

**NF VALIDATION**  
**Validation of alternative analytical methods**  
*Application in food microbiology*

## Summary report

Validation study according to the EN ISO 16140-2:2016

### **BAX® System PCR Assay for Genus *Listeria* 24E**

(certificate number QUA 18/06 – 07/08)

for detection of *Listeria* spp in food products and environmental samples (excluding samples from primary production)

#### Qualitative method

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This report consists of 69 pages, including 7 appendices.

Only copies including the totality of this report are authorised.

Competencies of the laboratory are certified by COFRAC accreditation for the analyses marked with the symbol♦.

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<b>1</b>	<b>INTRODUCTION</b>	<b>4</b>
<b>2</b>	<b>METHOD PROTOCOLS</b>	<b>5</b>
2.1	<b>Alternative method</b>	<b>5</b>
2.1.1	<i>Principle</i>	5
2.1.2	<i>Protocol</i>	5
2.1.3	<i>Restrictions</i>	6
2.2	<b>Reference methods</b>	<b>6</b>
2.3	<b>Study design</b>	<b>7</b>
<b>3</b>	<b>INITIAL VALIDATION AND EXTENSION STUDIES: RESULTS</b>	<b>7</b>
3.1	<b>Method comparison study</b>	<b>7</b>
3.1.1	<i>Sensitivity study</i>	7
3.1.2	<i>Relative level of detection</i>	17
3.1.3	<i>Inclusivity / exclusivity</i>	20
3.1.4	<i>Practicability</i>	21
3.1.5	<i>Method comparison study conclusion</i>	21
3.2	<b>Inter-laboratory study: organization and results</b>	<b>22</b>
3.2.1	<i>Study organization</i>	22
3.2.2	<i>Experimental parameters controls</i>	22
3.2.3	<i>Results analysis</i>	23
3.2.4	<i>Calculation and interpretation</i>	25
3.3	<b>General conclusion</b>	<b>28</b>
>	<i>Appendix 1 – Flow diagram of the alternative methods: BAX® System PCR Assay for Genus Listeria 24E and BAX® System X5 PCR Assay for Genus Listeria</i>	29
>	<i>Appendix 2 – Flow diagram of the reference method ISO 11290-1 (May 2017): Microbiology of the food chain - Horizontal method for the detection and enumeration of Listeria monocytogenes and other Listeria spp. - Part 1: detection method</i>	30
>	<i>Appendix 3 – Artificial contamination of samples</i>	31
>	<i>Appendix 4 – Sensitivity study: raw data</i>	36
>	<i>Appendix 5 - Relative level of detection: raw data</i>	54
>	<i>Appendix 6 – Inclusivity and exclusivity study: raw data</i>	61
>	<i>Appendix 7 - Results obtained by the collaborators (study realized by IPL, 2008)</i>	64

Quality Assurance documents related to this study can be consulted upon request from **Hygiena**.

The technical protocol and the result interpretation were realized according to the EN ISO 16140-2:2016 and the AFNOR technical rules (PR Revision 12).

<b>Validation protocols</b>	<ul style="list-style-type: none"> <li>▪ ISO 16140-1 (2016): Microbiology of the food chain - Method validation — <i>Part 1: Vocabulary</i></li> <li>▪ ISO 16140-2 (2016): Microbiology of the food chain - Method validation — <i>Part 2: Protocol for the validation of alternative (proprietary) methods against a reference method</i></li> <li><input checked="" type="checkbox"/> AFNOR technical rules (PR Revision 12)</li> </ul>
<b>Reference methods*</b>	<ul style="list-style-type: none"> <li>▪ NF EN ISO 11290-1/A1 (February 2005): Microbiology of food and animal feeding stuffs - Horizontal method for the detection and enumeration of <i>Listeria monocytogenes</i> - Part 1: detection method</li> <li>▪ ISO 11290-1 (May 2017): Microbiology of the food chain - Horizontal method for the detection and enumeration of <i>Listeria monocytogenes</i> and of <i>Listeria</i> spp.- Part 1: detection method</li> </ul>
<b>Alternative method</b>	<b>BAX® System PCR Assay for Genus <i>Listeria</i> 24E</b>
<b>Scope</b>	<input checked="" type="checkbox"/> Food products <input checked="" type="checkbox"/> Environmental samples ( <i>excluding samples from primary production</i> )
<b>Certification organism</b>	AFNOR Certification ( <a href="http://nf-validation.afnor.org/">http://nf-validation.afnor.org/</a> )

\* Analyses performed according to the COFRAC accreditation (Accreditation Testing n°1-0144, scope available on [www.cofrac.fr](http://www.cofrac.fr))

## 1 INTRODUCTION

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The **BAX® System PCR Assay for Genus *Listeria* 24E** was validated in July 2008 (certificate number QUA 18/06 – 07/08) according to the ISO 16140 (2003). Extension studies and renewal studies were performed:

<i>July 2008</i>	Initial validation for human food products, except smoked fishes, environmental samples (excluding samples from primary production) (IPL)
<i>January 2009</i>	Extension study: specific protocol for smoked fishes, as well as raw and cooked delicatessen (IPL)
<i>May 2011</i>	Extension to the 2.8 version of the BAX® software
<i>July 2012</i>	Renewal study (ADRIA)
<i>March 2016</i>	Renewal study (ADRIA)
<i>September 2016</i>	Extension study for using the BAX® System X5 instrument (Hygiena) with the BAX® System X5 PCR Assay for genus <i>Listeria</i> (Hygiena internal data)
<i>October 2017</i>	Extension study for a modification of the software as well as an update to be in agreement with the ISO 16140-2:2016 and the AFNOR technical rules (Revision 6). The current version of the software is version 3.6. The data generated during this study was interpreted with both versions (2.9 versus 3.6) (ADRIA).
<i>April 2020</i>	Renewal study (ADRIA)
<i>January 2024</i>	Extension study for the use of BAX® System software version 5.0 (Hygiena internal data)
<i>June 2024</i>	Renewal study (ADRIA)

## 2 METHOD PROTOCOLS

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### 2.1 Alternative method

#### 2.1.1 Principle

The BAX® System PCR Assay for Genus *Listeria* 24E and the BAX® System X5 PCR Assay for Genus *Listeria* are based on the gene amplification of a *Listeria* specific nucleic sequence by PCR technology.

The reagents necessary for the PCR reaction and for the internal control are included in one PCR tube.

The BAX® System Q7 and the BAX® System X5 PCR instruments are composed by a thermocycler and an optical module detecting the fluorescence. The software program analyses the level of fluorescence and provides results, *i.e.* positive or negative.

#### 2.1.2 Protocol

The flow diagrams are given in **Appendix 1**. The protocols are described below:

- Enrichment

- General protocol for food products (except smoked fish, raw and cooked delicatessen) and environmental samples: 26h ± 2h at 37°C ± 1°C in 24 LEB complete broth (ready to use) (BO1205S) or 24 LEB base (CM1107B) + selective supplement (SR0243E)
- Specific protocol for smoked fish, raw and cooked delicatessen: 26h ± 2h à 37°C ± 1°C in 24 LEB complete broth (ready to use) (BO1205S) + a non-selective supplement (BO1204M) or 24 LEB base (CM1107B) + selective supplement (SR0243E) + a non-selective supplement (BO1204M)

- DNA extraction step, amplification, detection

Step	BAX® System PCR Assay 24E	BAX® System X5 PCR Assay
Extraction	<ul style="list-style-type: none"> <li>○ Addition of 50 µl lysis reagents (40 µL agent 1 and 10 µL agent 2) in a cluster tube</li> <li>○ Addition of <b>0.5 ml</b> enriched sample in the cluster tube</li> <li>○ Incubation for 30 min at 37°C</li> <li>○ Addition of 200 µl lysis buffer + protease in a new cluster tube</li> <li>○ Transfer 5 µl of the heat-treated sample to the cluster tube</li> <li>○ Incubation for 30 min at 55°C and 10 min at 95°C</li> <li>○ Cooling for 5 min in a cooling block</li> </ul>	<ul style="list-style-type: none"> <li>○ Addition of 150 µl protease and 200 µl lysis Agent 2 to one 12 ml bottle of lysis buffer</li> <li>○ Transfer 200 µl prepared lysis reagent to each cluster tube</li> <li>○ Transfer <b>5 µl</b> enriched sample to the cluster tube</li> <li>○ Incubation for 30 min at 55°C and 10 min at 95°C</li> <li>○ Cooling for 5 min in a cooling block</li> </ul>
Amplification	<ul style="list-style-type: none"> <li>○ Transfer <b>50 µl</b> of the lysate in a PCR tube</li> </ul>	<ul style="list-style-type: none"> <li>○ Transfer <b>50 µl</b> of the lysate in a PCR tube</li> </ul>
Detection	<ul style="list-style-type: none"> <li>○ Run the PCR in the automate BAX® System Q7 instrument</li> </ul>	<ul style="list-style-type: none"> <li>○ Run the PCR in the automate BAX® System X5 instrument</li> </ul>

- Confirmation of positive result by:

- Using the conventional tests described by CEN or ISO, including purification step
- Streaking 10µl of 24 LEB enriched sample onto Brilliance Listeria Agar plates, incubated for 24-48 h at 37°C
- Streaking 0.1 ml onto agar Listeria plate according to Ottaviani and Agosti (O&A), incubated for 24-48 h at 37°C.

The presence of characteristic colonies is sufficient to confirm the presence of *Listeria* species.

### 2.1.3 Restrictions

*Listeria grayi* is excluded from the scope of the method.

## 2.2 Reference methods♦

The initial and extension studies were run using the EN ISO 11290-1/A1 (2005): Microbiology of food and animal feeding stuffs - Horizontal method for the detection and enumeration of *Listeria monocytogenes* - Part 1: detection method.

♦ Analyses performed according to the COFRAC accreditation (Accreditation Testing n°1-0144, scope available on [www.cofrac.fr](http://www.cofrac.fr))

The extension study (2017) was run using the ISO 11290-1 (May 2017): Microbiology of the food chain - Horizontal method for the detection and enumeration of *Listeria monocytogenes* and of *Listeria* spp. - Part 1: detection method.

*The modifications which occur in the version published in 2017 are considered as minor and have no impact on the previous data.*

For the renewal study, the reference method is the ISO 11290-1 (May 2017): Microbiology of the food chain - Horizontal method for the detection and enumeration of *Listeria monocytogenes* and of *Listeria* spp. - Part 1: detection method.

The flow diagrams are given in **Appendix 2**.

## 2.3 Study design

The study is **an unpaired study design** as the reference and the alternative methods have different enrichment procedures.

# 3 INITIAL VALIDATION AND EXTENSION STUDIES: RESULTS

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## 3.1 Method comparison study

*The method comparison study is a study performed by the expert laboratory to compare the alternative method with the reference method.*

*The study was carried out on a diversity of samples and strains representative of agri-food products. This does not constitute an exhaustive list of the different matrices included in the scope.*

*For any comment on the alternative method, please contact AFNOR Certification at <http://nf-validation.afnor.org/contact-2/>.*

### 3.1.1 Sensitivity study

*The sensitivity (SE) is the ability of the method to detect the analyte by either the reference or alternative method.*

### 3.1.1.1 Number and nature of samples

427 samples were analyzed in 2008 and 2009 by IPL. During the initial validation study, raw and cooked delicatessens were analyzed using the general protocol. These samples were removed from the study as a specific protocol is now dedicated to these products.

In agreement with the AFNOR Technical Committee, 24 samples were removed from the extension study interpretation due to an inoculation level higher than 10 CFU/sample.

In 2017, 93 additional samples were tested by ADRIA as part of the extension study.

Combining the different studies (initial and extension), 248 positive samples and 248 negative samples were obtained for a total of 496 samples. The repartition of samples per category and type is summarized in Table 1.

All the data provided in the report were obtained using the **BAX® System PCR Assay for Genus *Listeria* 24E**.

**Table 1 - Repartition per category and type**

Category		Type		Protocol	Positive samples	Negative samples	Total	
1	Composite foods	a	RTE	G	11	9	20	
		b	RTRH	G	11	11	22	
		c	Pastries, desserts, egg products	G	9	11	20	
		Total			31	31	62	
2	Meat products	a	Raw meat products (raw, frozen, seasoned)	G	14	8	22	
		b	RTRH, RTC	G	10	11	21	
		c	Delicatessen (raw and cooked)	S	28	28	56	
		Total			52	47	99	
3	Dairy products	a	Raw milk cheeses	G	10	11	21	
		b	Raw milk	G	10	10	20	
		c	Heat treated dairy products	G	11	21	32	
		Total			31	42	73	
4	Fruits and vegetables	a	Raw, non-processed (fresh or frozen)	G	7	13	20	
		b	Fresh, bagged	G	18	7	25	
		c	RTE, RTRH	G	16	15	31	
		Total			41	35	76	
5	Seafood and fishery products	a	Raw	G	16	22	38	
		b	Smoked	S	20	15	35	
		c	RTE, RTRH	G	17	10	27	
		Total			53	47	100	
6	Environmental samples	a	Process water	G	11	18	29	
		b	Surfaces	G	9	11	20	
		c	Dusts and residues	G	20	17	37	
		Total			40	46	86	
All categories				248	248	496		
Total General protocol (G)				200	205	405		
Total Specific protocol (S)				48	43	91		

The distribution of positive samples per target analyte is presented in Table 2.

**Table 2 - Distribution per target analyte**

Category	Listeria spp		Mix <i>Listeria</i> spp. + <i>Listeria monocytogenes</i>		Listeria spp mixed or alone		<i>Listeria monocytogenes</i>		Total positive samples
	Number of samples	%	Number of samples	%	Number of samples	%	Number of samples	%	
1	12	38,7%	3	9,7%	15	48,4%	16	51,6%	31
2	16	30,8%	13	25,0%	29	55,8%	23	44,2%	52
3	5	16,1%	10	32,3%	15	48,4%	16	51,6%	31
4	12	29,3%	4	9,8%	16	39,0%	25	61,0%	41
5	7	13,2%	8	15,1%	15	28,3%	38	71,7%	53
6	18	45,0%	5	12,5%	23	57,5%	17	42,5%	40
<b>Total</b>	<b>70</b>	<b>28,2%</b>	<b>43</b>	<b>17,3%</b>	<b>113</b>	<b>45,6%</b>	<b>135</b>	<b>54,4%</b>	<b>248</b>

### 3.1.1.2 Artificial contamination of samples

The strains were stressed using various injury protocols. For the spiking protocol, the injury efficiency was evaluated by comparing enumeration onto the selective medium (Palcam plates) and on a non-selective medium (TSYE plates).

