTGG-3' R 5'-CTTAGCCAACAAATCGTTACCG-3'	DNA: 300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 94°C 5 min Denaturation: 94°C 45s Annealing: 50°C 90s Extension: 72°C 90s D.A.E. Cycles: 35 cycles Final extension: 72°C 10 min	[7]
<i>nontuberculous mycobacteria, NTM</i>	16s rRNA 23S rRNA		Real-Q MTB & NTM Kit	DNA: 300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 95°C 10 min Denaturation: 95°C 15s Annealing & Extension: 67°C 45s D.A.E. Cycles: 45 cycles	[8]
<i>Stenotrophomonas maltophilia</i>	23S rRNA	278	F 5'GCTGGATTGGTTCTAGGAAAACGC3' R 5'ACGCAGTCACTCCTTGCG3'	DNA: 300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 94°C 5 min Denaturation: 94°C 45s Annealing: 68°C 45s Extension: 72°C 45s D.A.E. Cycles: 35 cycles Final extension: 72°C 10 min	[7]

<i>Gardmerella vaginalis</i>	16S rRNA		F- GV1 5'-TTACTGGTGTATCACTGTAAGG-3' R- GV3 5'-CCGTCACAGGCTGAACAGT-3'	DNA: 300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 94°C 5 min Denaturation: 94°C 45s Annealing: 55°C 45s Extension: 72°C 45s D.A.E. Cycles: 35 cycles Final extension: 72°C 10 min	[9]
<i>Chryseobacterium indologenes</i>	6S-23S rRNA ISR	561	Chry-16S-F 5'-GCATCAGCCATGGCGCGGTG-3' Chry-23S-R 5'-CCTTAACGATTTCTTTCCTA-3'	DNA: 300 ng Primer: 400 nM Master mix: 5 ul	Pre-denaturation: 94°C 5 min Denaturation: 94°C 45s Annealing: 55°C 45s Extension: 72°C 45s D.A.E. Cycles: 35 cycles Final extension: 72°C 10 min	[10]

Supplementary Table S2. Chemical analysis of fermented fruits from the 2nd stage of fermentation and final products

Process	Samples	pH ¹	°Brix ² (%)	Organic acid ³ (%)	Ethanol ⁴ (%)	Measurement time
	Ch1_2 nd	3.2	24.0	3.8	0.0	1 st month
	Ch2_2 nd	3.2	23.3	3.6	0.0	
	Ch3_2 nd	3.4	22.5	3.4	0.0	
	Ch1_2 nd	3.2	23.6	4.1	0.0	2 nd month
	Ch2_2 nd	3.4	23.2	3.7	0.0	
	Ch3_2 nd	3.3	22.2	3.5	0.0	

2 nd stage of fermentation	Ch1_2 nd	3.3	23.8	3.8	0.0	3 rd month
	Ch2_2 nd	3.4	23.2	3.6	2.2	
	Ch3_2 nd	3.4	22.2	3.4	2.4	
	Ch1_2 nd	3.3	23.6	4.3	0.0	4 th month
	Ch2_2 nd	3.4	23.2	4.1	0.0	
	Ch3_2 nd	3.4	22.2	3.8	0.7	
	Ch1_2 nd	3.3	23.2	4.4	0.0	5 th month
	Ch2_2 nd	3.2	23.0	4.0	0.0	
	Ch3_2 nd	3.3	22.0	4.0	0.0	
	Ch1_2 nd	3.2	23.4	4.3	0.0	6 th month
	Ch2_2 nd	3.3	23.0	4.0	1.7	
	Ch3_2 nd	3.3	22.0	3.9	1.4	
Product	Ch1_2 nd	3.2	23.4	4.3	0.0	7 th month
	Ch2_2 nd	3.3	23.0	4.1	1.1	
	Ch3_2 nd	3.3	22.0	4.0	0.0	
	Ch1_2 nd	3.2	22.8	4.5	0.0	8 th month
	Ch2_2 nd	3.3	23.0	4.0	0.0	
	Ch3_2 nd	3.3	22.0	3.9	1.7	
	Pr1_F	2.8	10.8	5.8	2.8	0 day
	Pr2_F	2.6	11.0	5.8	2.7	
	Pr3_F	2.7	11.2	5.9	2.7	

¹ The pH of samples is measured using a DO Microelectronic pH-Vision pH meter (model PHB-9901, Ai-On Industrial Corp., Taipei, Taiwan).

