

# Bacteriobiota and Chemical Changes during the Ripening of Traditional Fermented “Pirot ‘Ironed’ Sausage”

Svetlana Bogdanović <sup>1</sup>, Slaviša Stanković <sup>2</sup>, Tanja Berić <sup>2</sup>, Igor Tomasevic <sup>3,4,\*</sup>, Volker Heinz <sup>4</sup>, Nino Terjung <sup>4</sup> and Ivica Dimkić <sup>2,\*</sup>

<sup>1</sup> Agriculture and Food College of Applied Studies, Ćirila i Metodija 1, 18400 Prokuplje, Serbia

<sup>2</sup> Faculty of Biology, University of Belgrade, Studentski trg 16, 11158 Belgrade, Serbia

<sup>3</sup> Faculty of Agriculture, University of Belgrade, Nemanjina 6, 11080 Belgrade, Serbia

<sup>4</sup> DIL German Institute of Food Technologies, Prof.-v.-Klitzing-Str. 7, 49610 Quakenbrueck, Germany

\* Correspondence: tbigor@agrif.bg.ac.rs (I.T.); ivicad@bio.bg.ac.rs (I.D.)

**Citation:** Bogdanović, S.; Stanković, S.; Berić, T.; Tomasevic, I.; Heinz, V.; Terjung, N.; Dimkić, I. Bacteriobiota and Chemical Changes during the Ripening of Traditional Fermented “Pirot ‘Ironed’ Sausage”. *Foods* **2023**, *12*, x. <https://doi.org/10.3390/xxxxx>