The artificial contaminations are provided in **Appendix 3**.

71 samples were artificially contaminated. 59 gave a positive result. 189 samples were naturally contaminated.

The repartition of the positive samples per contamination level is given in Table 3.

**Table 3 - Repartition of the positive samples per contamination level**

Naturally contaminated	Artificially contaminated						Total	
	Spiking protocol			Seeding protocol				
	$\leq 5$	$5 < x \leq 10$	$10 < x \leq 30$	$\leq 3$	$3 < x \leq 10$	$10 < x \leq 30$		
Number of samples	189	14	17	13	15	0	248	
%	76,2%	5,6%	6,9%	5,2%	6,0%	0,0%	100,0%	

6.9% of the samples were inoculated between 3 or 5 CFU and 10 CFU, this is in agreement with the AFNOR technical rules.

**Taking into account all the studies, 76.2 % of the samples were naturally contaminated.**

### 3.1.1.3 Protocols applied during the validation study

> **Incubation time**

The minimum incubation time was applied:

- Enrichment step in 24 LEB: 24 h at 37°C,
- O&A plates: 24h to 48 h.

> **Confirmations**

For the initial study and extension study performed by IPL, 10 µl of the 24 LEB broth was streaked onto *Brilliance Listeria* Agar.

For the extension study performed in 2017, 0.1 ml of the 24 LEB broth was streaked onto O&A and onto Palcam and RAPID'L. *mono* plates if required.

The typical colonies were confirmed by biochemical gallery without purification step and by applying the tests described in the reference method after purification step.

### 3.1.1.4 Test results

Raw data per category are given in **Appendix 4**. The results are given in the Table 4.

**Table 4 – Summary of results obtained with the reference  
and the alternative methods**

Category		PA	NA*	PD	ND**	PPND	PPNA
1	Composite	18	30	5	7	1	1
2	Meat products	39	46	6	7	0	1
3	Dairy products	23	41	5	3	0	1
4	Fruits and vegetables	37	33	3	1	0	2
5	Seafood	41	44	9	3	0	3
6	Environmental samples	29	44	5	6	0	2
<b>All categories</b>		<b>187</b>	<b>238</b>	<b>33</b>	<b>27</b>	<b>1</b>	<b>10</b>
<b>General protocol</b>		<b>154</b>	<b>197</b>	<b>23</b>	<b>22</b>	<b>1</b>	<b>8</b>
<b>Specific protocol</b>		<b>33</b>	<b>41</b>	<b>10</b>	<b>5</b>	<b>0</b>	<b>2</b>

PA = positive agreement (R+/A+)

PD = positive deviation (R-/A+)

PP = positive presumptive non confirmed samples

\*: PPNA not included

NA = negative agreement (A-/R-)

ND = negative deviation (A-/R+)

\*\*: PPND not included

### 3.1.1.5 Calculation of the relative trueness (RT), the sensitivity (SE) and the false positive ratio (FPR)

The calculations are presented in Table 5.

**Table 5 – Calculation of the relative trueness (RT), the sensitivity (SE) and the false positive ratio (FPR)**

Category		Type	PA	NA*	PD	ND**	PPND	PPNA	SE <sub>alt</sub> %	SE <sub>ref</sub> %	RT %	FPR %	
1	Composite	a RTE	5	8	2	3	1	1	63,6	81,8	70,0	25	
		b RTRH	5	11	3	3	0	0	72,7	72,7	72,7	0,0	
		c Pastries, desserts, egg products	8	11	0	1	0	0	88,9	100,0	95,0	0,0	
		Total	18	30	5	7	1	1	74,2	83,9	79,0	6,5	
2	Meat products	a Raw meat products (raw, frozen, seasoned)	11	7	1	2	0	1	85,7	92,9	86,4	14,3	
		b RTRH, RTC	10	11	0	0	0	0	100,0	100,0	100,0	0,0	
		c Delicatessen (raw and cooked)	18	28	5	5	0	0	82,1	82,1	82,1	0,0	
		Total	39	46	6	7	0	1	86,5	88,5	86,9	2,1	
3	Dairy products	a Raw milk cheeses	6	11	2	2	0	0	80,0	80,0	81,0	0,0	
		b Raw milk	9	10	0	1	0	0	90,0	100,0	95,0	0,0	
		c Heat treated dairy products	8	20	3	0	0	1	100,0	72,7	90,6	5,0	
		Total	23	41	5	3	0	1	90,3	83,9	89,0	2,4	
4	Fruits and vegetables	a Raw, non-processed (fresh or frozen)	7	12	0	0	0	1	100,0	100,0	100,0	8,3	
		b Fresh, bagged	17	7	1	0	0	0	100,0	94,4	96,0	0,0	
		c RTE, RTRH	13	14	2	1	0	1	93,8	87,5	90,3	7,1	
		Total	37	33	3	1	0	2	97,6	92,7	94,7	5,7	
5	Seafood	a Raw	14	21	1	1	0	1	93,8	93,8	94,7	4,8	
		b Smoked	15	13	5	0	0	2	100,0	75,0	85,7	15,4	
		c RTE, RTRH	12	10	3	2	0	0	88,2	82,4	81,5	0,0	
		Total	41	44	9	3	0	3	94,3	83,0	88,0	6,4	
6	Environmental samples	a Process water	8	17	2	1	0	1	90,9	81,8	89,7	5,9	
		b Surfaces	8	11	0	1	0	0	88,9	100,0	95,0	0,0	
		c Dusts and residues	13	16	3	4	0	1	80,0	85,0	81,1	6,3	
		Total	29	44	5	6	0	2	85,0	87,5	87,2	4,3	
All categories			187	238	33	27	1	10	88,7	86,7	87,7	4,4	
General protocol (G)			154	197	23	22	1	8	88,5	88,5	88,6	4,4	
Specific protocol (S)			33	41	10	5	0	2	89,6	79,2	83,5	4,7	

\* PPNA not included

\*\* PPND not included

The results obtained for all the categories are the following (See **Table 6**).

**Table 6 - Summary of results**

		All categories	General protocol	Specific protocol
Sensitivity for the alternative method	$SE_{alt} = \frac{(PA + PD)}{(PA + ND + PD)} \times 100\%$	88.7 %	88.5 %	89.6 %
Sensitivity for the reference method	$SE_{ref} = \frac{(PA + ND +)}{(PA + ND + PD)} \times 100\%$	86.7 %	88.5 %	79.2 %
Relative trueness	$RT = \frac{(PA + NA)}{N} \times 100\%$	87.7 %	88.6 %	83.5 %
False positive ratio for the alternative method*	$FPR = \frac{(FP)}{NA} \times 100\%$ FP = PPNA + PPND	4.4 %	4.4 %	4.7 %

\* With    ND = ND + PPND  
           NA = NA + PPNA

### 3.1.1.6 Analysis of discordant results

28 negative deviations were observed, 23 with naturally contaminated samples and 5 with artificially contaminated samples. For 7 samples, the confirmatory tests concluded the presence of *Listeria* spp. in the enrichment broth, all were naturally contaminated.

Among the 33 samples in positive deviations, 9 were artificially contaminated and 24 naturally contaminated.

The negative deviations are given in Table 7 and the positive deviations in Table 8.





The analyses of discordant results according to the EN ISO 16140-2:2016 is the following (See **Table 9**).

**Table 9 - Analyses of discordant results**

Category	Type	PD	ND	PPND	N+	(ND+PPND)-PD	AL
1 Composite foods / Ready-to-eat and ready-to-reheat	a	2	3	1		2	
	b	3	3	0		0	
	c	0	1	0		1	
	Total	5	7	1	31	3	3
2 Meat Products	a	1	2	0		1	
	b	0	0	0		0	
	c	5	5	0		0	
	Total	6	7	0	52	1	3
3 Milks & Dairy products	a	2	2	0		0	
	b	0	1	0		1	
	c	3	0	0		-3	
	Total	5	3	0	31	-2	3
4 Vegetables	a	0	0	0		0	
	b	1	0	0		-1	
	c	2	1	0		-1	
	Total	3	1	0	41	-2	3
5 Seafood and fishery products	a	1	1	0		0	
	b	5	0	0		-5	
	c	3	2	0		-1	
	Total	9	3	0	53	-6	3
6 Environmental Samples	a	2	1	0		-1	
	b	0	1	0		1	
	c	3	4	0		1	
	Total	5	6	0	40	1	3
<b>All categories</b>		33	27	1	248	-5	8
<b>General protocol</b>		23	22	1	200	0	7
<b>Specific protocol</b>		10	5	0	48	-5	4

**The calculated values for ((ND + PPND) - PD) meet the Acceptability Limit (AL) for each category and for all the categories.**

### 3.1.1.7 Confirmations

Confirmations were carried out by streaking the 24 LEB broth (10 µl for the 2008 and 2009 studies, and 0.1 ml for this extension study) onto O&A plates. Characteristic colonies were observed for all samples after a 24 h incubation time of the O&A plates except for 12 samples.

For samples n° AA5, D19 and A9: characteristic colonies were observed only after 48 h incubation.

For samples n° U10, U12, C9, A11, A17, D17, E1 and 4865: characteristic colonies were observed after a subculture in Fraser 1.

For sample n° 4867: characteristic colonies were observed on the Palcam plate only.

### 3.1.1.8 PCR inhibition

No PCR inhibition was observed during the different studies.

### 3.1.2 Relative level of detection

*The relative level of detection is the level of detection at P = 0.50 (LOD<sub>50</sub>) of the alternative (proprietary) method divided by the level of detection at P = 0.50 (LOD<sub>50</sub>) of the reference method.*

The RLOD is defined as the ratio of the alternative and reference methods:

$$RLOD = \frac{LOD_{Alt.}}{LOD_{Ref.}}$$

#### 3.1.2.1 Experimental design

During studies carried out in 2008 and 2009, seven (matrix/strain) pairs were analyzed by the reference and the alternative methods.

Contaminations and enumerations were done according to the AFNOR technical rules (protocol for low level inoculation). The contamination levels were:

- Level 1: 0 CFU/g or ml,
- Level 2: level necessary to obtain 0 to 50 % positive,
- Level 3: level necessary to obtain 50 to 75 % positive,
- Level 4: level necessary to obtain 100 % positive.

For the extension study (2017), one matrix / strain pair was tested for the composite food category: a deli salad (Piémontaise) inoculated with *L. seeligeri* Ad1293 (isolated from parsley), using the protocol described in the EN ISO 16140-2:2016:

- 5 negative samples
- 20 samples inoculated at a level providing fractional positive results

- 5 samples inoculated at a higher level.

The matrix was stored for 48 h at 5°C ± 3°C after inoculation and before analysis.

The matrices tested and the strains inoculated are given in Table 10.

**Table 10 - Defined (matrix/strain) pairs for the RLOD determination**

Study	Category	Matrix	Inoculated strain	Origin	Protocol
Initial 2008	Dairy products	Raw milk	<i>L. ivanovii</i> L50	Dairy product	General
	Vegetables	Raw vegetable mix	<i>L. monocytogenes</i> 4b L58	Salad	General
	Environmental samples	Process water	<i>L. innocua</i> L175	Process water	General
Extension	Meat products	Rillettes	<i>L. monocytogenes</i> L49	Chipolatas	Specific
	Fishery products (smoked fish)	Smoked salmon	<i>L. monocytogenes</i> 1/2 a L5	Smoked salmon	Specific
Extension 2017	Composite foods	Deli Salad (Piemontaise)	<i>L. seeligeri</i> Ad1293	Parsley	General

### 3.1.2.2 Calculation and interpretation of the RLOD

The raw data are given in **Appendix 5**.

The RLOD calculations were performed using the Excel spreadsheet available at <http://standards.iso.org/iso/16140> - RLOD (clause 5-1-4-2 Calculation and interpretation of RLOD) version 15.08.2015. The RLOD are given in Table 11.

**Table 11 – Presentation of RLOD before and after confirmation of the alternative method results**

Name		RLOD	RLODL	RLODU	b=ln(RLOD)	sd(b)	z-Test statistic	p-value
Raw milk / <i>L.ivanovii</i> L150	0,888	0,392	2,015	-0,118	0,409	0,289	1,227	
	0,838	0,294	2,389	-0,177	0,524	0,338	1,264	
	0,859	0,348	2,116	-0,152	0,451	0,338	1,265	
Extension 2009	Rillettes / <i>L.monocytogenes</i> L49	1,000	0,406	2,462	0,000	0,450	0,000	1,000
	Raw fish fillet / <i>L.monocytogenes</i> L5	1,710	0,463	6,315	0,536	0,653	0,821	0,412
Extension 2017	Deli salad / <i>L. seeligeri</i> Ad1293	2,175	0,968	4,890	0,777	0,405	1,918	0,055
<b>Combined</b>		<b>1,175</b>	<b>0,818</b>	<b>1,688</b>	<b>0,162</b>	<b>0,181</b>	<b>0,893</b>	<b>0,372</b>

**The RLOD meet the Acceptability Limit fixed at 2.5 for an unpaired study design for each matrix/strain pair and for all matrices combined.**

The LOD<sub>50</sub> % calculations according to Wilrich & Wilrich POD-LOD calculation program - version 11, 2022-10-12 test are given in Table 7.

**Table 12 - LOD<sub>50</sub> results**

Category	(Strain / matrix) pair	Level of detection at 50% (CFU / test portion) according to Wilrich & Wilrich) <sup>1</sup>	
		Reference method	Alternative method
Initial 2008	3	Raw milk / <i>L.ivanovii</i> L 150	0.9 [0.5; 1.5]
	4	Raw vegetables mix / <i>L.monocytogenes</i> L58	0.8 [0.4; 1.4]
	6	Process water / <i>L.innocua</i> L175	0.9 [0.5; 1.6]
Extension 2009	2	Rillettes / <i>L.monocytogenes</i> L49	0.6 [0.4; 1.1]
	5	Raw fish fillet / <i>L.monocytogenes</i> L 5	0.5 [0.3; 0.8]
Extension 2017	1	Deli salad / <i>L. seeligeri</i> Ad1293	0.4 [0.2; 0.7]
<b>Combined</b>		<b>0.6 [0.5; 0.8]</b>	<b>0.6 [0.5; 0.8]</b>

**The LOD<sub>50</sub> varies from 0.4 to 0.9 CFU/sample size for the reference method and from 0.3 to 0.8 CFU/ sample size for the alternative method.**

<sup>1</sup> Wilrich, C., and P.-Th. Wilrich: Estimation of the POD function and the LOD of a qualitative microbiological measurement method. AOAC International **92** (2009) 1763 - 1772.