² °Brix (Brix) was measured using Brix meter (Lotus, China) following the manufacturer protocol. Brix is mainly made up of sugars but also includes other compounds.

³ Organic acids content was measured using the principle of acid-base neutralization titration of volumetric analysis. In brief, a 2 ml sample was taken in a beaker with the addition of distilled water to 20 ml (10 times dilution) and titrated with 0.1N NaOH standard solution. The initial and end of titration was recoded and finally, the below formula was used to determine the organic acid contents.

The amount of organic acid in fermented food (%) = (S-B) * F * O.A * D.F / Swt *100

S: The sample test consumed the number of 0.1N NaOH mL

B: The number of 0.1N NaOH mL consumed in the blank test

F: The titer of 0.1N NaOH (acceptable range: 0.9~1.1)

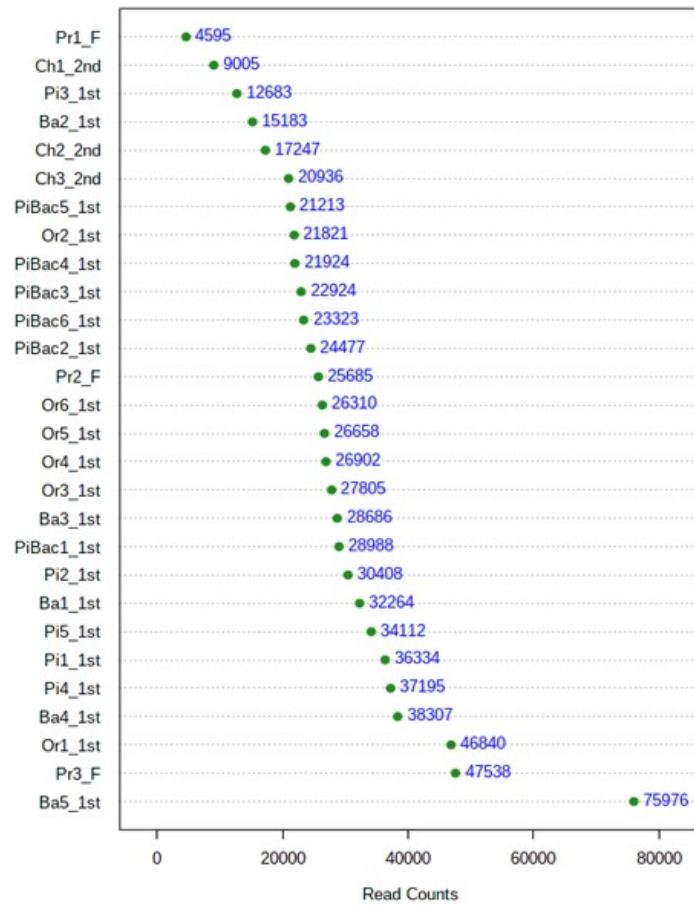
O.A: Organic acid (citric acid based: 0.0064)