Academic Editors: Jasna Mrvčić and Damir Stanzer

Received: 22 December 2022

Revised: 23 January 2023

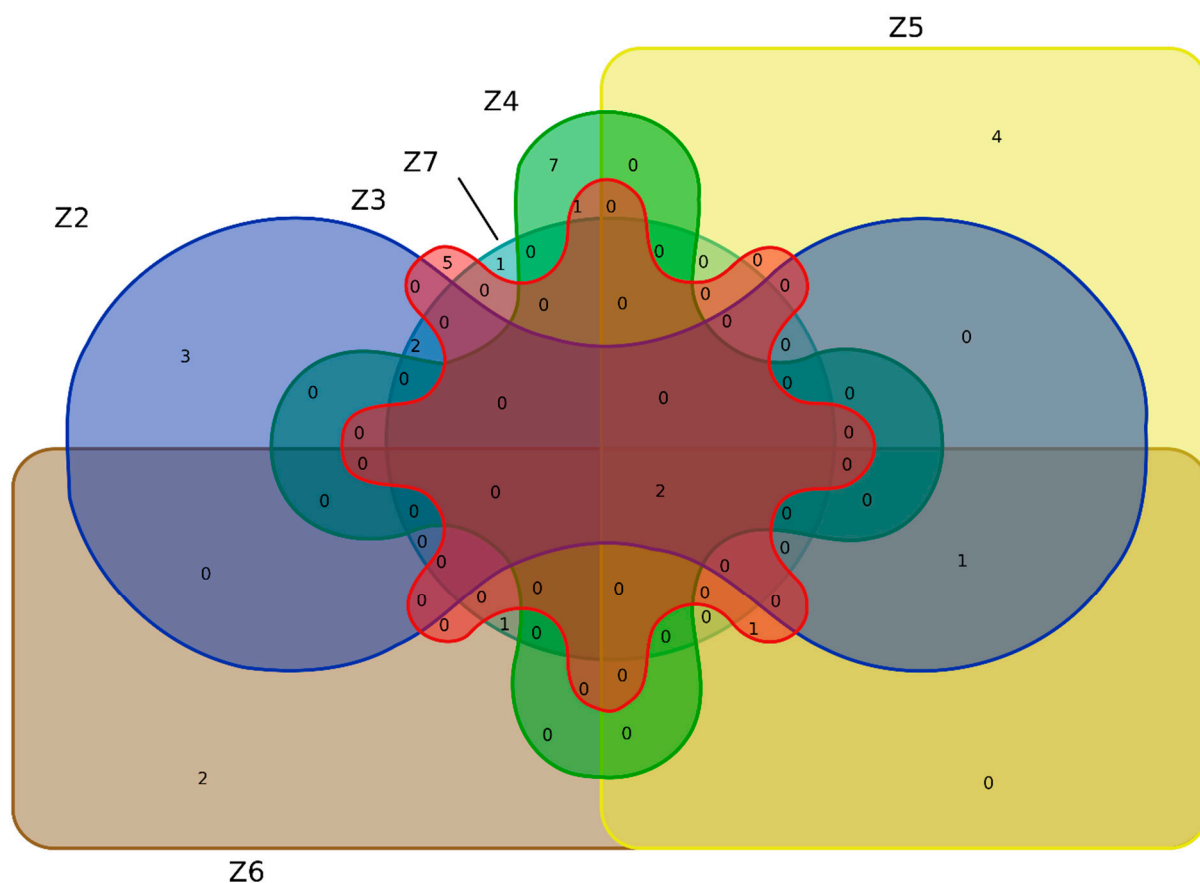
Accepted: 29 January 2023

Published: 3 February 2023



**Copyright:** © 2023 by the authors. Submitted for possible open access publication under the terms and conditions of the Creative Commons Attribution (CC BY) license (<https://creativecommons.org/licenses/by/4.0/>).





**Figure S2.** Venn diagram of common species between samples (Z2-Z6) throughout the ripening of “Pírot ‘ironed’ sausage”.

**Table S1.** A total number of the reads retention.

Sample	Input	Primer remove	Quality filter	Denoised F	Denoised R	Merged	Chimera remove	Final % of reads
Z1	169672	168669	156105	154649	154959	148160	127677	75.25
Z2	127184	117614	109826	109178	109328	105343	96909	76.20
Z3	148238	142399	133327	132649	132808	127148	114558	77.28
Z4	107113	104776	99454	98977	99092	95421	86161	80.44
Z5	166600	162465	152607	151885	152033	145402	128808	77.32
Z6	189000	184973	174132	173236	173503	166585	146679	77.61
Z7	145762	141171	133262	132584	132710	126288	108690	74.57
Total/average	1053569	1022067	958713	953158	954433	914347	809482	76.95

**Table S2.** Homology of the identified OTUs annotated based on the BLAST $n$  best bit score in the NCBI 16S database.

No.	Qseq ID	Sseq ID	Ident. (%)	Length	E value	Bit score	Species name
1	seq_001	gi 1397641705 ref NR_156814.1	100	427	0	789	<i>Photobacterium carnosum</i>
16	seq_004	gi 1397641705 ref NR_156814.1	99.766	427	0	784	
161	seq_033	gi 1397641705 ref NR_156814.1	99.766	427	0	784	
196	seq_040	gi 1397641705 ref NR_156814.1	99.532	427	0	778	
216	seq_044	gi 1397641705 ref NR_156814.1	99.766	427	0	784	
231	seq_047	gi 1397641705 ref NR_156814.1	99.532	427	0	778	
256	seq_052	gi 1397641705 ref NR_156814.1	99.532	427	0	778	
336	seq_068	gi 1397641705 ref NR_156814.1	99.766	427	0	784	
376	seq_076	gi 1397641705 ref NR_156814.1	99.532	427	0	778	
526	seq_106	gi 1397641705 ref NR_156814.1	99.766	427	0	784	
531	seq_107	gi 1397641705 ref NR_156814.1	99.766	427	0	784	
1176	seq_236	gi 1397641705 ref NR_156814.1	100	418	0	773	<i>Photobacterium phosphoreum</i>
1291	seq_259	gi 1397641705 ref NR_156814.1	100	427	0	789	
41	seq_009	gi 631252986 ref NR_114184.1	100	427	0	789	
276	seq_056	gi 631252986 ref NR_114184.1	99.766	427	0	784	<i>Photobacterium iliopiscarium</i>
56	seq_012	gi 631252986 ref NR_114184.1	99.766	427	0	784	
46	seq_010	gi 631250795 ref NR_111990.1	100	427	0	789	
51	seq_011	gi 631250795 ref NR_111990.1	99.766	427	0	784	<i>Photobacterium piscicola</i>
151	seq_031	gi 631250795 ref NR_111990.1	99.532	427	0	778	
406	seq_082	gi 631250795 ref NR_111990.1	99.766	427	0	784	
71	seq_015	gi 672239090 ref NR_125679.1	100	427	0	789	
11	seq_003	gi 631252623 ref NR_113821.1	100	427	0	789	<i>Lactobacillus sakei</i>
441	seq_089	gi 631252623 ref NR_113821.1	99.766	427	0	784	
446	seq_090	gi 631252623 ref NR_113821.1	99.766	427	0	784	
511	seq_103	gi 631252623 ref NR_113821.1	99.766	427	0	784	
716	seq_144	gi 631252623 ref NR_113821.1	99.766	427	0	784	
1876	seq_376	gi 631252623 ref NR_113821.1	100	427	0	789	<i>Lactobacillus algidus</i>
6	seq_002	gi 265678315 ref NR_028617.1	100	427	0	789	
1496	seq_300	gi 265678315 ref NR_028617.1	96.487	427	0	706	
1766	seq_354	gi 265678315 ref NR_028617.1	100	427	0	789	
1661	seq_333	gi 265678315 ref NR_028617.1	100	420	0	776	
121	seq_025	gi 343201712 ref NR_042438.1	99.766	427	0	784	<i>Lactobacillus graminis</i>
366	seq_074	gi 343201712 ref NR_042438.1	99.532	427	0	778	
881	seq_177	gi 343201712 ref NR_042438.1	99.532	427	0	778	
246	seq_050	gi 559795383 ref NR_104976.1	100	427	0	789	<i>Lactobacillus fuchuensis</i>
616	seq_124	gi 672238974 ref NR_125563.1	100	427	0	789	<i>Lactobacillus nenjiangensis</i>
1151	seq_231	gi 645319700 ref NR_117065.1	100	427	0	789	<i>Lactobacillus ultunensis</i>
1506	seq_302	gi 1491505234 ref NR_159082.1	99.766	427	0	784	<i>Lactobacillus allii</i>