### 3.1.3 Inclusivity / exclusivity

The inclusivity is the ability of the alternative method to detect the target analyte from a wide range of strains. The exclusivity is the lack of interference from a relevant range of non-target strains of the alternative method.

#### 3.1.3.1 Test protocols

The inclusivity and the exclusivity of the method are defined by analysis, respectively of 50 positive strains and 30 negative strains.

##### > Inclusivity

*Listeria* strains were tested with the complete BAX® System PCR Assay for Genus *Listeria* 24E protocol.

##### > Exclusivity

The different negative strains were inoculated in a nutrient broth at a level around  $10^5$  cells/mL. After incubation for 24 h at 37°C, the BAX® System PCR Assay for Genus *Listeria* 24E test was performed for each strain.

#### 3.1.3.2 Results

The raw data (in French) are given in **Appendix 6**.

67 strains of *Listeria* (36 *Listeria monocytogenes* and 31 *Listeria* spp strains (species different from *monocytogenes*) were tested:

- all the *Listeria monocytogenes* strains were detected with the BAX® Genus *Listeria* 24E method,
- all the *Listeria* different from *Listeria monocytogenes* gave a positive result with the BAX® System PCR Assay for Genus *Listeria* 24E method, except *Listeria seeligeri* L148 for which an inoculation level of  $10^5$  CFU was required to obtain a positive PCR result. 5 other strains of *Listeria seeligeri* were tested and gave positive results with a low inoculation level.

No cross reaction was observed with the 31 non-*Listeria* strains.

### 3.1.4 Practicability

The alternative method practicability was evaluated according to the AFNOR criteria relative to method comparison study.

<b>Storage, modalities of use after first use</b>	The storage temperature is between 2 to 8°C. Expiration date is shown on the kit package on the different reagent vials. - The reconstituted agent 1 should be stored at room temperature for 3 months maximum - the mixture agent 1 + agent 2 shall be used within 4 hours - lysis reagent + protease, once reconstituted, should be stored 15 days at 2-8°C		
<b>Time-to-result</b>	Step	<b>Alternative method</b>	<b>Reference method</b>
	Enrichment step	Day 0	Day 0
	Transfer to selective broths (Fraser)	/	Day 1
	BAX® procedure	Day 1	/
	Test result	<b>Day 1</b>	/
	<b>Negative result (if test is negative)</b>		
	Streaking of selective broths on selective media	Day 1	Day 1 and Day 3
	Reading the plates Confirmation tests	Day 2 to Day 3	Day 2 to Day 5
	<b>Negative result</b> (after streaking and negative confirmation if done, for example with a positive BAX® test)	<b>Day 2 to Day 3</b>	<b>Day 3 to Day 6</b>
<b>Common steps with the reference method</b>	<b>Positive result</b> Confirmation by the reference method tests (GRAM, catalase test) Confirmation on <i>Brilliance Listeria</i> Agar	Day 3	Day 3 to Day 6
		<b>Day 2 to Day 3</b>	

### 3.1.5 Method comparison study conclusion

The method comparison study conclusions are:

- The **BAX® System PCR Assay for Genus *Listeria* 24E** shows satisfying sensitivity results.
- The RLOD meet the Acceptability Limit for all the tested matrix/strain pairs.
- The alternative method is specific and selective.

## 3.2 Inter-laboratory study: organization and results

### 3.2.1 Study organization

Samples were sent to 13 laboratories. A pasteurized milk was inoculated with a *Listeria monocytogenes* strain (origin « raw milk cheese »).

2 sets of 24 samples were prepared per laboratory, one for the BAX® System PCR Assay for Genus *Listeria* 24E method and the other for the reference method. Each set of samples was divided in 3 levels of contamination, with 8 samples per level.

### 3.2.2 Experimental parameters controls

#### 3.2.2.1 Contamination levels

The contamination levels and the sample codification were the following (see Table 13).

**Table 13 - Contamination levels**

Level	Samples	Theoretical target level (b/25 g)	True level (b/25 g sample)	Low limit / 25 g sample	High limit / 25 g sample
Level 0 (L0)	3-4-9-10-11-12-19-20	0	0		
Low level (L1)	1-2-5-6-13-14-21-22	3	4.8	1.2	11.0
High level (L2)	7-8-15-16-17-18-23-24	30	48	36	63

#### 3.2.2.2 Logistic conditions

Temperature conditions are given in Table 14.

**Table 14 - Sample temperatures at receipt**

Laboratory	Reception Temperatures (°C)		Comments
	Communicated by the laboratory	Recorded by the probe	
A	3.5	1.9	/
B	4.0	5.1	/
C	2.0	2.0	/
D	7.2	6.8	/
E	0.4	0.9	/
F	10.0	5.2	/
G	7.3	6.6	/
H	17.0	15.9	Delivery at D2
I	15.5	14.5	Delivery at D2
J	Not communicated	13.6	Delivery at D2
K	Not communicated	8.2	/
L	5.6	4.7	/
M	7.2	6.5	/

Among the 13 laboratories, 3 received their samples at D2 at a temperature above 8.4°C. Labs I and J didn't carry out the analyses.

Due to the delivery conditions, the results of 10 laboratories were used for interpretation.

### 3.2.3 Results analysis

#### 3.2.3.1 Expert laboratory results

The results obtained by the expert laboratory are given in Table 15.

**Table 15 – Results obtained by the expert Lab.**

Level	Reference method	Alternative method
L0	0/8	0/8
L1	8/8	8/8
L2	8/8	8/8

### 3.2.3.2 Results of the collaborative laboratories retained for interpretation

The raw data are given in **Appendix 6**.

> **Aerobic mesophilic flora enumeration**

Depending on the Lab results, the enumeration levels varied from < 1 to 54 000 CFU/ml.

> **Listeria spp. detection**

The results obtained are provided in Table 16 (reference method) and Table 17 (alternative method).

**Table 16 - Positive results by the reference method**

Collaborator	Contamination level		
	L0	L1	L2
A	0	8	8
B	0	8	8
C	0	8	8
D	0	8	8
E	0	8	8
F	0	8	8
G	0	8	8
K	0	8	8
L	0	8	8
M	0	8	8
<b>TOTAL</b>	<b>P<sub>0</sub> = 0</b>	<b>P<sub>1</sub> = 80</b>	<b>P<sub>2</sub> = 80</b>

**Table 17 - Positive results (before and after confirmation)  
by the alternative method**

Collaborators	Contamination level					
	L0		L1		L2	
	Before confirmation	After confirmation	Before confirmation	After confirmation	Before confirmation	After confirmation
A	0	0	8	8	8	8
B	0	0	8	8	8	8
C	0	0	8	8	8	8
D	0	0	8	8	8	8
E	0	0	8	8	8	8
F	0	0	8	8	8	8
G	0	0	8	8	8	8
K	0	0	7	7	8	8
L	0	0	8	8	8	8
M	0	0	8	8	8	8
<b>TOTAL</b>	<b>P<sub>0</sub> = 0</b>	<b>CP<sub>0</sub> = 0</b>	<b>P<sub>1</sub> = 79</b>	<b>CP<sub>1</sub> = 79</b>	<b>P<sub>2</sub> = 80</b>	<b>CP<sub>2</sub> = 80</b>

### 3.2.4 Calculation and interpretation

#### 3.2.4.1 Calculation of the specificity percentage (SP)

The percentage specificities (SP) of the reference method and of the alternative method, using the data after confirmation, based on the results of level L0 are the following (See Table 18).

**Table 18 - Percentage specificity**

Specificity for the reference method	$SP_{ref} = \left(1 - \left(\frac{P_0}{N_-}\right)\right) \times 100 \% =$	100.0 %
Specificity for the alternative method	$SP_{alt} = \left(1 - \left(\frac{CP_0}{N_-}\right)\right) \times 100 \% =$	100.0 %

N: number of all L0 tests

P<sub>0</sub> = total number of false-positive results obtained with the blank samples before confirmation

CP<sub>0</sub> = total number of false-positive results obtained with the blank samples

**3.2.4.2 Calculation of the sensitivity ( $SE_{alt}$ ), the sensitivity for the reference method ( $SE_{ref}$ ), the relative trueness (RT) and the false positive ratio for the alternative method (FPR)**

Fractional positive results were not obtained for this study. The calculations were done with the results obtained for Level 1. A summary of the results obtained by the collaborators with the reference and the alternative methods is provided in Table 19.

**Table 19 - Summary of the obtained results with the reference method and the alternative method for Level 1**

Response	Reference method positive (R+)	Reference method negative (R-)
Alternative method positive (A+)	Positive agreement (A+/R+) <b>PA = 79</b>	Positive deviation (R-/A+) <b>PD = 0</b>
Alternative method negative (A-)	Negative deviation (A-/R+)	Negative agreement (A-/R-) <b>NA = 0</b>

Based on the data summarized in Table 19, the values of sensitivity of the alternative and reference methods, as well as the relative trueness and false positive ratio for the alternative method taking account the confirmations, are the following (See Table 20).

**Table 20 - Sensitivity, relative trueness and false positive ratio percentages**

Sensitivity for the alternative method:	$SE_{alt} = \frac{(PA+PD)}{(PA+PD+ND)} \times 100\% =$	98.8 %
Sensitivity for the reference method:	$SE_{ref} = \frac{(PA+ND)}{(PA+PD+ND)} \times 100\% =$	100.0 %
Relative trueness	$RT = \frac{(PA+NA)}{N} \times 100\% =$	98.8 %
False positive ratio for the alternative method	$FPR = \frac{FP}{NA} \times 100\% =$	0

**3.2.4.3 Interpretation of data**

For an **unpaired study design**, the difference between (ND – PD) is calculated for the level(s) where fractional recovery is obtained (so  $L_1$  and possibly  $L_2$ ). The observed value found for (ND – PD) shall not be higher than the AL. The AL is defined as  $[(ND - PD)_{max}]$  and calculated per level where fractional recovery is obtained as described below using the following three parameters:

$$(p+)_{ref} = \frac{P_x}{N_x}$$

where

$P_x$  = number of samples with a positive result obtained with the reference method at level  $x$  ( $L_1$  or  $L_2$ ) for all the collaborators

$N_x$  = number of samples tested at level  $x$  ( $L_1$  or  $L_2$ ) with the reference method by all the collaborators

$$(p+)_{\text{alt}} = \frac{CP_x}{N_x}$$

where

$CP_x$  = number of samples with a confirmed positive result obtained with the alternative method at level  $x$  ( $L_1$  or  $L_2$ ) for all the collaborators;

$N_x$  = number of samples tested at level  $x$  ( $L_1$  or  $L_2$ ) with the alternative method by all the collaborators.

$$(ND - PD)_{\text{max}} = \sqrt{3N_x \times ((p+)_{\text{ref}} + (p+)_{\text{alt}} - 2((p+)_{\text{ref}} \times (p+)_{\text{alt}}))}$$

where

$N_x$  = number of samples tested for level  $x$  ( $L_1$  or  $L_2$ ) with the reference method by all the collaborators.

The AL is not met when the observed value is higher than the AL. When the AL is not met, investigations should be made (e.g. root cause analysis) in order to provide an explanation of the observed results. Based on the AL and the additional information, it is decided whether the alternative method is regarded as not fit for purpose. The reasons for acceptance of the alternative method when the AL is not met shall be stated in the study report.

In this study, fractional recovery was observed at Level 1. The calculations are the following, according to the EN ISO 16140-2:2016 (See Table 21).

**Table 21 - Calculations**

$N_x$	80
$(p+)_{\text{ref}}$	1.0
$(p+)_{\text{alt}}$	1.0
$AL = (ND - PD) \text{ max}$	4.77
$ND - PD$	1
Conclusion	$ND - PD < AL$

**The ISO 16140-2 (2016) requirements are fulfilled as  $(ND - PD)$  meet the AL.**

### 3.2.4.4 Evaluation of the LOD<sub>50%</sub>, LOD<sub>95%</sub> and RLOD between laboratories

The LOD<sub>50%</sub>, the LOD<sub>95%</sub> and the RLOD was calculated using the EN ISO 16140-2:2016 Excel spreadsheet available at [https://standards.iso.org/iso/16140/-5/ed-1/en/RLOD\\_inter-lab-study\\_16140-2\\_AnnexF\\_ver1\\_28-06-2017.xls](https://standards.iso.org/iso/16140/-5/ed-1/en/RLOD_inter-lab-study_16140-2_AnnexF_ver1_28-06-2017.xls). The results are used only for information (see Table 22).

**Table 22 - LOD<sub>50%</sub>, LOD<sub>95%</sub> and RLOD**

Method	LOD 50%	LOD 95%	RLOD
Reference	/	/	/
Alternative	0.76 [0.48;1.19]	3.28 [2.08;5.16]	

The calculation for LOD<sub>50%</sub> (reference method), LOD<sub>95%</sub> (reference method) and RLOD is not possible as all the samples gave positive results using the reference method for both inoculation levels (L1 and L2).

### 3.2.4.5 Inter-laboratory study conclusion

The data and interpretations comply with the EN ISO 16140-2:2016 requirements. The alternative method is considered equivalent to the ISO standard.

## 3.3 General conclusion

The **method comparison study conclusions** are:

- The **BAX® System PCR Assay for Genus *Listeria* 24E** shows satisfying sensitivity results.
- The RLOD meet the Acceptability Limit for all the tested matrix/strain pairs.
- The alternative method is specific and selective.

The **inter-laboratory study conclusions** are:

- The data and interpretations comply with the EN ISO 16140-2:2016 requirements. **The BAX® System PCR Assay for Genus *Listeria* 24E is considered equivalent to the ISO standard.**

Quimper, 10 July 2024

Astrid CARIOU

Manager

Validation of Alternative methods

I hereby attest to the validation of the verification of the conformity of the report (opinion and interpretation).

