

## **SUPPLEMENTAL FILE**

### **Ratio of histamine-producing/non-histamine-producing subgroups of *Tetragenococcus halophilus* determines the histamine accumulation during spontaneous fermentation of soy sauce**

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Running title: Strain specificity related to histamine accumulation

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Supplemental files include:

Supplemental figures: Fig. S1-S7;

Supplemental tables: Table S1-S2;

## Supplementary figures:

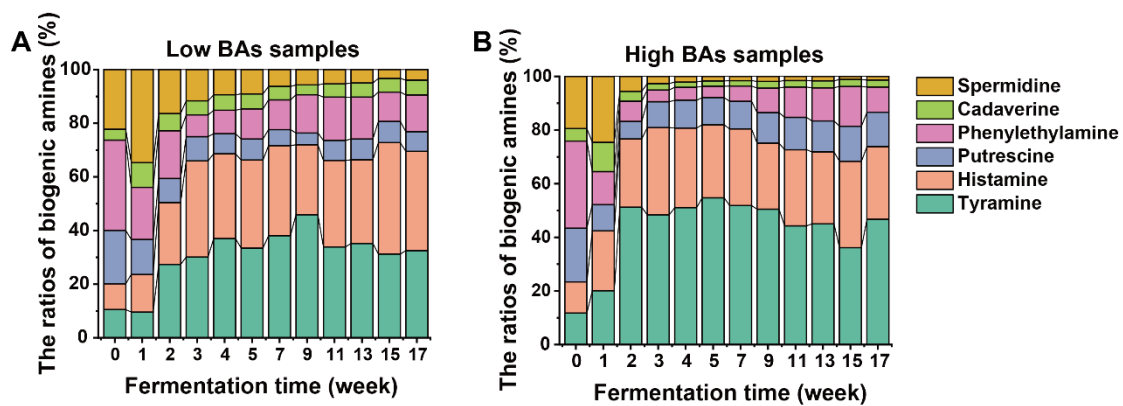


Fig. S1 The profiles of diverse BAs production in low-BAs soy sauce samples (A) and high-BAs soy sauce samples (B).

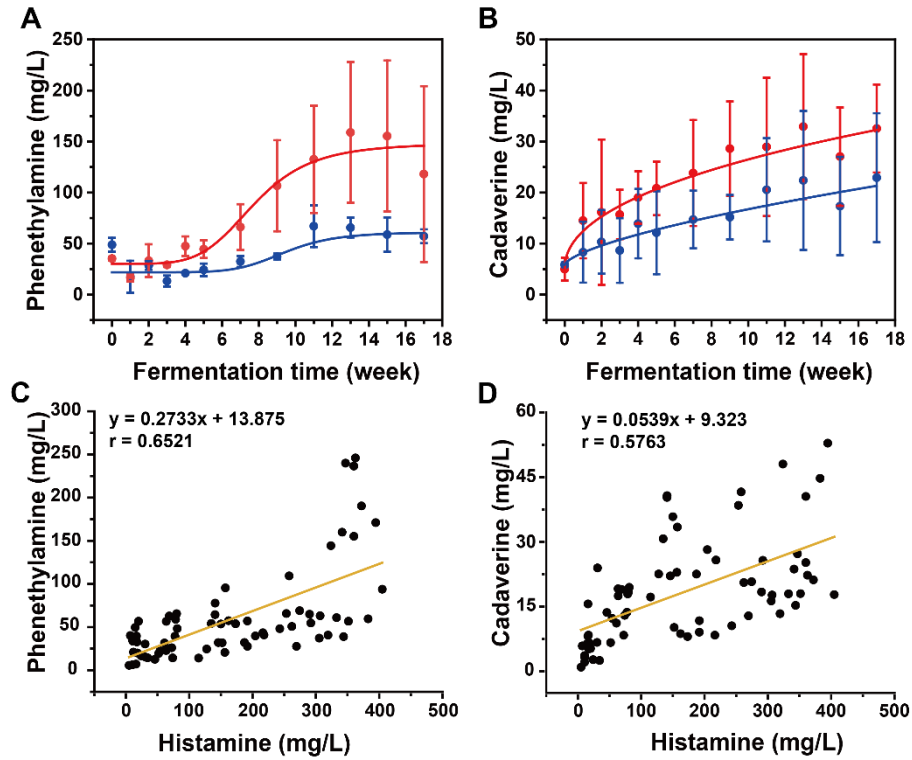


Fig. S2 Biogenic amines during the soy sauce fermentation. Dynamic of biogenic amines accumulation in high- (red line) and low- (blue line) BAs samples (A) Phenethylamine, (B) Cadaverine. Correlation analysis of phenethylamine (C) and cadaverine (D) with histamine. Data are indicated as the average of three samples  $\pm$  standard deviation.