1676	seq_336	gi 631253167 ref NR_114365.1	99.063	427	0	767	<i>Lactobacillus iwatensis</i>
1761	seq_353	gi 343198639 ref NR_043148.1	100	427	0	789	<i>Lactobacillus oligofermentans</i>
1816	seq_364	gi 631253053 ref NR_114251.1	98.829	427	0	761	<i>Lactobacillus senmaizukei</i>
2006	seq_402	gi 1199303358 ref NR_147709.1	98.829	427	0	761	<i>Lactobacillus wasatchensis</i>
21	seq_005	gi 959494902 ref NR_133769.1	100	427	0	789	<i>Leuconostoc gelidum</i>
2001	seq_401	gi 959494902 ref NR_133769.1	99.766	427	0	784	
1591	seq_319	gi 959494902 ref NR_133769.1	98.656	372	0	660	
61	seq_013	gi 1269801505 ref NR_074997.2	100	427	0	789	
101	seq_021	gi 1441204190 ref NR_157602.1	100	427	0	789	<i>Leuconostoc mesenteroides</i>
791	seq_159	gi 631250774 ref NR_109004.1	99.766	427	0	784	<i>Leuconostoc pseudomesenteroides</i>
166	seq_034	gi 343200124 ref NR_040811.1	100	427	0	789	<i>Leuconostoc carnosum</i>
1011	seq_203	gi 343202334 ref NR_042620.1	100	427	0	789	<i>Leuconostoc holzapfelii</i>
1786	seq_358	gi 631252057 ref NR_113255.1	100	427	0	789	<i>Leuconostoc lactis</i>
26	seq_006	gi 566085118 ref NR_108858.1	99.061	426	0	765	<i>Weissella fabalis</i>
146	seq_030	gi 566085118 ref NR_108858.1	98.826	426	0	760	
316	seq_064	gi 566085118 ref NR_108858.1	98.826	426	0	760	
321	seq_065	gi 566085118 ref NR_108858.1	98.826	426	0	760	
1886	seq_378	gi 566085118 ref NR_108858.1	99.061	426	0	765	
96	seq_020	gi 265678736 ref NR_029041.1	100	427	0	789	<i>Weissella koreensis</i>
236	seq_048	gi 265678736 ref NR_029041.1	99.766	427	0	784	
296	seq_060	gi 265678736 ref NR_029041.1	99.766	427	0	784	
832	seq_167	gi 310975060 ref NR_036924.1	100	427	0	789	<i>Weissella cibaria</i>
191	seq_039	gi 631252761 ref NR_113959.1	100	427	0	789	<i>Lactococcus raffinolactis</i>
516	seq_104	gi 631252761 ref NR_113959.1	99.766	427	0	784	
901	seq_181	gi 631252761 ref NR_113959.1	99.766	427	0	784	
273	seq_055	gi 631252762 ref NR_113960.1	100	427	0	789	<i>Lactococcus lactis</i>
358	seq_072	gi 631252070 ref NR_113268.1	100	427	0	789	<i>Lactococcus garvieae</i>
76	seq_016	gi 343204087 ref NR_043739.1	100	427	0	789	<i>Lactococcus piscium</i>
156	seq_032	gi 631252600 ref NR_113798.1	100	427	0	789	<i>Carnobacterium divergens</i>
696	seq_140	gi 343201367 ref NR_042093.1	99.532	427	0	780	<i>Carnobacterium gallinarum</i>
991	seq_199	gi 636560400 ref NR_116460.1	100	427	0	789	<i>Carnobacterium jeotgali</i>
126	seq_026	gi 631252389 ref NR_113587.1	100	427	0	789	<i>Brochothrix thermosphacta</i>
176	seq_036	gi 631252148 ref NR_113346.1	98.135	429	0	747	<i>Acinetobacter lwoffii</i>
291	seq_059	gi 631252148 ref NR_113346.1	97.902	429	0	741	
1346	seq_270	gi 631252148 ref NR_113346.1	100	428	0	791	
326	seq_066	gi 645320413 ref NR_117624.1	99.065	428	0	769	<i>Acinetobacter johnsonii</i>
456	seq_092	gi 645320413 ref NR_117624.1	99.533	428	0	780	
566	seq_114	gi 645320413 ref NR_117624.1	99.766	428	0	785	