**Appendix 1 – Flow diagram of the alternative methods: BAX® System PCR Assay for Genus *Listeria* 24E and BAX® System X5 PCR Assay for Genus *Listeria***

- 25 g + 225 ml of 24 LEB compete broth (ready to use broth) (BO1205S)  
or 24 LEB base (CM1107B) + selective supplement (SR0243E)

(General protocol)

OR

- 25 g + 225 ml of 24 LEB complete broth (ready to use) (BO1205S)  
+ a non-selective supplement (BO1204M)

or 24 LEB base (CM1107B) + selective supplement (SR0243E)  
+ a non-selective supplement (BO1204M)

(Specific protocol for smoked fish, raw and cooked delicatessen)

↓

Incubation for 26 h ± 2 h at 37°C

↓

BAX System 24E	BAX System X5
<p>40 µl lysis Agent 1 + 10 µl lysis Agent 2 ↓</p> <p>50 µl combined agents in a Cluster tube + 0.5 ml enriched sample ↓</p> <p>30 min at 37°C ↓</p> <p>200 µl reagent (150 µl protease + 12 ml buffer lysis) in a cluster tube + 5 µl heat-treated enriched sample ↓</p> <p>55°C for 30 min 95°C for 10 min Cool at 2 - 8°C for 5 min ↓</p> <p>PCR on <b>50 µl</b> lysate ↓</p> <p>Automatic measurement of the fluorescence using the BAX® Q7</p>	<p>150 µl protease + 200 µl Agent 2 + 12 ml lysis buffer ↓</p> <p>200 µl reagent in a Cluster tube + 5 µl enriched sample ↓</p> <p>55°C for 30 min 95°C for 10 min Cool at 2 - 8°C for 5 min ↓</p> <p>PCR on <b>50 µl</b> lysate ↓</p> <p>Automatic measurement of the fluorescence using the BAX® X5</p>

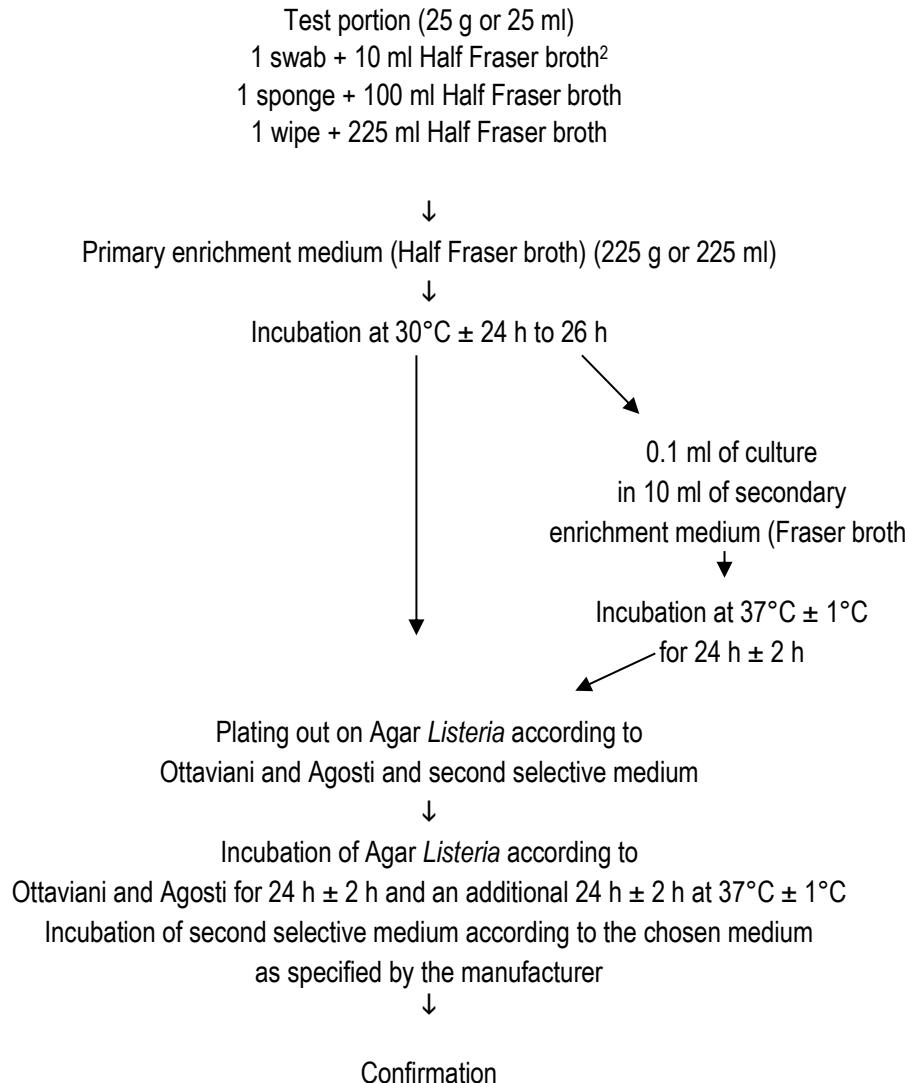
↓

Confirmation of positive result by:

- Using the conventional tests described by CEN or ISO, including purification step
- Streaking 10µl of 24 enriched sample onto Brilliance Listeria Agar plates, incubated for 24-48 h at 37°C
- Streaking 0.1 ml onto agar Listeria plate according to Ottaviani and Agosti (O&A), incubated for 24-48 h at 37°C.

The presence of characteristic colonies is sufficient to confirm the presence of *Listeria* species.

**Appendix 2 – Flow diagram of the reference method**  
**ISO 11290-1 (May 2017): Microbiology of the food chain -**  
**Horizontal method for the detection and enumeration of *Listeria monocytogenes* and other *Listeria* spp. - Part 1: detection method**



Target	Gram	Catalase	Beta hemolysis	CAMP test	Carbohydrates
<i>Listeria</i> spp	x	x			

<sup>2</sup> For sampling after cleaning process pre-moisten  
 - 1 swab + 1 ml broth universal neutralizing (+ 9 ml Half-Fraser)  
 - 1 sponge + 10 ml broth universal neutralizing (+ 90 ml Half-Fraser)  
 - 1 wipe + BPW + 10 % neutralizing agent (+ 225 ml Half-Fraser)

### Appendix 3 – Artificial contamination of samples

Laboratory	Sample N°	Products	Cat. (Initial classification)	Artificial contamination (IPL Study)					Global result
				Strain	Origin	Injury protocol	Injury measure- ment	Inoculation level/sample	
IPL	L8	Cordon bleu with vegetables	MP3	<i>Listeria monocytogenes</i> 4b	Salad	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,2	1,2	+
IPL	L18	Beef balls with tomato sauce	MP3	<i>Listeria monocytogenes</i> 1/2c	Beef	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,6	16	+
IPL	L19	Scallop cream	MP3	<i>Listeria monocytogenes</i> 1/2c	Beef	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,6	14	+
IPL	L12	Pollock fillet	SF1	<i>Listeria monocytogenes</i> 1/2a	Smoked salmon	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,3	4,5	+
IPL	L13	Fish with bordelaise sauce	SF3	<i>Listeria monocytogenes</i> 1/2a	Smoked salmon	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,3	4	+
IPL	L14	Frozen breaded fish	SF3	<i>Listeria monocytogenes</i> 1/2a	Smoked salmon	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,3	3,5	+
IPL	L15	Frozen breaded fish	SF3	<i>Listeria monocytogenes</i> 1/2a	Fish shell	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,8	2,8	+
IPL	L17	Salmon with hollandaise sauce	SF3	<i>Listeria monocytogenes</i> 1/2a	Fish shell	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,8	5,6	+
IPL	L20	Salmon dish	SF3	<i>Listeria monocytogenes</i> 1/2a	Fish shell	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,8	8,4	+
IPL	H5	Brie cheese	DP1	<i>Listeria monocytogenes</i> 1/2b	Maroille (raw milk cheese)	45 min at 55°C, 30 minutes at -80°C	0,9	9	+
IPL	H6	Raw milk cheese	DP1	<i>Listeria monocytogenes</i> 1/2b	Maroille (raw milk cheese)	45 min at 55°C, 30 minutes at -80°C	0,9	16,2	+
IPL	H8	Brie cheese	DP1	<i>Listeria monocytogenes</i> 1/2b	Maroille (raw milk cheese)	45 min at 55°C, 30 minutes at -80°C	0,9	12,6	+
IPL	J21	Camembert pasteurized milk cheese	DP1	<i>Listeria monocytogenes</i> 4e	Reblochon	45 min at 55°C, 30 minutes at -80°C	0,6	10	+

Laboratory	Sample N°	Products	Cat. (Initial classification)	Artificial contamination (IPL Study)					Global result
				Strain	Origin	Injury protocol	Injury measure- ment	Inoculation level/sample	
IPL	J22	Camembert de Normandie raw milk cheese	DP1	<i>Listeria monocytogenes</i> 4e	Reblochon	45 min at 55°C, 30 minutes at -80°C	0,6	9	+
IPL	J23	Reblochon raw milk cheese	DP1	<i>Listeria monocytogenes</i> 4e	Reblochon	45 min at 55°C, 30 minutes at -80°C	0,6	8	+
IPL	N15	Soft cheese	DP1	<i>Listeria monocytogenes</i> 4b	Munster (rind)	40% NaCl, 48 h at +4°C	0,5	3,2	+
IPL	N16	Camembert cheese	DP1	<i>Listeria monocytogenes</i> 4b	Munster (rind)	40% NaCl, 48 h at +4°C	0,5	6,4	+
IPL	N17	Light cheese	DP1	<i>Listeria monocytogenes</i> 4b	Munster (rind)	40% NaCl, 48 h at +4°C	0,5	9,6	+
IPL	N18	Gouda cheese	DP1	<i>Listeria monocytogenes</i> 4b	Munster (rind)	40% NaCl, 48 h at +4°C	0,5	12,8	+
IPL	N19	Brie de Meaux raw milk cheese	DP1	<i>Listeria monocytogenes</i> 1/2c	Munster (rind)	40% NaCl, 48 h at +4°C	0,6	0,5	-
IPL	J24	Blue cheese	DP2	<i>Listeria monocytogenes</i> 4e	Reblochon	45 min at 55°C, 30 min at -80°C	0,6	10	+
IPL	F10	Skim milk powder	DP3	<i>Listeria monocytogenes</i> 1/2a	Munster (rind)	60 min at 55°C, 30 min at -80°C	>2,5	1,5	-
IPL	F11	Skim milk powder	DP3	<i>Listeria monocytogenes</i> 1/2a	Munster (rind)	60 min at 55°C, 30 min at -80°C	>2,5	2,5	-
IPL	N3	Raw milk	DP3	<i>Listeria monocytogenes</i>	Coulommiers (raw milk cheese)	40% NaCl, 48 h at +4°C	0,5	9	+
IPL	N20	Cream	DP3	<i>Listeria monocytogenes</i> 1/2c	Munster (rind)	40% NaCl, 48 h at +4°C	0,6	0,8	+
IPL	P1	Chocolate pastry Versaillais	DP3	<i>Listeria monocytogenes</i> 1/2a	Munster (rind)	40% NaCl, 48 h at +4°C	0,4	14	+
IPL	P2	Princesse des îles pastry	DP3	<i>Listeria monocytogenes</i> 1/2a	Munster (rind)	40% NaCl, 48 h at +4°C	0,4	21	+
IPL	P16	Milk powder	DP3	<i>Listeria monocytogenes</i> 1/2b	Maroille (raw milk cheese)	20% NaCl, 24 h at -80°C	0,7	4	+

Laboratory	Sample N°	Products	Cat. (Initial classification)	Artificial contamination (IPL Study)					Global result
				Strain	Origin	Injury protocol	Injury measure- ment	Inoculation level/sample	
IPL	P23	Whipped cream puff	DP3	<i>Listeria monocytogenes</i> 1/2b	Maroille (raw milk cheese)	20% NaCl, 24 h at -80°C	0,7	6	+
IPL	R12	Whipped cream puff	DP3	<i>Listeria innocua</i>	Munster	45 min at 55°C, 30 min at -80°C	ND	0,7	-
IPL	R16	Raw milk	DP3	<i>Listeria monocytogenes</i>	Munster	45 min at 55°C, 30 min at -80°C	0,2	1,5	-
IPL	U1	Chocolate profiteroles	DP3	<i>Listeria monocytogenes</i>	Mozzarella	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	1,4	3	+
IPL	U2	Chocolate mousse	DP3	<i>Listeria monocytogenes</i>	Mozzarella	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	1,4	6	+
IPL	U3	Saint Honoré pastry	DP3	<i>Listeria monocytogenes</i>	Mozzarella	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	1,4	9	+
IPL	L9	Broccoli	VG2	<i>Listeria monocytogenes</i> 4b	Salad	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,2	1,2	+
IPL	L11	Grated carrots	VG2	<i>Listeria monocytogenes</i> 4b	Salad	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,2	2,4	+
IPL	L23	Carrots and Celery	VG2	<i>Listeria monocytogenes</i> 1/2a	Sauté potatoes	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,4	16,1	+
IPL	Y3	Broccoli	VG2	<i>Listeria monocytogenes</i> 1/2	Potatoes	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,7	2,8	+
IPL	Z11	Mixed vegetables	VG2	<i>Listeria monocytogenes</i>	Fried potatoes	45 min at 55°C, 30 min at -80°C	0,4	15	+
IPL	L10	Rice salad	VG3	<i>Listeria monocytogenes</i> 4b	Salad	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,2	1,8	+
IPL	P19	Cucumber salad	VG3	<i>Listeria monocytogenes</i> 1/2a	Sauté potatoes	20% NaCl, 24 h at -80°C	1,4	1,2	-
IPL	P20	Grated carrots salad	VG3	<i>Listeria monocytogenes</i> 1/2a	Sauté potatoes	20% NaCl, 24 h at -80°C	1,4	2,4	-
IPL	P22	Rice salad	VG3	<i>Listeria monocytogenes</i> 1/2a	Sauté potatoes	20% NaCl, 24 h at -80°C	1,4	4,8	+
IPL	P25	Celery with mayonnaise	VG3	<i>Listeria monocytogenes</i> 1/2a	Sauté potatoes	20% NaCl, 24 h at -80°C	1,4	3,6	-