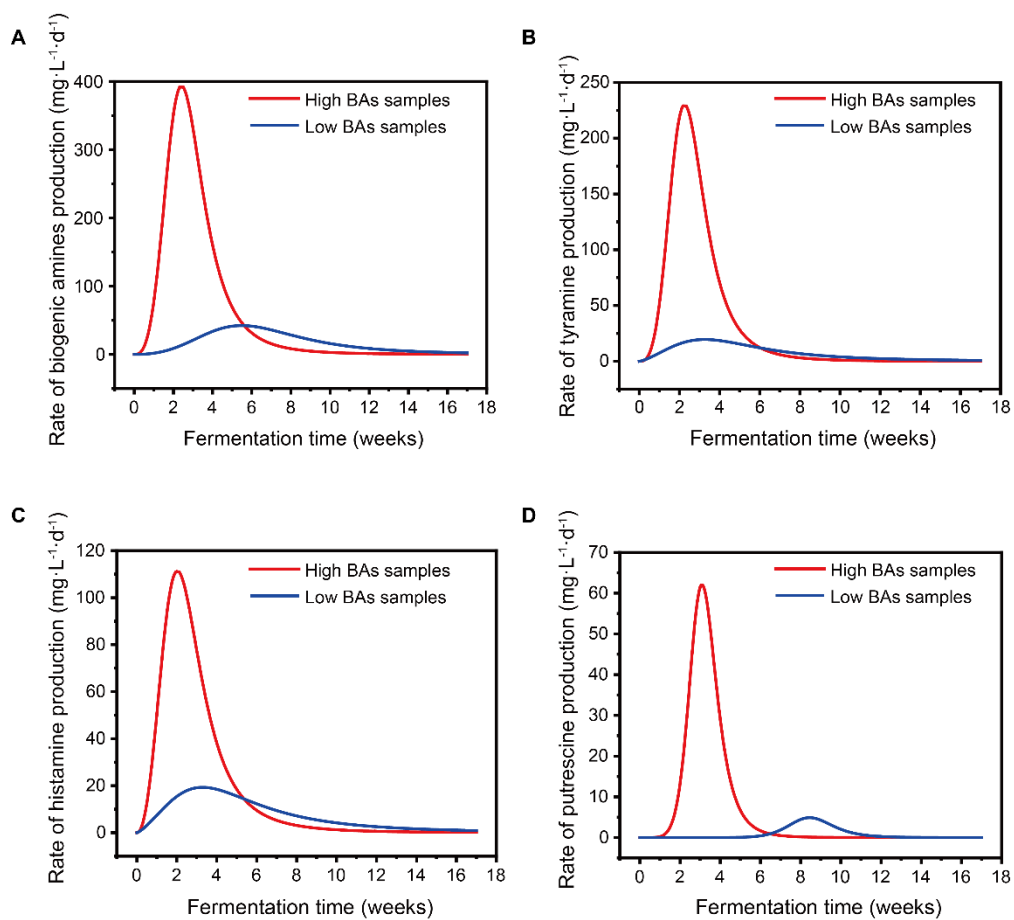


Fig. S3 BAs production rates during soy sauce fermentation. (A) The total of biogenic amines; (B) Tyramine; (C) Histamine; (D) Putrescine.