586	seq_118	gi 645320413 ref NR_117624.1	99.766	428	0	785	
916	seq_184	gi 645320413 ref NR_117624.1	99.533	428	0	780	
921	seq_185	gi 645320413 ref NR_117624.1	99.065	428	0	769	
986	seq_198	gi 645320413 ref NR_117624.1	99.766	428	0	785	
1051	seq_211	gi 645320413 ref NR_117624.1	99.533	428	0	780	
1071	seq_215	gi 645320413 ref NR_117624.1	98.832	428	0	763	
751	seq_151	gi 1341394069 ref NR_153741.1	99.533	428	0	780	<i>Acinetobacter celticus</i>
1701	seq_341	gi 1341394069 ref NR_153741.1	98.832	428	0	763	
846	seq_170	gi 645320481 ref NR_117677.1	97.436	429	0	730	<i>Acinetobacter baumannii</i>
1741	seq_349	gi 645320481 ref NR_117677.1	97.669	429	0	736	
651	seq_131	gi 961554986 ref NR_133953.1	97.902	429	0	741	
1386	seq_278	gi 961554986 ref NR_133953.1	100	428	0	791	<i>Acinetobacter gandensis</i>
1566	seq_314	gi 961554986 ref NR_133953.1	98.14	429	0	747	
976	seq_196	gi 1137647852 ref NR_145641.1	99.533	428	0	780	<i>Acinetobacter albensis</i>
1276	seq_256	gi 1137647852 ref NR_145641.1	98.598	428	0	758	
806	seq_162	gi 1146059109 ref NR_145841.1	99.766	428	0	785	<i>Acinetobacter movanagherensis</i>
891	seq_179	gi 1146059109 ref NR_145841.1	98.364	428	0	752	
941	seq_189	gi 645320415 ref NR_117626.1	100	428	0	791	<i>Acinetobacter guillouiae</i>
1261	seq_253	gi 645320419 ref NR_117629.1	99.533	428	0	780	<i>Acinetobacter tjernbergiae</i>
1296	seq_260	gi 1230874683 ref NR_148844.1	97.196	428	0	725	<i>Acinetobacter dispersus</i>
1356	seq_272	gi 1397641879 ref NR_156989.1	98.832	428	0	763	<i>Acinetobacter defluvii</i>
691	seq_139	gi 1277396356 ref NR_152069.1	98.598	428	0	758	<i>Acinetobacter pragensis</i>