Laboratory	Sample N°	Products	Cat. (Initial classification)	Artificial contamination (IPL Study)					Global result
				Strain	Origin	Injury protocol	Injury measure- ment	Inoculation level/sample	
IPL	Y1	Zucchinis flan	VG3	<i>Listeria monocytogenes</i> 1/2	Potatoes	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,7	2	-
IPL	Y2	Spinaches with cream	VG3	<i>Listeria monocytogenes</i> 1/2	Potatoes	48 h at 4°C, 30 min at 55°C, 30 min at -80°C	0,7	2,4	-
IPL	Z9	Mashed celery	VG3	<i>Listeria monocytogenes</i>	Fried potatoes	45 min at 55°C, 30 min at -80°C	0,4	10	+
IPL	Z10	Lenses salad	VG3	<i>Listeria monocytogenes</i>	Fried potatoes	45 min at 55°C, 30 min at -80°C	0,4	12,5	+
IPL	K9	Water of wash lines pea	EN1	<i>Listeria monocytogenes</i>	Environmental sample	45 min at 55°C, 30 min at -80°C	2,4	8,6	+
IPL	X1	Process water fish retail	EN1	<i>Listeria monocytogenes</i> 1/2a	Smoked salmon	45 min at 55°C, 30 min at -80°C	0,8	8,1	+
IPL	X2	Water process	EN1	<i>Listeria monocytogenes</i>	Perch fillet	45 min at 55°C, 30 min at -80°C	0,6	9	+
IPL	X5	Water process	EN1	<i>Listeria monocytogenes</i>	Spinach	45 min at 55°C, 30 min at -80°C	0,4	11	+
IPL	X3	Sponge	EN2	<i>Listeria monocytogenes</i> 1/2a	Smoked salmon	45 min at 55°C, 30 min at -80°C	0,8	16,2	+
IPL	X4	Scraps from fish retail outlet	EN3	<i>Listeria monocytogenes</i>	Perch fillet	45 min at 55°C, 30 min at -80°C	0,6	18	+

Laboratory	Sample N°	Products	Artificial contamination (ADRIA Study)					Global result
			Strain	Origin	Injury protocol	Injury measurement	Inoculation level/sample	
ADRIA	5150	Deli-salad	<i>L. innocua</i> Ad1230	Salad	Seeding 48h 2-8°C	/	1-0-1-1-0 (0,6)	-
ADRIA	5152	Deli-salad	<i>L. innocua</i> Ad1230	Salad	Seeding 48h 2-8°C	/	1-0-1-1-0 (0,6)	+
ADRIA	5154	Deli-salad	<i>L. innocua</i> Ad1230	Salad	Seeding 48h 2-8°C	/	1-0-1-1-0 (0,6)	+
ADRIA	5690	RTRH food	<i>L. welshimeri</i> Ad1175	RTRH food	Seeding 48h 2-8°C	/	1-0-1-1-0 (0,6)	+
ADRIA	5692	RTRH food	<i>L. welshimeri</i> Ad1175	RTRH food	Seeding 48h 2-8°C	/	1-0-1-1-0 (0,6)	+
ADRIA	5694	RTRH food	<i>L. welshimeri</i> Ad1175	RTRH food	Seeding 48h 2-8°C	/	1-0-1-1-0 (0,6)	+
ADRIA	5698	Pizza	<i>L. welshimeri</i> Ad1175	RTRH food	Seeding 48h 2-8°C	/	1-0-1-1-0 (0,6)	+
ADRIA	5926	Sandwich with ham	<i>L. welshimeri</i> Ad1204	Pork meat	Seeding 48h 2-8°C	/	1-2-3-0-0 (1,2)	+
ADRIA	5927	Deli-salad with ham	<i>L. welshimeri</i> Ad1204	Pork meat	Seeding 48h 2-8°C	/	1-2-3-0-0 (1,2)	+
ADRIA	5928	Deli-salad with chicken	<i>L. innocua</i> Ad1227	Poultry meat	Seeding 48h 2-8°C	/	2-1-1-0-1 (1,0)	+
ADRIA	5929	Deli-salad with chicken	<i>L. innocua</i> Ad1227	Poultry meat	Seeding 48h 2-8°C	/	2-1-1-0-1 (1,0)	+
ADRIA	5930	Sandwich with ham	<i>L. welshimeri</i> Ad1670	Pork meat	Seeding 48h 2-8°C	/	1-0-0-2-0 (0,6)	+
ADRIA	5931	Deli-salad with ham	<i>L. welshimeri</i> Ad1670	Pork meat	Seeding 48h 2-8°C	/	1-0-0-2-0 (0,6)	+
ADRIA	5932	Deli-salad with chicken	<i>L. ivanovii</i> Ad2465	Poultry meat	Seeding 48h 2-8°C	/	0-1-4-1-2 (1,6)	+
ADRIA	5933	Deli-salad with chicken	<i>L. ivanovii</i> Ad2465	Poultry meat	Seeding 48h 2-8°C	/	0-1-4-1-2 (1,6)	-
ADRIA	5950	Surface sample	<i>L. monocytogenes</i> Ad1271	Environmental sample	Seeding 48h 2-8°C	/	2-1-1-0-2 (1,2)	+
ADRIA	6278	Mushrooms	<i>L. innocua</i> Ad1176	Vegetables	Seeding 48h 2-8°C	/	4-2-3-0-1 (2,0)	+

## Appendix 4 – Sensitivity study: raw data (IPL and ADRIA Studies)

### IPL Legend

#### Total bacteria growth

Ø : no growth

L = low

M = medium

H = high

#### Distribution of flora

A = pure culture of suspicious colonies

B = mix with a majority of suspicious colonies

C = mix with a minority of suspicious colonies

D = mix with rare suspicious colonies

E = absence of suspicious colonies

(x) : x characteristic colonies of *Listeria* if  $x \leq 5$

- (L)(M)(H) (A to D) : presence of blue colonies without halo

ND: negative deviation

PD: positive deviation

PPNA: positive presumptive negative agreement

### ADRIA Legend

H: characteristic *Listeria* colonies without halo

H+: characteristic *Listeria* colonies with halo

-: no typical colonies but presence of background microflora

st: plate without any colony

i: PCR inhibition

PA: positive agreement

NA: negative agreement

ND: negative deviation

PD: positive deviation

PPNA: positive presumptive negative agreement

PPND : positive presumptive negative deviation

NC: non-characteristic colony on TSYEA

d: doubtful colony

\*: new extraction

NI: No identification

ni : No isolated colony

Analyses performed according to the COFRAC accreditation (Accreditation Testing n°1-0144, scope available on [www.cofrac.fr](http://www.cofrac.fr)) (ADRIA, Expert laboratory)

COMPOSITE FOODS																				
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1						Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E						Agreement	Protocol	Category	Type	
				FRASER 1/2		FRASER		CONFIRMATION		RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB			IDENTIFICATION	FINAL RESULT				
				O&A1	P1	O&A2	P2	IDENTIFICATION				Brilliance 24h (IPL)	Brilliance 48h (IPL)	O&A 24h (Adria)	O&A 48h (Adria)					
IPL	K4	Sandwich thon	Tuna sandwich	+LA	+LA	+MA	+MA	<i>L. monocytogenes</i>	+	+	+LB	+MB	<i>L. monocytogenes</i>	+	PA	G	1	a		
ADRIA	4866	Allumette poulet fumé	RTE chicken meat	H-d	+ (4)	H-	+	<i>L. welshimeri</i>	+	-/-	H+ (2)	H+ (2)	<i>L. monocytogenes</i>	-	ND	G	1	a		
ADRIA	5150	Salade jambon emmental	Deli-salad	-	-	st	st	/	-	-	-	-	/	-	NA	G	1	a		
ADRIA	5152	Salade jambon emmental	Deli-salad	H-	+	H-	+	<i>L. innocua</i>	+	-	-	-	/	-	ND	G	1	a		
ADRIA	5154	Salade poulet emmental	Deli-salad	-	-	st	st	/	-	+	H-	H-	<i>L. innocua</i>	+	PD	G	1	a		
ADRIA	5388	Sandwich poulet crudités	Sandwich	-	st	st	st	/	-	+/-	-	- (- after subculture in Fraser)	/	-	PPNA	G	1	a		
ADRIA	5390	Tartinable fromage saumon	RTE food	-	-	-	-	/	-	-	-	-	/	-	NA	G	1	a		
ADRIA	5392	Salade pomme de terre hareng	RTE food	st	st	st	st	/	-	-	-	-	/	-	NA	G	1	a		
ADRIA	5926	Sandwich jambon beurre	Sandwich with ham	st	st	st	st	/	-	+	H-	H-	<i>L. welshimeri</i>	+	PD	G	1	a		
ADRIA	5927	Piémontaise au jambon	Deli-salad with ham	H-	+	H-	+	<i>L. welshimeri</i>	+	+	H-	H-	<i>L. welshimeri</i>	+	PA	G	1	a		
ADRIA	5928	Salade césar	Deli-salad with chicken	H-	+	H-	+	<i>L. innocua</i>	+	+	H-	H-	<i>L. innocua</i>	+	PA	G	1	a		
ADRIA	5929	Salade au poulet	Deli-salad with chicken	H-	+	H-	+	<i>L. innocua</i>	+	+	H-	H-	<i>L. innocua</i>	+	PA	G	1	a		
ADRIA	5930	Sandwich jambon beurre	Sandwich with ham	H-	+	H-	+	<i>L. welshimeri</i>	+	+	H-	H-	<i>L. welshimeri</i>	+	PA	G	1	a		
ADRIA	5931	Piémontaise au jambon	Deli-salad with ham	H+	+	H+	+	<i>L. monocytogenes</i>	+	+/-	st	- (- after subculture in Fraser)	/	-	PPND	G	1	a		
ADRIA	5932	Salade césar	Deli-salad with chicken	-	-	H+	+	<i>L. ivanovii</i>	+	-	-	-	/	-	ND	G	1	a		
ADRIA	5933	Salade au poulet	Deli-salad with chicken	-	-	-	st	/	-	-	-	-	/	-	NA	G	1	a		
ADRIA	5934	Club sandwich rosette	Sandwich with ham	-	-	-	-	/	-	-	-	-	/	-	NA	G	1	a		
ADRIA	5935	Piémontaise au jambon	Deli-salad with ham	st	-	-	-	/	-	-	st	st	/	-	NA	G	1	a		
ADRIA	5936	Piémontaise au jambon	Deli-salad with ham	st	st	st	st	/	-	-	st	st	/	-	NA	G	1	a		
ADRIA	5937	Salade au poulet	Deli-salad with chicken	-	-	-	st	/	-	-	-	-	/	-	NA	G	1	a		
IPL	P11	Pie	Ø	Ø	Ø	-LE	/	-	-	-LE	/	/	-	NA	G	1	b			
ADRIA	4864	Poêlée de pommes de terre	RTRH potatoes	-	-	-	-	/	-	-	-	-	/	-	NA	G	1	b		
ADRIA	4865	Galette soja tomates	RTRH food	H-d	-	-	-	NC	-	+	-	- (H+ after subculture in Fraser 1)	<i>L. monocytogenes</i> after subculture in Fraser 1	+	PD	G	1	b		
ADRIA	4867	Poêlée de riz au poulet	RTRH chicken meat and rice	H-d	-	H+/H-	+	<i>L. monocytogenes</i> / <i>L. innocua</i>	+	+	H-d	H-d (+on Palcam)	<i>L. monocytogenes</i>	+	PA	G	1	b		

COMPOSITE FOODS																				
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1						Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E						Agreement	Protocol	Category	Type	
				FRASER 1/2		FRASER		CONFIRMATION		RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB			FINAL RESULT					
				O&A1	P1	O&A2	P2	IDENTIFICATION				Brilliance 24h (IPL)	Brilliance 48h (IPL)	<i>IDENTIFICATION</i>						
ADRIA	4869	Sauce colombo	RTRH dressing	-	-	-	-	/	-	-	-	-	-	/	-	NA	G	1	b	
ADRIA	4870	Galette de blé noir épaisse	RTRH food	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+/H-d	H+/H-d	<i>L. monocytogenes</i> / NC	+	PA	G	1	b		
ADRIA	4871	Galette quinoa provençale	RTRH food	H-d	-	-	-	NC, NI	-	-	H-d	H-d	NI	-	NA	G	1	b		
ADRIA	4875	Blanquette de poulet	RTRH food	st	-	-	-	/	-	-	-	-	-	-	NA	G	1	b		
ADRIA	4876	Paella	Paella	H+ (2)	-	H+	+	<i>L. monocytogenes</i>	+	+	H-d	H-d	<i>L. innocua</i>	+	PA	G	1	b		
ADRIA	4877	Galette boulgour lentilles	RTRH food	H-d	-	-	-	NC	-	-	-	-	/	-	NA	G	1	b		
ADRIA	4878	Tajine olives	RTRH food	-	-	-	-	/	-	-	-	-	/	-	NA	G	1	b		
ADRIA	4958	Brandade de morue parmentier	RTRH food	-	-	-	-	/	-	-	-	-	/	-	NA	G	1	b		
ADRIA	5384	Riz au thon olives tomates	RTRH food	H+	+	H+	+	<i>L. monocytogenes</i>	+	-	-	-	/	-	ND	G	1	b		
ADRIA	5385	Pommes de terre salardaise	RTRH food	-	-	-	-	/	-	-	-	-	/	-	NA	G	1	b		
ADRIA	5386	Pommes de terre au beurre	RTRH food	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+	H+	<i>L. monocytogenes</i>	+	PA	G	1	b		
ADRIA	5387	Bouchée à la reine	RTRH food	st	st	st	st	/	-	-	-	-	/	-	NA	G	1	b		
ADRIA	5389	Friand au fromage	RTRH food	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+	H+	<i>L. monocytogenes</i>	+	PA	G	1	b		
ADRIA	5391	Friand au fromage		st	-	-	-	/	-	-	-	-	/	-	NA	G	1	b		
ADRIA	5690	Croissant au jambon	RTRH food	-	-	-	-	/	-	+	H-	H-	<i>L. welshimeri</i>	+	PD	G	1	b		
ADRIA	5692	Croque monsieur fromage jambon	RTRH food	H-	+	H-	+	<i>L. welshimeri</i>	+	-	-	-	/	-	ND	G	1	b		
ADRIA	5694	Galette blé noir jambon emmental	RTRH food	-	-	-	-	/	-	+	H-	H-	<i>L. welshimeri</i>	+	PD	G	1	b		
ADRIA	5698	Pizza jambon fromages	Pizza	H-	+	H-	+	<i>L. innocua</i>	+	-	-	-	/	-	ND	G	1	b		
IPL	J8		Cookie dough	-ME	-ME	+MB	+MB	<i>L. monocytogenes</i>	+	-	-ME	/	/	-	ND	G	1	c		
IPL	O1		Chocolate pastry Opéra	+MB	+MD	+MB	+MB	<i>L. monocytogenes</i>	+	+	-LB	+LB	<i>L. monocytogenes</i>	+	PA	G	1	c		
IPL	P1		Chocolate pastry Versaillaïs	+LB	+LB	+MA	+MA	<i>L. monocytogenes</i>	+	+	+MB	+MB	<i>L. monocytogenes</i>	+	PA	G	1	c		
IPL	P2		Princesse des îles pastry	+LB	+LB	+MA	+MA	<i>L. monocytogenes</i>	+	+	+MB	+MB	<i>L. monocytogenes</i>	+	PA	G	1	c		
IPL	P23		Whipped cream puff	Ø	Ø	+MA	+MA	<i>L. monocytogenes</i>	+	+	+LB	+LB	<i>L. monocytogenes</i>	+	PA	G	1	c		
IPL	P26		Chocolate cream pastry	Ø	Ø	-LE	Ø	/	-	-	-ME	/	/	-	NA	G	1	c		
IPL	R12		Whipped cream puff	Ø	Ø	-LE	-LE	/	-	-	-ME	-ME	/	-	NA	G	1	c		
IPL	R17		Flan	Ø	Ø	Ø	Ø	/	-	-	-LE	-LE	/	-	NA	G	1	c		
IPL	U1		Chocolate profiteroles	+MB	+MB	+MA	+MA	<i>L. monocytogenes</i>	+	+	+MB	+MA	<i>L. monocytogenes</i>	+	PA	G	1	c		
IPL	U2		Chocolate mousse	+MA	+MA	+MA	+MB	<i>L. monocytogenes</i>	+	+	+MA	+MA	<i>L. monocytogenes</i>	+	PA	G	1	c		
IPL	U3		Saint Honoré pastry	+MA	+MA	+MA	+MA	<i>L. monocytogenes</i>	+	+	+LB	+MB	<i>L. monocytogenes</i>	+	PA	G	1	c		
ADRIA	4863	Plaque à croissant	Pastry	-	-	-	-	/	-	-	-	-	/	-	NA	G	1	c		
ADRIA	4868	Omelette orient	Omelette	st	-	-	-	/	-	-	-	-	/	-	NA	G	1	c		

