

Supplementary data, Table S1, S2, and S3

Table S1. Primers used for mPCR to determine the presence of *Listeria* genus, and serogroup *L. monocytogenes* in the study (Doumith et al., 2004)

PCR assay	Target Gene	Product Size (bp)	Primer Sequences (5'-3')
mPCR1	<i>ORF2110</i>	597	<i>ORF2110-F: AGTGGACAATTGATTGGTGAA</i> <i>ORF2110-R: CATCCATCCCTTACTTTGGAC</i>
	<i>ORF2819</i>	471	<i>ORF2819-F: AGCAAAATGCCAAACTCGT</i> <i>ORF2819-R: CATCACTAAAGCCTCCCATTG</i>
	<i>Imo1118</i>	691	<i>Imo1118-F: AGGGGTCTTAAATCCTGGAA</i> <i>Imo1118-R: CGGCTTGTTGCGCATACTTA</i>
	<i>Imo0737</i>	906	<i>Imo0737-F: AGGGCTTCAAGGACTTACCC</i> <i>Imo0737-R: ACGATTTCTGCTTGCCATTC</i>
	<i>Prs</i>	370	<i>prs-F: GCTGAAGAGATTGCGAAAGAAG</i> <i>prs-R: CAAAGAAACCTTGGATTTGCGG</i>

Table S2. Primers used for mPCR speciation in this study (Ryu et al., 2013)

Species	Gene	Primer	Sequences (5'-3')	PCR Product Size (bp)
<i>Listeria</i> genus	<i>Prs</i>	<i>prs-F</i>	GCTGAAGAGATTGCGAAAGAAG	370
		<i>prs-R</i>	CAAAGAAACCTTGGATTTGCGG	
<i>L. grayi</i>	<i>Oxidoreductase</i>	<i>JOgrayi-F</i>	GCGGATAAAGGTGTTTCGGGTCAA	201
		<i>JOgrayi-R</i>	ATTTGCTATCGTCCGAGGCTAGG	
<i>L. innocua</i>	<i>lin0464</i>	<i>lin0464-F</i>	CGCATTTATCGCCAAAACCTC	749
		<i>lin0464-R</i>	TCGTGACATAGACGCGATTG	
<i>L. ivanovii</i>	<i>namA</i>	<i>liv22-228-F</i>	CGAATTCCTTATTCACTTGAGC	463
		<i>liv22-228-R</i>	GGTGCTGCGAACTTAACTCA	
<i>L. monocytogenes</i>	<i>lmo1030</i>	<i>lmo1030-F</i>	GCTTGTATTCACCTGGATTTGTCTGG	509
		<i>lmo1030-R</i>	ACCATCCGCATATCTCAGCCAACT	
<i>L. seeligeri</i>	<i>lmo033</i>	<i>lseelin-F</i>	GTACCTGCTGGGAGTACATA	673
		<i>lseelin-R</i>	CTGTCTCCATATCCGTACAG	
<i>L. welshimeri</i>	<i>scrA</i>	<i>lwe1801-F</i>	CGTGGCACAATAGCAATCTG	281
		<i>lwe1801-R</i>	GACATGCCTGCTGAACTAGA	

Table S3. Primer sequences, PCR preparation, and PCR condition used for virulence gene detection in this study

PCR assay	Target Gene	Product Size (bp)	Primer Sequences (5' 3')	PCR Preparation	PCR Condition	References
mPCR1	<i>inlB</i>	376	<i>inlB</i> -F: <i>GATATTGTGCCACTTTCAGGTT</i>	12.5 µL 2× <i>DreamTaq</i> master mix,	2 min at 94°C,	Liu et al. (2007)
			<i>inlB</i> -R: <i>CCTCTTTCAGTGGTTGGGTT</i>	5 µL nuclease-free water,	35 cycles of 94°C for 30 s,	
	<i>plcA</i>	1484	<i>plcA</i> -F: <i>CTGCTTGAGCGTTCATGTCTCATCCC</i>	5 µL template DNA,	55°C for 30 s,	
			<i>plcA</i> -R: <i>ATGGGTTTTCACTCTCCTTCTAC</i>	3 µL primer mix for mPCR2	72°C for 1 min and	
	<i>hlyA</i>	456	<i>hlyA</i> -F: <i>GTTAATGAACCTACAAGACCTTCC</i> <i>hlyA</i> -R: <i>ACCGTTCTCCACCATTCCCA</i>		a final extension at 72°C for 10 min.	
<i>actA</i>	839	<i>actA</i> -F: <i>TCGCCGCGGAAATTAATAAAGA</i> <i>actA</i> -R: <i>ACGAAGGAACCGGGCTGCTAG</i>				
mPCR2	<i>iap</i>	131	<i>iap</i> -F: <i>ACAAGCTGCACCTGTTGCAG</i> <i>iap</i> -R: <i>TGACAGCGTGTGTAGTAGCA</i>			Liu et al. (2007)
	<i>inlA</i>	800	<i>inlA</i> -F: <i>ACGAGTAACGGGACAAATGC</i> <i>inlA</i> -R: <i>CCCGACAGTGGTGCTAGATT</i>			
	<i>inlC</i>	517	<i>inlC</i> -F: <i>AATTCCCACAGGACACAACC</i> <i>inlC</i> -R: <i>CGGGAATGCAATTTTCACTA</i>			
	<i>inlJ</i>	238	<i>inlJ</i> -F: <i>TGTAACCCCGCTTACACACAGTT</i> <i>inlJ</i> -R: <i>AGCGGCTTGGCAGTCTAATA</i>			