---

**Table S3.** The list of total samples of culturable bacterial communities.

Coded name	Species name	Closest accession number
<b>Intestine</b>		
C1	<i>Enterococcus faecium</i>	NR 115764
C2	<i>Enterococcus faecium</i>	NR 115764
C3	<i>Enterococcus durans</i>	NR 113900
C4	<i>Enterococcus durans</i>	NR 113900
C5	<i>Lactobacillus sakei</i>	NR 113821
haC1	<i>Enterococcus hirae</i>	NR 114783
haC3	<i>Enterococcus faecium</i>	NR 115764
haC8	<i>Bacillus toyonensis/wiedmannii/sanguinis</i>	NR 121761/NR 152692/NR 175555
<b>0 day</b>		
0/1	<i>Leuconostoc mesenteroides</i>	NR 074957
0/2	<i>Leuconostoc mesenteroides</i>	NR 074957
0/5	<i>Leuconostoc mesenteroides</i>	NR 074957
0/6	<i>Leuconostoc mesenteroides</i>	NR 074957
0/7	<i>Leuconostoc mesenteroides</i>	NR 074957
0/8	<i>Leuconostoc mesenteroides</i>	NR 074957
0/9	<i>Leuconostoc mesenteroides</i>	NR 074957
0/10	<i>Leuconostoc mesenteroides</i>	NR 074957
0/11	<i>Leuconostoc mesenteroides</i>	NR 074957
0/12	<i>Leuconostoc mesenteroides</i>	NR 074957
0/13	<i>Leuconostoc mesenteroides</i>	NR 074957
0/14	<i>Leuconostoc mesenteroides</i>	NR 074957
0/15	<i>Leuconostoc mesenteroides</i>	NR 074957
0/16	<i>Leuconostoc mesenteroides</i>	NR 074957
0/17	<i>Leuconostoc mesenteroides</i>	NR 074957
0/18	<i>Leuconostoc mesenteroides</i>	NR 074957
0/19	<i>Leuconostoc mesenteroides</i>	NR 074957
0/20	<i>Leuconostoc mesenteroides</i>	NR 074957
0/21	<i>Leuconostoc mesenteroides</i>	NR 074957
0/22	<i>Leuconostoc mesenteroides</i>	NR 074957
ha0/0	<i>Lactobacillus sakei</i>	NR 113821
ha0/1	<i>Bacillus tropicus/paramycoides/luti</i>	NR 157736/NR 157734/NR 157730
ha0/2	<i>Marichromatium purpuratum</i>	NR 116469
ha0/3	<i>Bacillus toyonensis/wiedmannii/sanguinis</i>	NR 121761/NR 152692/NR 175555
ha0/10	<i>Enterococcus faecium</i>	NR 115764
ha0/12	<i>Bacillus tropicus/paramycoides/luti</i>	NR 157736/NR 157734/NR 157730
ha0/13	<i>Macroccoccus caseolyticus</i>	NR 119262
ha0/20	<i>Corynebacterium lipophiloflavum</i>	NR 026370
<b>2<sup>nd</sup> day</b>		
2/1	<i>Leuconostoc mesenteroides</i>	NR 074957
2/2	<i>Leuconostoc mesenteroides</i>	NR 074957

2/3	<i>Leuconostoc mesenteroides</i>	NR 074957
2/5	<i>Leuconostoc mesenteroides</i>	NR 074957
2/7	<i>Leuconostoc mesenteroides</i>	NR 074957
2/8	<i>Leuconostoc mesenteroides</i>	NR 074957
2/9	<i>Leuconostoc mesenteroides</i>	NR 074957
2/10	<i>Leuconostoc mesenteroides</i>	NR 074957
2/11	<i>Leuconostoc mesenteroides</i>	NR 074957
2/12	<i>Leuconostoc mesenteroides</i>	NR 074957
2/13	<i>Leuconostoc mesenteroides</i>	NR 074957
2/15	<i>Leuconostoc mesenteroides</i>	NR 074957
2/16	<i>Leuconostoc mesenteroides</i>	NR 074957
2/17	<i>Leuconostoc mesenteroides</i>	NR 074957
2/18	<i>Leuconostoc mesenteroides</i>	NR 074957
2/19	<i>Leuconostoc mesenteroides</i>	NR 074957
2/20	<i>Leuconostoc mesenteroides</i>	NR 074957
ha2/1	<i>Lactobacillus sakei</i>	NR 113821
ha2/2	<i>Serratia proteamaculans</i>	NR 025341
ha2/4	<i>Hafnia alvei</i>	NR 044729
ha2/7	<i>Hafnia paralvei</i>	NR 116898
ha2/9	<i>Carnobacterium divergens</i>	NR 113798
ha2/10	<i>Serratia proteamaculans</i>	NR 025341
ha2/13	<i>Bacillus licheniformis</i>	NR 118996
ha2/17	<i>Macrococcus canis</i>	NR 156154
ha2/20	<i>Pectobacterium wasabiae</i>	NR 118294
<hr/>		
7 <sup>th</sup> day		
7/1	<i>Leuconostoc mesenteroides</i>	NR 074957
7/2	<i>Leuconostoc mesenteroides</i>	NR 074957
7/3	<i>Leuconostoc mesenteroides</i>	NR 074957
7/4	<i>Leuconostoc mesenteroides</i>	NR 074957
7/5	<i>Leuconostoc mesenteroides</i>	NR 074957
7/6	<i>Leuconostoc mesenteroides</i>	NR 074957
7/7	<i>Leuconostoc mesenteroides</i>	NR 074957
7/8	<i>Leuconostoc mesenteroides</i>	NR 074957
7/9	<i>Leuconostoc mesenteroides</i>	NR 074957
7/10	<i>Leuconostoc mesenteroides</i>	NR 074957
7/11	<i>Leuconostoc mesenteroides</i>	NR 074957
7/12	<i>Leuconostoc mesenteroides</i>	NR 074957
7/14	<i>Weissella cibaria</i>	NR 036924
7/15	<i>Leuconostoc mesenteroides</i>	NR 074957
7/16	<i>Leuconostoc mesenteroides</i>	NR 074957
7/17	<i>Leuconostoc mesenteroides</i>	NR 074957
7/18	<i>Leuconostoc mesenteroides</i>	NR 074957
7/19	<i>Leuconostoc mesenteroides</i>	NR 074957