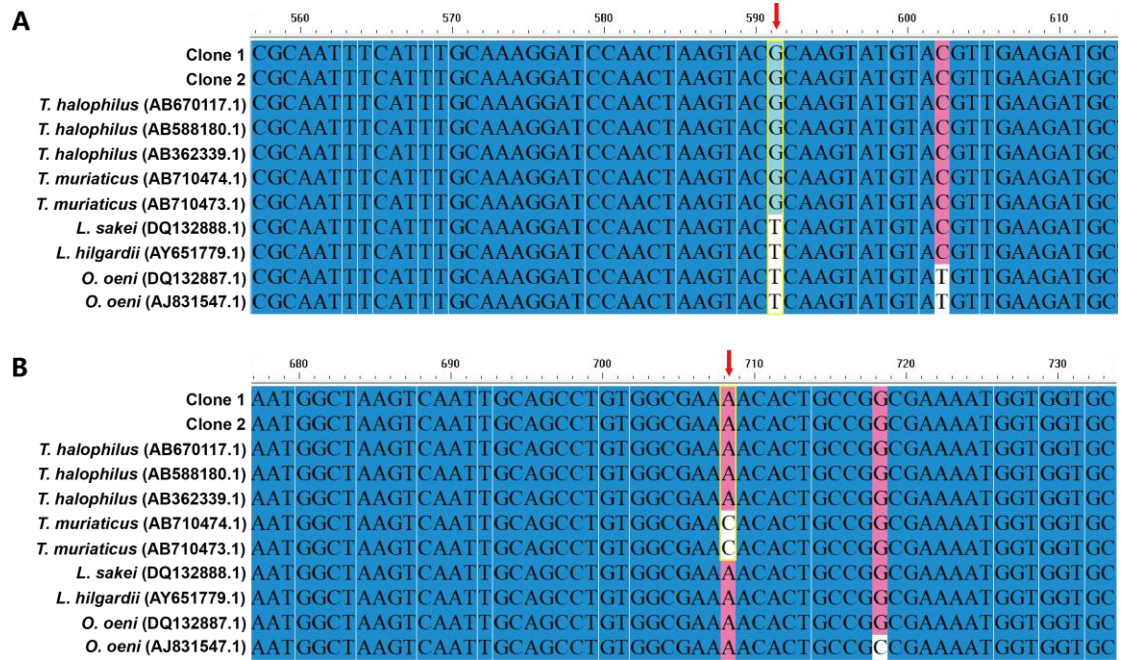


Fig. S4 Partial clone sequence alignment of *hdcA* gene clone library. The red arrows show the key differential bases of the different sequences. Note: Reference genomes from the same species retained only samples with sequence differences.