COMPOSITE FOODS																					
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1						Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E						Agreement	Protocol	Category	Type		
				FRASER 1/2		FRASER		CONFIRMATION		RESULT <i>Listeria</i> spp	CONFIRMATION 24 LEB			FINAL RESULT							
				O&A1	P1	O&A2	P2	IDENTIFICATION			Brilliance 24h (IPL)	Brilliance 48h (IPL)	IDENTIFICATION								
ADRIA	4872	Mille feuilles	Pastry	st	st	st	st	/	-	-	-	-	/	-	NA	G	1	c			
ADRIA	4873	Omelette	Omelette	-	-	-	-	/	-	-	-	-	/	-	NA	G	1	c			
ADRIA	4874	Eclair au chocolat	Pastry	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+	H+	<i>L. monocytogenes</i>	+	PA	G	1	c			
ADRIA	4946	Blanc d'œuf en neige	Egg white	st	st	st	-	/	-	-	-	-	/	-	NA	G	1	c			
ADRIA	4947	Œuf entier liquide pasteurisé	Pasteurized liquid whole egg	st	st	st	st	/	-	-	st	st	/	-	NA	G	1	c			
ADRIA	4948	Œuf entier liquide pasteurisé	Pasteurized liquid whole egg	st	st	st	st	/	-	-	st	-	/	-	NA	G	1	c			
ADRIA	4949	Jaune d'œuf liquide pasteurisé	Pasteurized liquid yolk egg	st	st	st	st	/	-	-	st	st	/	-	NA	G	1	c			

MEAT PRODUCTS																				
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1						Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E						Agreement	Protocol	Category	Type	
				FRASER 1/2		FRASER		CONFIRMATION		RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB			<i>IDENTIFICATION</i>	FINAL RESULT				
				O&A1	P1	O&A2	P2	IDENTIFICATION				Brilliance 24h (IPL)	Brilliance 48h (IPL)	O&A 48h (Adria)						
IPL	A2	Beef minced meat		-LA	+LA	-MA	+MB	<i>L.innocua</i>		+	+	-MB	-MA	<i>L.innocua</i>	+	PA	G	2	a	
IPL	A3	Pork chops		-LA	+LA	-MA	+MB	<i>L.welshimeri</i>		+	+	-MA	-MA	<i>L.welshimeri</i>	+	PA	G	2	a	
IPL	A10	Thigh of guinea fowl		+MB	+MB	+MB	+MB	<i>L.monocytogenes</i> <i>L.welshimeri</i>		+	+	+MB	+HB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	2	a	
IPL	B8	Rumsteak		Ø	Ø	Ø	Ø	/		-	-	-LE	-LE	/	-	NA	G	2	a	
IPL	D4	Mincemeat		-LA	+LA	+MB	+MA	<i>L.innocua</i> ( <i>Bacillus</i> )		+	+	-MB	-MB	<i>L.innocua</i>	+	PA	G	2	a	
IPL	I11	Chicken leg		+LA	+LB	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>		+	+	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	2	a	
IPL	J12	Duck leg		Ø	Ø	Ø	Ø	/		-	+	Ø	Ø	/	-	PPNA	G	2	a	
IPL	K5	Chicken leg		Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	2	a	
IPL	K7	Chopped steak		Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	2	a	
IPL	M5	Beef mincemeat		+LA	+LA	+MA	+MA	<i>L.lavanovii</i>		+	+	+MA	+MB	<i>L.lavanovii</i>	+	PA	G	2	a	
IPL	M6	Beef mincemeat		+LA	+LA	+MA	+MA	<i>L.lavanovii</i>		+	-	-ME	/	/	-	ND	G	2	a	
IPL	N11	Horse mincemeat		+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	2	a	
IPL	P12	Beef mincemeat		+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+LA	+LB	<i>L.monocytogenes</i>	+	PA	G	2	a	
IPL	V8	Chicken		+LB	+LB	-MB	+MB	<i>L.monocytogenes</i> <i>L.welshimeri</i>		+	+	-MB	Fraser : -MA RLM : +HD	<i>L.monocytogenes</i> <i>L.welshimeri</i>	+	PA	G	2	a	
IPL	V9	Duck		-LA(2)	+LA(1)	-LA	+LA	<i>L.welshimeri</i>		+	+	-MB	-MB	<i>L.welshimeri</i>	+	PA	G	2	a	
IPL	AA1	Ground beef		-ME	-ME	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	2	a	
IPL	AA3	Pork chops		-LE	-LE	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	2	a	
IPL	I10	Pork offals		+LA	+LB	+MA	+MB	<i>L.monocytogenes</i>		+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	2	a	
ADRIA	6279	Steack haché	Ground beef	H- (3)	-	H-	+	<i>L.welshimeri</i>		+	-	-	-	/	-	ND	G	2	a	
ADRIA	6280	Steack haché	Ground beef	-	-	-	-	/		-	+	H-	H-	<i>L.innocua</i>	+	PD	G	2	a	
ADRIA	6456	Cote d'agneau	Lamb meat	-	-	-	-	/		-	-	-	-	/	-	NA	G	2	a	
ADRIA	6457	Escalope de veau	Veal meat	-	-	-	-	/		-	-	-	-	/	-	NA	G	2	a	
IPL	A4	Hamburger		+LA	+LB	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>		+	+	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	2	b	
IPL	A5	Beef balls		+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	2	b	
IPL	B1	Mincemeat		Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	2	b	
IPL	B5	Hamburger		Ø	Ø	-MA	+MA	<i>L.welshimeri</i>		+	+	-MA	-MA	<i>L.grayi</i>	+	PA	G	2	b	
IPL	D1	Hamburger		-LE	-LE	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	2	b	
IPL	J1	Tomato burger		Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	2	b	
IPL	K1	Hamburger		Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	2	b	
IPL	O5	Horse meat with curry		-LA(2)	+LA(2)	-MA	+HB	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	2	b	
IPL	AA2	Hamburger with onions		-LE	-LE	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	2	b	
IPL	B9	Foie gras		+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+MA	+HA	<i>L.monocytogenes</i>	+	PA	G	2	b	





DAIRY PRODUCTS																					
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1					Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E					FINAL RESULT	Agreement	Protocol	Category	Type			
				FRASER 1/2		FRASER		CONFIRMATION	RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB										
				O&A1	P1	O&A2	P2	IDENTIFICATION			O&A 24h (Adria)	O&A 48h (Adria)	IDENTIFICATION								
IPL	C11		Maroilles raw milk cheese	-LE	Ø	-LE	Ø	/	-	-	+LD	+LD	<i>L.innocua</i>	-	NA	G	3	a			
IPL	C16		Epoisses raw milk cheese	+LB	+LB	+MB	+MB	<i>L.monocytogenes L.innocua</i>	+	-	-ME	/	/	-	ND	G	3	a			
IPL	D3		Maroilles raw milk cheese	+LB	+LB	+MB	+MB	<i>L.monocytogenes L.innocua</i>	+	+	+MB	+MB	<i>L.monocytogenes L.innocua</i>	+	PA	G	3	a			
IPL	E1		Maroilles raw milk cheese	+LA	+LA	+LB	+LB	<i>L.monocytogenes L.innocua</i>	+	+	+MB	+MB	<i>L.monocytogenes L.innocua</i>	+	PA	G	3	a			
IPL	E2		Maroilles raw milk cheese	+MA	+MA	+MB	+MB	<i>L.monocytogenes L.innocua</i>	+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	3	a			
IPL	E5		Comté raw milk cheese	-MA	+MB	-MB	+MB	<i>L.innocua</i>	+	-	-ME	-ME	Ø	-	ND	G	3	a			
IPL	H6		Raw milk cheese	+MA	+MA	+MA	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	3	a			
IPL	J22		Camembert de Normandie raw milk cheese	Ø	-LE	Ø	Ø	/	-	+	+LA	+MB	<i>L.monocytogenes</i>	+	PD	G	3	a			
IPL	J23		Reblochon raw milk cheese	-LE	-ME	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes L.innocua</i>	+	PA	G	3	a			
IPL	K12		Camembert raw milk cheese	-LE	-LE	Ø	-LE	/	-	-	-LE	/	/	-	NA	G	3	a			
IPL	K13		Camembert de Normandie raw milk cheese	Ø	-ME	-ME	-ME	/	-	-	-HE	/	/	-	NA	G	3	a			
IPL	N19		Brie de Meaux raw milk cheese	Ø	Ø	-LE	-LE	/	-	-	-ME	/	/	-	NA	G	3	a			
IPL	P8		Brie de Meaux raw milk cheese	Ø	Ø	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	3	a			
IPL	P13		Reblochon raw milk cheese	-MA	+MA	-MA	+MA	<i>L.innocua</i>	+	+	-MB	-MB	<i>L.innocua</i>	+	PA	G	3	a			
IPL	Q7		Reblochon raw milk cheese	-ME	Ø	-LE	Ø	/	-	-	-ME	/	/	-	NA	G	3	a			
IPL	Q8		Epoisses raw milk cheese	-LE	-LE	-LE	-LE	/	-	+	-MD	-MD	<i>L.welshimeri</i>	+	PD	G	3	a			
IPL	T7		Munster raw milk cheese	Ø	-LE	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	3	a			
IPL	T12		Roquefort cheese	Ø	Ø	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	3	a			
ADRIA	4950	Brie de Meaux lait cru	Raw milk cheese	st	st	-	st	/	-	-	-	-	/	-	NA	G	3	a			
ADRIA	4951	Morbier lait cru de vache	Raw milk cheese	-	-	-	-	/	-	-	-	-	/	-	NA	G	3	a			