7/20	<i>Leuconostoc mesenteroides</i>	NR 074957
ha7/0	<i>Lactobacillus sakei</i>	NR 113821
ha7/3	<i>Citrobacter murlinae</i>	NR 028688
ha7/7	<i>Pantoea agglomerans</i> / <i>Enterobacter ludwigii</i> / <i>Enterobacter cloacae</i> subsp. <i>dissolvens</i>	NR 111998/NR 042349/NR 118011
ha7/8	<i>Enterococcus pallens</i>	NR 043794
ha7/10	<i>Lactococcus lactis</i> subsp. <i>hordniae</i>	NR 113958
ha7/11	<i>Enterobacter aerogenes</i>	NR 113614
ha7/13	<i>Macrococcus canis</i>	NR 156154
ha7/15	<i>Kocuria kristinae</i>	NR 026199
ha7/20	<i>Pantoea agglomerans</i> / <i>Enterobacter ludwigii</i> / <i>Enterobacter cloacae</i> subsp. <i>dissolvens</i>	NR 111998/NR 042349/NR 118011

#### 10<sup>th</sup> day

10/1	<i>Leuconostoc mesenteroides</i>	NR 074957
10/4	<i>Leuconostoc mesenteroides</i>	NR 074957
10/6	<i>Leuconostoc mesenteroides</i>	NR 074957
10/7	<i>Leuconostoc mesenteroides</i>	NR 074957
10/8	<i>Leuconostoc mesenteroides</i>	NR 074957
10/9	<i>Leuconostoc mesenteroides</i>	NR 074957
10/10	<i>Leuconostoc mesenteroides</i>	NR 074957
10/11	<i>Leuconostoc mesenteroides</i>	NR 074957
10/12	<i>Leuconostoc mesenteroides</i>	NR 074957
10/13	<i>Leuconostoc mesenteroides</i>	NR 074957
10/14	<i>Leuconostoc mesenteroides</i>	NR 074957
10/15	<i>Leuconostoc mesenteroides</i>	NR 074957
10/16	<i>Leuconostoc mesenteroides</i>	NR 074957
10/17	<i>Leuconostoc mesenteroides</i>	NR 074957
10/18	<i>Leuconostoc mesenteroides</i>	NR 074957
ha10/1	<i>Enterococcus faecium</i>	NR 115764
ha10/3	<i>Lactobacillus sakei</i>	NR 113821
ha10/4	<i>Leuconostoc mesenteroides</i>	NR 074957
ha10/6	<i>Enterococcus faecium</i>	NR 115764
ha10/7	<i>Carnobacterium divergens</i>	NR 113798
ha10/8	<i>Enterococcus casseliflavus</i>	NR 104560
ha10/11	<i>Moraxella osloensis</i>	NR 113392
ha10/13	<i>Carnobacterium maltaromaticum</i>	NR 044710
ha10/15	<i>Lactobacillus sakei</i> subsp. <i>carnosus</i>	NR 104208