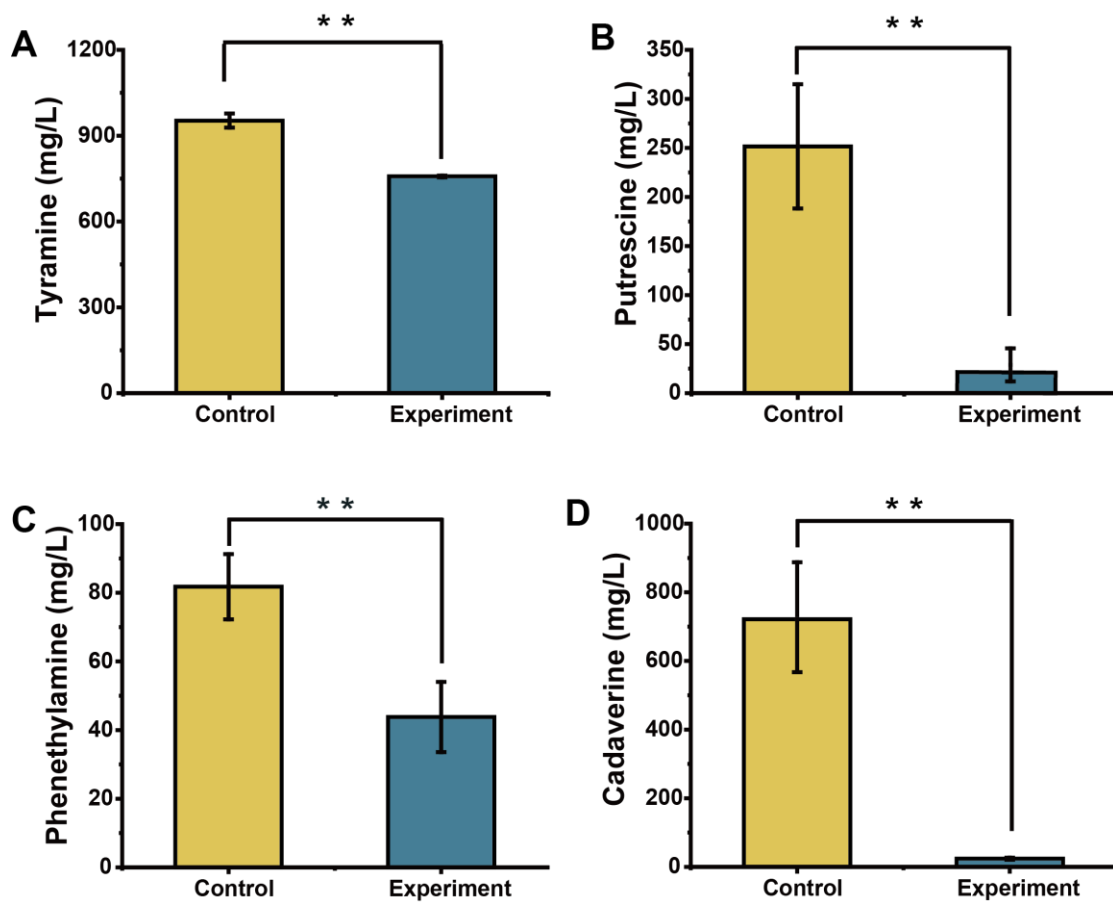


Fig. S5 Four BAs control by artificially modifying the ratio of histamine-producing to non-histamine-producing *T. halophilus* strains during soy sauce fermentation. (A) Tyramine; (B) Putrescine; (C) Phenethylamine; (D) Cadaverine. ANOVA: \*\* means  $p < 0.01$

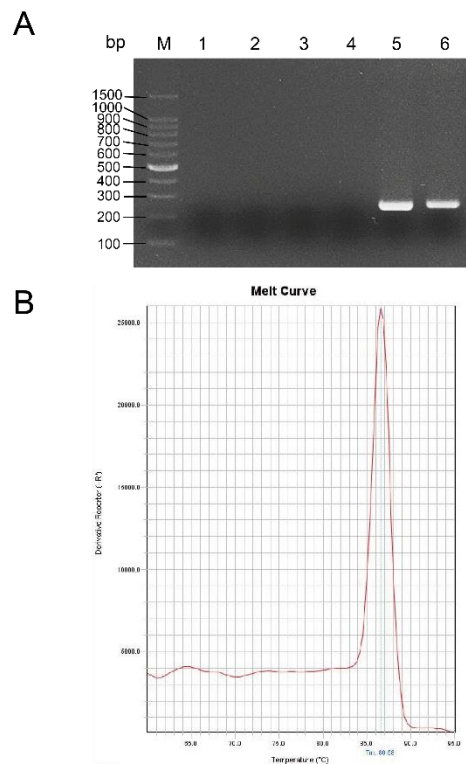


Fig. S6 PCR and qPCR were used to verify the specificity of the primers TetRT-F1 and TetRT-R1, which was used for the quantitative of *T. halophilus*. A: 16 S rRNA amplification of 6 kinds of dominant soy sauce bacteria by using primers TetRT-F1 and TetRT-R1. M. marker; 1. *Weissella paramesenteroides*, 2. *Weissella cibaria*, 3. *Weissella viridescens*, 4. *Weissella confuse*, 5. Histamine-producing *T. halophilus*, 6. Non-histamine-producing *T. halophilus*. B: Melt curve of qPCR.



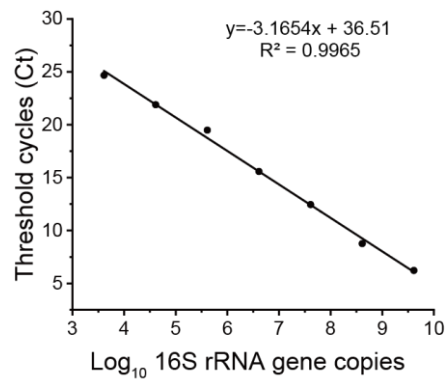


Fig. S7 Real-time quantitative PCR (qPCR) analysis of 10-fold serial dilutions of control plasmid DNA. Cycle threshold values (Ct) are plotted against the calculated plasmid copy number. The equation and R2 value of the regression line are indicated.

## Supplementary tables:

Table 1 Maximum histamine production capacity of the reported histamine-producing microorganisms.

Histamine-producing Microorganisms	Maximum production of histamine (mg/L)	Culture Conditions	References
<i>Morganella morganii</i>	6000	TSB medium	(1)
<i>Tetragenococcus muriaticus</i>	4870	MRSB medium	(2)
<i>Photobacterium damsela</i> subsp. <i>Damsela</i>	3670	TSB medium	(1)
<i>Klebsiella pneumoniae</i>	3416.43	HDB medium	(3)
<i>Photobacterium phosphoreum</i>	3140	TSB medium	(1)
<i>Tetragenococcus halophilus</i>	3000	HB medium	(4)
<i>Morganella psychrotolerans</i>	3000	Canned tuna (fish)	(5)
<i>Klebsiella oxytoca</i>	2231	Broth	(6)
<i>Staphylococcus epidermidis</i>	1945.9	HEB medium	(7)
<i>Photobacterium kishitanii</i>	1545	LSW-70 broth	(8)
<i>Staphylococcus capitis</i>	1260	TSB medium	(9)
<i>Klebsiella aerogenes</i>	1000	Histidine broth	(10)
<i>Streptococcus thermophilus</i>	540	LM17 medium	(11)