DAIRY PRODUCTS																							
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1						Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E						FINAL RESULT	Agreement	Protocol	Category	Type			
				FRASER 1/2		FRASER		CONFIRMATION		RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB			IDENTIFICATION								
				O&A1	P1	O&A2	P2	IDENTIFICATION				Brilliance 24h (IPL)	Brilliance 48h (IPL)	O&A 24h (Adria)	O&A 48h (Adria)								
ADRIA	4952	Kalbach lait cru	Raw milk cheese	-	st	-	-	/	-	-	-	-	-	-	/	-	NA	G	3	a			
IPL	B11		Raw milk	Ø	Ø	Ø	Ø	/	-	-	-LE	Ø	/	/	/	-	NA	G	3	b			
IPL	C17		Raw milk	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	/	/	-	NA	G	3	b			
IPL	C20		Raw milk	-LE	Ø	Ø	Ø	/	-	-	Ø	/	/	/	/	-	NA	G	3	b			
IPL	N3		Raw milk	+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MA	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	<i>L.monocytogenes</i> <i>L.innocua</i>	+ + <i>L.innocua</i>	+	PA	G	3	b			
IPL	N4		Raw milk	+LA	+LB	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	+	-LD	-LD	<i>L.innocua</i>	<i>L.innocua</i>	+ + <i>L.innocua</i>	+	PA	G	3	b			
IPL	R2		Raw milk	-LE	-LE	-ME	-ME	/	-	-	-ME	-ME	/	/	/	-	NA	G	3	b			
IPL	R16		Raw milk	-LE	-LE	-LE	-LE	/	-	-	-ME	-ME	/	/	/	-	NA	G	3	b			
ADRIA	4942	Lait cru fermier	Raw milk	H-	+	H-	+	<i>L. innocua</i>	+	-	-	-	/	/	/	-	ND	G	3	b			
ADRIA	4943	Lait cru fermier	Raw milk	st	st	-	-	/	-	-	-	-	/	/	/	-	NA	G	3	b			
ADRIA	4944	Lait cru fermier	Raw milk	st	st	-	-	/	-	-	-	-	/	/	/	-	NA	G	3	b			
ADRIA	4945	Lait cu de vache jersiaise	Raw milk	-	st	-	-	/	-	-	-	-	/	/	/	-	NA	G	3	b			
ADRIA	5395	Lait cru de brebis	Raw milk	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+	H+	<i>L. monocytogenes</i>	<i>L. monocytogenes</i>	+ + <i>L. monocytogenes</i>	+	PA	G	3	b			
ADRIA	5396	Lait cru de brebis	Raw milk	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+	H+	<i>L. monocytogenes</i>	<i>L. monocytogenes</i>	+ + <i>L. monocytogenes</i>	+	PA	G	3	b			
ADRIA	5397	Lait cru de brebis	Raw milk	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+/H-	H+/H-	<i>L.monocytogenes</i> <i>L.innocua</i>	<i>L.monocytogenes</i> <i>L.innocua</i>	+ + H+/H-	+	PA	G	3	b			
ADRIA	5398	Lait cru de brebis	Raw milk	st	-	-	-	/	-	-	-	-	/	/	/	-	NA	G	3	b			
ADRIA	5399	Lait cru de brebis	Raw milk	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+	H+	<i>L. monocytogenes</i>	<i>L. monocytogenes</i>	+ + <i>L. monocytogenes</i>	+	PA	G	3	b			
ADRIA	5400	Lait cru de brebis	Raw milk	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+	H+	<i>L. monocytogenes</i>	<i>L. monocytogenes</i>	+ + <i>L. monocytogenes</i>	+	PA	G	3	b			
ADRIA	5401	Lait cru de brebis	Raw milk	H+ (8)	+	H+	+	<i>L. monocytogenes</i>	+	+	H+	H+	<i>L. monocytogenes</i>	<i>L. monocytogenes</i>	+ + <i>L. monocytogenes</i>	+	PA	G	3	b			
ADRIA	5402	Lait cru de brebis	Raw milk	H+	+	H+	+	<i>L. monocytogenes</i>	+	+	H+/H-	H+/H-	<i>L.monocytogenes</i> <i>L.innocua</i>	<i>L.monocytogenes</i> <i>L.innocua</i>	+ + H+/H-	+	PA	G	3	b			
ADRIA	5959	Lait cru fermier	Raw milk	-	-	-	-	/	-	-	-	-	/	/	/	-	NA	G	3	b			
IPL	H5	Brie cheese	+LA	+LA	+HA	+HA	<i>L.monocytogenes</i>	+	+	+HA	+MA	<i>L.monocytogenes</i> <i>L.innocua</i>	<i>L.monocytogenes</i> <i>L.innocua</i>	+ + <i>L. innocua</i>	+	PA	G	3	c				
IPL	H8	Brie cheese	+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	<i>L.monocytogenes</i>	+ + <i>L. monocytogenes</i>	+	PA	G	3	c				
IPL	J21	Camembert pasteurized milk cheese	Ø	-LE	Ø	Ø	/	-	+	+LB	+MB	<i>L.monocytogenes</i>	<i>L.monocytogenes</i>	+ + <i>L. monocytogenes</i>	+	PD	G	3	c				
IPL	K14	Camembert cheese	-LE	-LE	-HE	-HE	/	-	-	-ME	/	/	/	/	-	NA	G	3	c				

DAIRY PRODUCTS																					
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1					Alternative method: BAX® System PCR Assay for Genus Listeria 24E					FINAL RESULT	Agreement	Protocol	Category	Type			
				FRASER 1/2		FRASER		CONFIRMATION	RESULT <i>Listeria spp</i>	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB										
				O&A1	P1	O&A2	P2	IDENTIFICATION			O&A 24h (Adria)	O&A 48h (Adria)	IDENTIFICATION								
IPL	N15		Soft cheese	+MB	+MB	+LA	+LB	<i>L.monocytogenes</i>	+	+	+MD	+MB	<i>L.monocytogenes</i>	+	PA	G	3	c			
IPL	N16		Camembert cheese	-LE	Ø	Ø	Ø	/	-	+	+LB	+MB	<i>L.monocytogenes</i>	+	PD	G	3	c			
IPL	N17		Light cheese	+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+HB	<i>L.monocytogenes</i>	+	PA	G	3	c			
IPL	N18		Gouda cheese	+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+LB	+MB	<i>L.monocytogenes</i>	+	PA	G	3	c			
IPL	P9		Coulommier pasteurized milk cheese	Ø	Ø	Ø	Ø	/	-	-	-LE	/	/	-	NA	G	3	c			
IPL	P14		Soft cheese	-LE	-LE	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	3	c			
IPL	C14		Ossau Iraty cheese	+MA	+MA	+MB	+MB	<i>L.monocytogenes L.innocua</i>	+	+	-MB	+MB	<i>L.monocytogenes L.innocua</i>	+	PA	G	3	c			
IPL	J24		Blue cheese	Ø	-LE	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	3	c			
IPL	K15		Goat's cheese	Ø	-LE	-LE	-ME	/	-	-	Ø	/	/	-	NA	G	3	c			
IPL	P5		Ewe's milk cheese	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	3	c			
IPL	P7		Angeroux	Ø	Ø	Ø	Ø	/	-	-	-LE	/	/	-	NA	G	3	c			
IPL	P10		Goat's cheese	-ME	-LE	-LE	-LE	/	-	-	-LE	/	/	-	NA	G	3	c			
IPL	Q9		Goat's cheese	-LE	-LE	Ø	Ø	/	-	-	-LE	/	/	-	NA	G	3	c			
IPL	R18		Petit Bethmale cheese	-LE	Ø	-LE	-LE	/	-	-	-ME	-ME	/	-	NA	G	3	c			
IPL	T3		Goat's cheese	Ø	Ø	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	3	c			
IPL	T4		Goat's cheese	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	3	c			
IPL	T8		Goat's cheese	-LE	Ø	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	3	c			
IPL	T14		Goat's cheese	-LE	-LE	-LE	-LE	/	-	-	-ME	/	/	-	NA	G	3	c			
IPL	T18		Goat's cheese	-LE	-LE	-ME	-ME	/	-	-	-LE	/	/	-	NA	G	3	c			
IPL	F10		Skim milk powder	-ME	-ME	-ME	-ME	/	-	+	-HE	-ME	/	-	PPN A	G	3	c			
IPL	F11		Skim milk powder	-LE	-LE	Ø	-LE	/	-	-	-ME	Fraser : Ø	/	-	NA	G	3	c			
IPL	G1		Milk powder	-LE	-LE	Ø	-ME	/	-	-	Ø	/	/	-	NA	G	3	c			
IPL	G2		Milk powder	-LE	-LE	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	3	c			
IPL	G3		Milk powder	Ø	Ø	-LE	-LE	/	-	-	-ME	/	/	-	NA	G	3	c			
IPL	G4		Milk powder	-ME	-LE	-LE	-ME	/	-	-	-ME	/	/	-	NA	G	3	c			
IPL	G5		Milk powder	-ME	-LE	-LE	-ME	/	-	-	-ME	/	/	-	NA	G	3	c			
IPL	N20		Cream	+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+LA	+LA	<i>L.monocytogenes</i>	+	PA	G	3	c			
IPL	P16		Milk powder	-LE	-LE	-LE	-LE	/	-	+	+LB	+LB	<i>L.monocytogenes</i>	+	PD	G	3	c			

FRUITS AND VEGETABLES																							
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1						Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E						FINAL RESULT	Agreement	Protocol	Category	Type			
				FRASER 1/2		FRASER		CONFIRMATION		RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB			IDENTIFICATION								
				O&A1	P1	O&A2	P2	IDENTIFICATION				Brilliance 24h (IPL)	Brilliance 48h (IPL)										
IPL	Q1	Frozen green peas		Ø	Ø	Ø	Ø	/	-	-	-LE	/	/	-	NA	G	4	a					
IPL	R1	Frozen spinach		-MA	+MA	-MA	+MA	<i>L.innocua</i>	+	+	-MB	-MB	<i>L.innocua</i>	+	PA	G	4	a					
IPL	R14	Frozen coco beans		Ø	Ø	-ME	-ME	/	-	-	-ME	-ME	<i>Bacillus</i>	-	NA	G	4	a					
IPL	S12	Frozen spinach		+MA	+MA	+MB*	+MB*	<i>L.monocytogenes</i> <i>L.innocua</i>	+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	4	a					
IPL	T15	Frozen carrots		-LE	-LE	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	4	a					
IPL	T17	Frozen leeks		Ø	-LE	-ME	-ME	/	-	-	Ø	/	/	-	NA	G	4	a					
IPL	T19	Frozen onions		Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	4	a					
IPL	V6	Frozen spinaches		Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	4	a					
IPL	V10	Frozen cauliflower		Ø	Ø	Ø	Ø	/	-	-	-LE	/	/	-	NA	G	4	a					
IPL	V11	Frozen zucchini		Ø	-LE	-ME	-ME	/	-	-	-LE	/	/	-	NA	G	4	a					
IPL	F6	Green peas		Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	4	a					
IPL	F7	Spinach		+MA	+LA	+MA	+MB	<i>L.monocytogenes</i>	+	+	+MA	+MB	<i>L.monocytogenes</i>	+	PA	G	4	a					
IPL	J15	Green peas		+LB	+LB	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	a					
IPL	L9	<b>Broccoli</b>		+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+HA	+HA	<i>L.monocytogenes</i>	+	PA	G	4	a					
IPL	P3	Red cabbage		Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	4	a					
IPL	P4	White cabbage		Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	4	a					
IPL	P18	Radish		Ø	Ø	Ø	Ø	/	-	-	-LE	/	/	-	NA	G	4	a					
IPL	Y3	<b>Broccoli</b>		Ø	-LE	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	a					
IPL	Y10	Cauliflower		Ø	Ø	Ø	-LE	/	-	+	-LE	-LE (5b) Fraser : Ø	/	-	PPNA	G	4	a					
ADRIA	6278	<b>Champignon blanc</b>	Mushrooms	H-	+	H-	+	<i>L. innocua</i>	+	+	H-	H-	<i>L. innocua</i>	+	PA	G	4	a					
IPL	B10	Frozen French fries		+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+HB	<i>L.monocytogenes</i> <i>L.grayi</i>	+	PA	G	4	b					
IPL	C2	Frozen French fries		+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	C9	Frozen French fries		+MB	+MB	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	C13	Frozen French fries		+MA	+MA	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	+	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	4	b					
IPL	C15	Frozen French fries		-LE	-LE	-ME	-ME	/	-	+	+MA	+MB	<i>L.monocytogenes</i>	+	PD	G	4	b					
IPL	E11	Frozen French fries		+MA	+LA	+MB	+LB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	F4	Frozen French fries		+MB	+MB	+HB	+HB	<i>L.monocytogenes</i>	+	+	+MA	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	F9	Frozen French fries		+MA	+HB	+MB	+HB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	J13	Frozen French fries		+MB	+MB	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	Q6	Frozen French fries		+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MA	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	L11	<b>Grated carrots</b>		+LA	+LA	+MA	+HA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	L23	<b>Carrots and Celery</b>		+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	P21	Salad		Ø	Ø	Ø	-LE	/	-	-	Ø	/	/	-	NA	G	4	b					
IPL	T5	Grated carrots		Ø	Ø	Ø	Ø	/	-	-	-LE	/	/	-	NA	G	4	b					
IPL	W5	Salad		+LA	+LA	+MB	+MB	<i>Livanovii</i>	+	+	+LB	+LD	<i>Livanovii</i>	+	PA	G	4	b					
IPL	W6	Salad		+LA	+LA	+MB	+MB	<i>Livanovii</i>	+	+	+MB	+HB	<i>Livanovii</i>	+	PA	G	4	b					
IPL	W7	Salad		-MA	+MA	-MA	+MA	<i>L.innocua</i>	+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	4	b					
IPL	W8	Salad		-MA	+MA	-MA	+MA	<i>L.innocua</i>	+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	4	b					
IPL	W9	Salad		Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	4	b					
IPL	Z11	<b>Mixed vegetables</b>		Ø	Ø	+MA	+MA	<i>L.monocytogenes</i>	+	+	+HB	+MB	<i>L.monocytogenes</i>	+	PA	G	4	b					
IPL	T0-1	Mixed vegetables		+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	4	b					
ADRIA	6281	Duo de carottes	Carrots mix	st	st	-	-	/	-	-	-	-	/	-	NA	G	4	b					
ADRIA	6282	Poireaux émincés	Sliced leeks	st	-	-	-	/	-	-	-	-	/	-	NA	G	4	b					