#### 14<sup>th</sup> day

14/3	<i>Leuconostoc mesenteroides</i>	NR 074957
14/4	<i>Leuconostoc pseudomesenteroides</i>	NR 040814
14/5	<i>Leuconostoc mesenteroides</i>	NR 074957
14/6	<i>Leuconostoc mesenteroides</i>	NR 074957
14/7	<i>Leuconostoc mesenteroides</i>	NR 074957
14/9	<i>Leuconostoc mesenteroides</i>	NR 074957
14/10	<i>Leuconostoc mesenteroides</i>	NR 074957

14/11	<i>Leuconostoc mesenteroides</i>	NR 074957
14/12	<i>Leuconostoc mesenteroides</i>	NR 074957
14/13	<i>Leuconostoc mesenteroides</i>	NR 074957
14/14	<i>Leuconostoc mesenteroides</i>	NR 074957
14/16	<i>Leuconostoc mesenteroides</i>	NR 074957
14/17	<i>Leuconostoc mesenteroides</i>	NR 074957
14/18	<i>Leuconostoc mesenteroides</i>	NR 074957
ha14/1	<i>Lactobacillus sakei</i>	NR 113821
ha14/2	<i>Lactobacillus sakei</i>	NR 113821
ha14/3	<i>Carnobacterium divergens</i>	NR 113798
ha14/4	<i>Shigella sonney</i>	NR 104826
ha14/7	<i>Enterococcus faecium</i>	NR 115764
ha14/10	<i>Lactococcus garvieae</i>	NR 113268

---

**21<sup>st</sup> day**

---

21/1	<i>Leuconostoc mesenteroides</i>	NR 074957
21/2	<i>Leuconostoc mesenteroides</i>	NR 074957
21/3	<i>Leuconostoc mesenteroides</i>	NR 074957
21/4	<i>Leuconostoc mesenteroides</i>	NR 074957
21/5	<i>Leuconostoc mesenteroides</i>	NR 074957
21/6	<i>Leuconostoc rapi</i>	NR 136799
21/7	<i>Leuconostoc mesenteroides</i>	NR 074957
21/8	<i>Leuconostoc mesenteroides</i>	NR 074957
21/10	<i>Leuconostoc mesenteroides</i>	NR 074957
21/11	<i>Leuconostoc mesenteroides</i>	NR 074957
21/13	<i>Leuconostoc mesenteroides</i>	NR 074957
21/14	<i>Lactococcus garvieae</i>	NR 113268
21/15	<i>Leuconostoc mesenteroides</i>	NR 074957
21/16	<i>Leuconostoc mesenteroides</i>	NR 074957
21/17	<i>Leuconostoc mesenteroides</i>	NR 074957
ha21/1	<i>Lactobacillus sakei</i>	NR 113821
ha21/4	<i>Bacillus tropicus/paramycoides/luti</i>	NR 157736/NR 157734/NR 157730
ha21/5	<i>Lactobacillus sakei</i>	NR 113821
ha21/6	<i>Lactobacillus sakei</i>	NR 113821
ha21/7 (1)	<i>Bacillus toyonensis/wiedmannii/sanguinis</i>	NR 121761/NR 152692/NR 175555
ha21/7 (2)	<i>Bacillus toyonensis/wiedmannii/sanguinis</i>	NR 121761/NR 152692/NR 175555
ha21/11	<i>Lactobacillus sakei</i>	NR 113821
ha21/14	<i>Lactobacillus sakei</i>	NR 113821

---

\*the coded names without prefixes represent isolates grown on MRS medium

**Table S4.** Physical-chemical and technological parameters as the mean differences of statistical significance throughout the ripening of “Pirrot ‘ironed’ sausage”.