D.F: Dilution factor

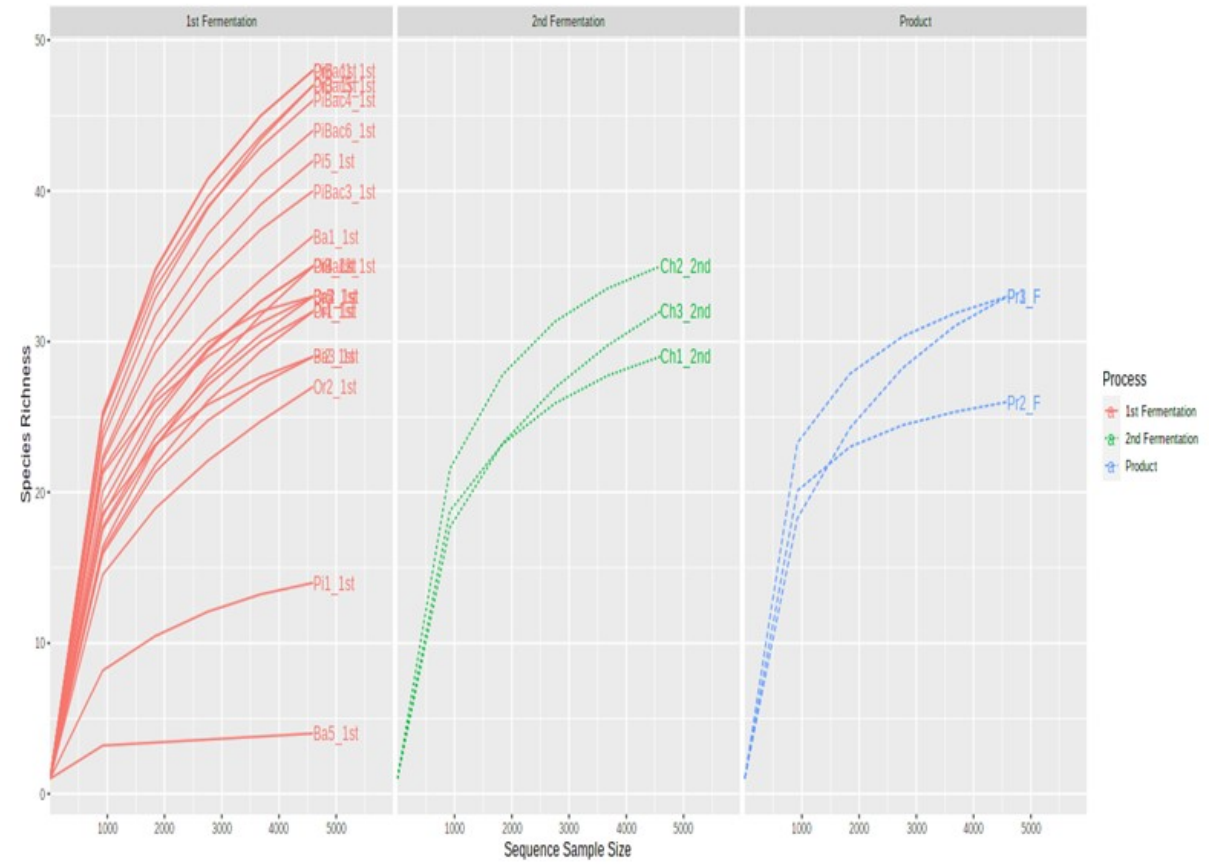
Swt: sample weight (mL)

Additionally, the manufacture company also uses Agilent Hi-Plex H column with UV detection to separate, identify, and estimate organic acids, including oxalic acid, citric acid, tartaric acid, malic acid, succinic acid, formic acid, and fumaric acid, when they need to show the product quality report.

⁴ Ethanol contents were measured using ethyl alcohol Meter (Ethyl alcohol meter PAL-33S, Atago Co., LTD.) following the manufacture protocol.



A



B

Supplementary Figure S1. Library size overview (A) and alpha rarefaction plot (B) among the fermentation stages and the final products.

Supplementary Table S3. The sequence good's coverage at the minimum sequence depth associated with the fermented samples

samples	no. se- quence	goods
Pr1_F	4595	99.89119
Pr2_F	4595	99.93471
Pr3_F	4595	99.8259
Or1_1st	4595	99.8259
Or2_1st	4595	99.76061
Or3_1st	4595	99.6518
Pi- Bac1_1st	4595	99.71708
Pi- Bac2_1st	4595	99.67356
Pi- Bac3_1st	4595	99.76061
Pi1_1st	4595	99.93471
Pi2_1st	4595	99.8259
Ba1_1st	4595	99.69532
Ba2_1st	4595	99.8259
Ba3_1st	4595	99.86942
Ch1_2nd	4595	99.89119
Ch2_2nd	4595	99.86942
Ch3_2nd	4595	99.78237
Or4_1st	4595	99.78237
Or5_1st	4595	99.93471
Or6_1st	4595	99.71708
Pi- Bac4_1st	4595	99.69532

Pi-Bac5_1st	4595	99.6518
Pi-Bac6_1st	4595	99.71708
Pi3_1st	4595	99.78237
Pi4_1st	4595	99.73885
Pi5_1st	4595	99.71708
Ba4_1st	4595	99.73885
Ba5_1st	4595	99.97824

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