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<i>Lentilactobacillus saerimneri</i>	522.41	MRSB medium	(12)
<i>Lentilactobacillus parabuchneri</i>	470.16	MRSB medium	(13)
<i>Lentilactobacillus buchneri</i>	452.38	MRSB medium	(12)
<i>Lentilactobacillus hilgardii</i>	235	H-MDBmod medium	(14)
<i>Oenococcus oeni</i>	99	H-MDBmod medium	(14)
<i>Clostridium perfringens</i>	41.1	Decarboxylating broth	(15)

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Table S2 Histamine production of *T. halophilus* in MRS medium containing 1% histidine and soy sauce simulated fermentation system

NO.	Strain	source	<i>hdcA</i> gene	MRS+1% histidine (mg/L)	Simulated soy sauce fermentation (mg/L)
1	<i>T. halophilus</i> H178	High-BAs samples	+	1287.48	555.85
2	<i>T. halophilus</i> H128	High-BAs samples	+	1286.32	609.61
3	<i>T. halophilus</i> H168	High-BAs samples	+	1241.23	655.97
4	<i>T. halophilus</i> L64	Low BAs samples	+	1221.43	635.27
5	<i>T. halophilus</i> H181	High-BAs samples	+	1220.83	620.39
6	<i>T. halophilus</i> H189	High-BAs samples	+	1209.31	617.33
7	<i>T. halophilus</i> H124	High-BAs samples	+	1207.35	599.12
8	<i>T. halophilus</i> H42	High-BAs samples	+	1202.8	520.63
9	<i>T. halophilus</i> H29	High-BAs samples	+	1200.25	515.77
10	<i>T. halophilus</i> H125	High-BAs samples	+	1184.64	559.59
11	<i>T. halophilus</i> L100	Low BAs samples	+	1181.65	684.25
12	<i>T. halophilus</i> H159	High-BAs samples	+	1170.62	633.9
13	<i>T. halophilus</i> H156	High-BAs samples	+	1168.66	558.12
14	<i>T. halophilus</i> L45	Low BAs samples	+	1164.55	655.06
15	<i>T. halophilus</i> L137	Low BAs samples	+	1153.3	540.84
16	<i>T. halophilus</i> L106	Low BAs samples	+	1151.65	560.61
17	<i>T. halophilus</i> L134	Low BAs samples	+	1151.61	650.03

18	<i>T. halophilus</i> L192	Low BAs samples	+	1146.98	512.07
19	<i>T. halophilus</i> H45	High-BAs samples	+	1145.96	655.45
20	<i>T. halophilus</i> L40	Low BAs samples	+	1125.15	557.12
21	<i>T. halophilus</i> H121	High-BAs samples	+	1116.27	583.21
22	<i>T. halophilus</i> H131	High-BAs samples	+	1113.86	633.19
23	<i>T. halophilus</i> H52	High-BAs samples	+	1110.21	596.89
24	<i>T. halophilus</i> L143	Low BAs samples	+	1108.21	593.93
25	<i>T. halophilus</i> H32	High-BAs samples	+	1104.75	670.82
26	<i>T. halophilus</i> L31	Low BAs samples	+	1103.76	634.41
27	<i>T. halophilus</i> H180	High-BAs samples	+	1091.05	622.49
28	<i>T. halophilus</i> L15	Low BAs samples	+	1088.45	508.68
29	<i>T. halophilus</i> H99	High-BAs samples	+	1085.82	611.1
30	<i>T. halophilus</i> L123	Low BAs samples	+	1081.43	563.37
31	<i>T. halophilus</i> H115	High-BAs samples	+	1074.06	664.31
32	<i>T. halophilus</i> H103	High-BAs samples	+	1061.32	654.11
33	<i>T. halophilus</i> H111	High-BAs samples	+	1037.37	521.32
34	<i>T. halophilus</i> H114	High-BAs samples	+	1034.16	594.84
35	<i>T. halophilus</i> H48	High-BAs samples	+	1023.55	504.65
36	<i>T. halophilus</i> H47	High-BAs samples	+	1008.36	508.14
37	<i>T. halophilus</i> L205	Low BAs samples	+	898.45	612.5

Note: Data for non-histamine-producing strains were not shown.

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