Dependent Variable	(I) Ripening	(J) Ripening	Mean Difference (I-J)	Sig.	Dependent Variable	(I) Ripening	(J) Ripening	Mean Difference (I-J)	Sig.	Dependent Variable	(I) Ripening	(J) Ripening	Mean Difference (I-J)	Sig.
pH	0 day	7 <sup>th</sup> day	0.08111*	0.040	L*	0 day	7 <sup>th</sup> day	3.72333	0.451	L*	0 day	7 <sup>th</sup> day	-0.23667	0.999
		14 <sup>th</sup> day	0.10333*	0.009			14 <sup>th</sup> day	11.91333*	0.002			14 <sup>th</sup> day	0.89667	0.877
		21 <sup>st</sup> day	0.21333*	0.000			21 <sup>st</sup> day	10.58000*	0.004			21 <sup>st</sup> day	1.21667	0.714
		28 <sup>th</sup> day	0.49333*	0.000			28 <sup>th</sup> day	8.33000*	0.019			28 <sup>th</sup> day	3.02667	0.061
	7 <sup>th</sup> day	14 <sup>th</sup> day	0.02222	0.874		7 <sup>th</sup> day	14 <sup>th</sup> day	8.19000*	0.021		7 <sup>th</sup> day	14 <sup>th</sup> day	1.13333	0.761
		21 <sup>st</sup> day	0.13222*	0.002			21 <sup>st</sup> day	6.85667	0.056			21 <sup>st</sup> day	1.45333	0.575
		28 <sup>th</sup> day	0.41222*	0.000			28 <sup>th</sup> day	4.60667	0.269			28 <sup>th</sup> day	3.26333*	0.042
	14 <sup>th</sup> day	21 <sup>st</sup> day	0.11000*	0.006		14 <sup>th</sup> day	21 <sup>st</sup> day	-1.33333	0.968		14 <sup>th</sup> day	21 <sup>st</sup> day	0.32	0.997
		28 <sup>th</sup> day	0.39000*	0.000			28 <sup>th</sup> day	-3.58333	0.486			28 <sup>th</sup> day	2.13	0.247
	21 <sup>st</sup> day	28 <sup>th</sup> day	0.28000*	0.000		21 <sup>st</sup> day	28 <sup>th</sup> day	-2.25	0.825		21 <sup>st</sup> day	28 <sup>th</sup> day	1.81	0.381
a <sub>w</sub>	0 day	7 <sup>th</sup> day	0.01144	0.132	a*	0 day	7 <sup>th</sup> day	2.76667*	0.002	a*	0 day	7 <sup>th</sup> day	-0.28	0.996
		14 <sup>th</sup> day	0.04667*	0.000			14 <sup>th</sup> day	5.08000*	0.000			14 <sup>th</sup> day	6.10000*	0.000
		21 <sup>st</sup> day	0.08667*	0.000			21 <sup>st</sup> day	6.49667*	0.000			21 <sup>st</sup> day	9.15333*	0.000
		28 <sup>th</sup> day	0.11667*	0.000			28 <sup>th</sup> day	7.87000*	0.000			28 <sup>th</sup> day	12.51667*	0.000
	7 <sup>th</sup> day	14 <sup>th</sup> day	0.03522*	0.000		7 <sup>th</sup> day	14 <sup>th</sup> day	2.31333*	0.007		7 <sup>th</sup> day	14 <sup>th</sup> day	6.38000*	0.000
		21 <sup>st</sup> day	0.07522*	0.000			21 <sup>st</sup> day	3.73000*	0.000			21 <sup>st</sup> day	9.43333*	0.000
		28 <sup>th</sup> day	0.10522*	0.000			28 <sup>th</sup> day	5.10333*	0.000			28 <sup>th</sup> day	12.79667*	0.000
	14 <sup>th</sup> day	21 <sup>st</sup> day	0.04000*	0.000		14 <sup>th</sup> day	21 <sup>st</sup> day	1.41667	0.106		14 <sup>th</sup> day	21 <sup>st</sup> day	3.05333*	0.024
		28 <sup>th</sup> day	0.07000*	0.000			28 <sup>th</sup> day	2.79000*	0.002			28 <sup>th</sup> day	6.41667*	0.000
	21 <sup>st</sup> day	28 <sup>th</sup> day	0.03000*	0.000		21 <sup>st</sup> day	28 <sup>th</sup> day	1.37333	0.121		21 <sup>st</sup> day	28 <sup>th</sup> day	3.36333*	0.013
Fat (%)	0 day	7 <sup>th</sup> day	-0.27667	0.999	b*	0 day	7 <sup>th</sup> day	1.59	0.236	b*	0 day	7 <sup>th</sup> day	0.49667	0.960
		14 <sup>th</sup> day	-2.67	0.138			14 <sup>th</sup> day	2.54000*	0.031			14 <sup>th</sup> day	9.99333*	0.000
		21 <sup>st</sup> day	-2.72667	0.127			21 <sup>st</sup> day	5.08667*	0.000			21 <sup>st</sup> day	14.13333*	0.000