Laboratory		Sample N°	Products (French name)	Products	FRUITS AND VEGETABLES								Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E							Agreement	Protocol	Category	Type				
					Reference method: ISO 11290-1/A1		FRASER 1/2		FRASER		CONFIRMATION		RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB			IDENTIFICATION	FINAL RESULT								
					O&A1	P1	O&A2	P2	IDENTIFICATION						Brilliance 24h (IPL)	Brilliance 48h (IPL)	O&A 24h (Adria)	O&A 48h (Adria)									
ADRIA	6284	Trio de poivrons	Peppers mix		st	-	-	-	/		-		-		-	-	-	/	-	NA	G	4	b				
ADRIA	6285	Lait cru	Raw milk		st	st	st	st	/		-		-		-	-	-	/	-	NA	G	4	b				
IPL	C10		Frozen fried potatoes	+LA	+LA	+MB	+MB		<i>L.monocytogenes</i>		+		+		+MD	+MB		<i>L.monocytogenes</i>	+	PA	G	4	c				
IPL	T16		Frozen marinated peppers	Ø	Ø	Ø	Ø		/		-		-		Ø	/		/	-	NA	G	4	c				
IPL	C3		Fried mixed vegetables	+LA	+LA	+MA	+MB		<i>L.monocytogenes</i>		+		+		+MA	+MA		<i>L.monocytogenes</i>	+	PA	G	4	c				
IPL	C4		Cakes cauliflower - broccoli	-LA	+LA	+MB	+MB		<i>L.innocua</i> <i>L.monocytogenes</i>		+		+		+MB	-MA		<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	4	c				
IPL	D5		Potatoes	+MA	+MA	+MB	+MB		<i>L.monocytogenes</i>		+		+		+MA	+MA		<i>L.monocytogenes</i>	+	PA	G	4	c				
IPL	F8		Tabbouleh	Ø	Ø	Ø	Ø		/		-		-		-LE	/		/	-	NA	G	4	c				
IPL	I12		Fried mixed vegetables	+MA	+MB	+MB	+MB		<i>L.monocytogenes</i>		+		-		-ME	-ME		/	-	ND	G	4	c				
IPL	L10		Rice salad	+MA	+MA	+MA	+HA		<i>L.monocytogenes</i>		+		+		+MB	+MB		<i>L.monocytogenes</i>	+	PA	G	4	c				
IPL	P17		Mashed potatoes	Ø	Ø	Ø	Ø		/		-		-		Ø	/		/	-	NA	G	4	c				
IPL	P19		Cucumber salad	-LE	-LE	-ME	-ME		/		-		-		-LE	/		/	-	NA	G	4	c				
IPL	P20		Grated carrots salad	Ø	Ø	Ø	Ø		/		-		-		Ø	/		/	-	NA	G	4	c				
IPL	P22		Rice salad	Ø	Ø	-LE	-ME		/		-		-		+LA	+MB		<i>L.monocytogenes</i>	+	PD	G	4	c				
IPL	P24		Fried mixed vegetables	Ø	Ø	Ø	Ø		/		-		-		-LE	/		/	-	NA	G	4	c				
IPL	P25		Celery with mayonnaise	Ø	Ø	Ø	Ø		/		-		-		Ø	Fraser : Ø		/	-	PPNA	G	4	c				
IPL	Q3		Fried mixed vegetables	-LE	-LE	-ME	-ME		/		-		-		-ME	/		/	-	NA	G	4	c				
IPL	Q4		Fried mixed vegetables	-LE	-LE	Ø	Ø		/		-		-		-MD	-MD		<i>L.innocua</i>	+	PD	G	4	c				
IPL	Q5		Fried mixed vegetables	Ø	Ø	-ME	-ME		/		-		-		-ME	/		/	-	NA	G	4	c				
IPL	T6		Guacamole	Ø	Ø	Ø	Ø		/		-		-		Ø	/		/	-	NA	G	4	c				
IPL	T9		Beetroot salad	Ø	Ø	Ø	Ø		/		-		-		Ø	/		/	-	NA	G	4	c				
IPL	T10		Houmous	-LE	-LE	Ø	-LE		/		-		-		-LE	/		/	-	NA	G	4	c				
IPL	U10		Salad with cabbage and carrots	-LB	+LB	-MB	+MB		<i>L.welshimeri</i>		+		+		-ME	Fraser : -MA		<i>L.welshimeri</i>	+	PA	G	4	c				
IPL	U11		Fruit salad	-LB	+LB	-MB	+MB		<i>L.welshimeri</i>		+		+		-LB	-LB		<i>L.welshimeri</i>	+	PA	G	4	c				
IPL	U12		Zucchini salad	-LB	+LB	-LB	+LB		<i>L.welshimeri</i>		+		+		-ME	Fraser : -MD		<i>L.welshimeri</i>	+	PA	G	4	c				
IPL	W14		Salad with potatoes	+LB	+LB	+MB	+MB		<i>L.livanovii</i>		+		+		+MB	+LB		<i>L.livanovii</i>	+	PA	G	4	c				
IPL	Y1		Zucchinis flan	Ø	Ø	Ø	Ø		/		-		-		-LE	/		/	-	NA	G	4	c				
IPL	Y2		Spinaches with cream	Ø	Ø	-LE	-LE		/		-		-		-LE	/		/	-	NA	G	4	c				
IPL	Y6		Cauliflower flan	Ø	Ø	Ø	Ø		/		-		-		Ø	/		/	-	NA	G	4	c				
IPL	Y8		Tomatoes salad	Ø	+LD	-MA	+MB		<i>L.welshimeri</i>		+		+		-MB	-LB		<i>L.welshimeri</i>	+	PA	G	4	c				
IPL	Z8		Mashed potatoes	+LB	+LB	+MA	+MB		<i>L.monocytogenes</i>		+		+		+MB	+HB		<i>L.monocytogenes</i>	+	PA	G	4	c				
IPL	Z9		Mashed celery	+MA	+LD	+MB	+MB		<i>L.monocytogenes</i>		+		+		+HA	+MA		<i>L.monocytogenes</i>	+	PA	G	4	c				
IPL	Z10		Lenses salad	+MA	+MA	+MB	+MB		<i>L.monocytogenes</i>		+		+		+HA	+MB		<i>L.monocytogenes</i>	+	PA	G	4	c				

Laboratory	Sample N°	Products (French name)	Products	SEAFOOD								Alternative method: BAX® System PCR Assay for Genus Listeria 24E						Agreement	Protocol	Category	Type			
				Reference method: ISO 11290-1/A1				CONFIRMATION			RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB			IDENTIFICATION	FINAL RESULT							
				FRASER 1/2		FRASER		O&A2	P2	IDENTIFICATION			Brilliance 24h (IPL)	Brilliance 48h (IPL)										
				O&A1	P1	O&A2	P2	IDENTIFICATION		O&A 24h (Adria)	O&A 48h (Adria)													
IPL	B7		Whiting fillet	-MB	+MB	-MA	+MA	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	5	a					
IPL	C6		Herring fillet	-MB	+MB	-MA	+MA	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	5	a					
IPL	C7		Cod fillet	Ø	Ø	Ø	Ø	/		-	+	+HD	+MB	<i>L.monocytogenes</i>	+	PD	G	5	a					
IPL	C8		Haddock fillet	Ø	-LE	Ø	-LE	/		-	-	-ME	-ME	/	-	NA	G	5	a					
IPL	C18		Salmon fillet	Ø	-LE	Ø	Ø	/		-	-	-HE	/	/	-	NA	G	5	a					
IPL	D15		Herring fillet	Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	5	a					
IPL	D17		Herring fillet	Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	5	a					
IPL	E7		Herring fillet	-LA	+LA	-LD	+LA(1)	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	5	a					
IPL	E10		Cod fillet	Ø	Ø	Ø	Ø	/		-	-	-LE	-LE	/	-	NA	G	5	a					
IPL	F5		Fish cubes	Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	5	a					
IPL	I8		Herring fillet	+4LA	+4LB	+MA	+MA	<i>L.monocytogenes</i>		+	+	+LB	+LD	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	J3		Herring fillet	Ø	-LE	Ø	Ø	/		-	+	-LE	-LE	/	-	PPNA	G	5	a					
IPL	L12		Pollock fillet	+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+LB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	V7		Whiting fillet	Ø	Ø	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AB3		Halibut fillet	Ø	Ø	Ø	Ø	/		-	-	+LD?	-LE	<i>Bacillus</i>	-	NA	G	5	a					
IPL	AB11		Tilapia fillet	-MA	+MA	-MA	+MA	<i>L.innocua</i>		+	+	-MA	-MB	<i>L.innocua</i>	+	PA	G	5	a					
IPL	AC1		Cod fillet	Ø	Ø	-ME	-ME	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AC2		Salmon fillet	Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	5	a					
IPL	AC4		Fish fillet	-LE	-LE	-LE	-LE	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AC5		Hake fillet	-LE	-LE	-LE	-LE	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AD10		Saithe fillet	Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	5	a					
IPL	AD14		Haddock fillet	Ø	Ø	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AD16		Whiting fillet	Ø	Ø	Ø	-ME	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AD17		Perch fillet	Ø	Ø	-LE	-LE	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AD18		Fish fillet	Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	5	a					
IPL	AD19		Salmon fillet	Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	5	a					
IPL	E8		Shrimps	Ø	Ø	+LD	+LB	<i>L.monocytogenes</i>		+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	I1		Mixed shellfish	+LD	+MD	+MA	+MA	<i>L.monocytogenes</i> <i>L.innocua</i>		+	+	+LB	+MD	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	5	a					
IPL	I9		Shrimps	+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>		+	+	+HB	+MA	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	J9		Shrimps	+LA	+LB	+MA	+MA	<i>L.monocytogenes</i>		+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	J16		Cocktail seafood	+LA	+LD	+LB	+LB	<i>L.monocytogenes</i>		+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	O9		Cocktail seafood	+LB	+LD	+MB	+MB	<i>L.monocytogenes</i>		+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	AA5		Cocktail seafood	+LB	+LD(1)	+MA	+MA	<i>L.monocytogenes</i>		+	-	-ME	+MD(5b)	<i>L.monocytogenes</i>	-	ND	G	5	a					
IPL	AB4		Shrimps	Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	5	a					
IPL	AB5		Shrimps	Ø	Ø	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AB8		Shrimps	Ø	Ø	+MD	+LD	<i>L.monocytogenes</i>		+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	AB9		Shrimps	Ø	Ø	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	5	a					
IPL	AB12		Gambas	+LA	-ME	+MB	+MB	<i>L.monocytogenes</i>		+	+	+MB	+LB	<i>L.monocytogenes</i>	+	PA	G	5	a					
IPL	A8		Smoked salmon	+MA	+MA	+MA	+MA	<i>L.monocytogenes</i> <i>L.innocua</i>		+	+	+MA	+MA	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	S	5	b					
IPL	A9		Smoked salmon	Ø	Ø	Ø	Ø	/		-	+	-ME	+MA	<i>L.monocytogenes</i>	+	PD	S	5	b					
IPL	A11		Smoked salmon	Ø	Ø	Ø	Ø	/		-	-	Ø	Fr: +LA	<i>L.monocytogenes</i>	-	NA	S	5	b					

SEAFOOD																				
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1						Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E						Agreement	Protocol	Category	Type	
				FRASER 1/2		FRASER		CONFIRMATION		RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB			IDENTIFICATION	FINAL RESULT				
				O&A1	P1	O&A2	P2	IDENTIFICATION				Brilliance 24h (IPL)	Brilliance 48h (IPL)	O&A 24h (Adria)	O&A 48h (Adria)					
IPL	A12		Smoked salmon	Ø	Ø	Ø	Ø	/	-	-	-LE			/	-	NA	S	5	b	
IPL	A13		Smoked salmon Scotland	Ø	Ø	Ø	Ø	/	-	-	Ø	Ø Fr: Ø		/	-	NA	S	5	b	
IPL	A14		Smoked salmon Scotland	+MA	+LA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	A16		Smoked herring	Ø	Ø	Ø	Ø	/	-	+	Ø	Ø Fr: Ø		/	-	PPNA	S	5	b	
IPL	A17		Salmon tartar	-MA	+LA	-HA	+HA	<i>L.innocua</i>	+	+	-ME	Ø F=+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	S	5	b		
IPL	A18		Smoked salmon	+MA	+LB	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MA	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	B1		Smoked salmon	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	S	5	b		
IPL	B2		Smoked salmon	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	S	5	b		
IPL	B3		Smoked salmon	Ø	Ø	Ø	Ø	/	-	-	+LD	+LD	<i>L.monocytogenes</i>	-	NA	S	5	b		
IPL	B4		Smoked salmon	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	S	5	b		
IPL	B5		Smoked herring	Ø	-LE	Ø	-LE	/	-	-	-ME	/	/	-	NA	S	5	b		
IPL	B6		Salmon roulade	Ø	Ø	Ø	-LE	/	-	+	+MB	+MB	<i>L.monocytogenes</i>	+	PD	S	5	b		
IPL	B7		Smoked salmon	Ø	Ø	Ø	Ø	/	-	+	+MB	+MB	<i>L.monocytogenes</i>	+	PD	S	5	b		
IPL	C16		Smoked sprats	+MD	+MD	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	C17		Smoked sprats	+MD	+MD	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	D6		Smoked haddock	+LB	+LB	°LB	+LB(2)	<i>L.monocytogenes</i> <i>L.innocua</i>	+	+	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	S	5	b		
IPL	D12		Smoked herring	Ø	-LE	Ø	-LE	/	-	-	-ME	/	/	-	NA	S	5	b		
IPL	D17		Smoked salmon	Ø	Ø	Ø	-LE	/	-	+	-LE	-LE Fr : +LB	<i>L.monocytogenes</i>	+	PD	S	5	b		
IPL	D18		Smoked trout	+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	E1		Smoked salmon	Ø	+LA(2)	+MA	+MB	<i>L.monocytogenes</i>	+	+	Ø	Ø Fr : +MA	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	E2		Smoked salmon	Ø	Ø	Ø	Ø	/	-	+	Ø	Ø Fr : Ø	/	-	PPNA	S	5	b		
IPL	E3		Smoked salmon	+LA	+LA	+MA	+MB	<i>L.monocytogenes</i>	+	+	+LB	+LB	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	E4		Smoked trout	+LA	+MB	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	E5		Smoked trout	+LA	+MA	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	E6		Smoked salmon Scotland	+LA(1)	+LB	+MA	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MA	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	E7		Smoked trout	+LA	+LA	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	E8		Smoked halibut	+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MA	<i>L.monocytogenes</i>	+	PA	S	5	b		
IPL	E9		Smoked salmon Scotland	Ø	Ø	Ø	Ø	/	-	+	+MA	+MA	<i>L.monocytogenes</i>	+	PD	S	5	b		
IPL	E10		Smoked salmon Ireland	Ø	Ø	Ø	Ø	/	-	-	+LB	+LB	<i>L.monocytogenes</i>	-	NA	S	5	b		
ADRIA	5940	Filets de maquereaux fumés	Smoked fish	st	-	st	st	/	-	-	st	-	/	-	NA	S	5	b		
ADRIA	5941	Hareng fumé	Smoked fish	st	st	st	st	/	-	-	st	-	/	-	NA	S	5	b		
ADRIA	5942	Trite fumée	Smoked salmon	st	st	st	st	/	-	-	st	-	/	-	NA	S	5	b		

Laboratory	Sample N°	Products (French name)	Products	SEAFOOD											Agreement	Protocol	Category	Type				
				Reference method: ISO 11290-1/A1					Alternative method: BAX® System PCR Assay for Genus <i>Listeria</i> 24E													
				FRASER 1/2		FRASER		CONFIRMATION	RESULT <i>Listeria</i> spp	BAX® <i>Listeria</i>	CONFIRMATION 24 LEB		IDENTIFICATION	FINAL RESULT								
				O&A1	P1	O&A2	P2	IDENTIFICATION			Brilliance 24h (IPL)	Brilliance 48h (IPL)										
IPL	AA4		Breaded fish fillet	+MA	+MB	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	c				
IPL	AB6		Plateau of shellfish	-ME	-ME	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	5	c				
IPL	AB10		Plateau of shellfish	+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	5	c				
IPL	D13		Fish with bordelaise sauce	+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>	+	-	-LE	-LE	Ø	-	ND	G	5	c				
IPL	D20		Raw marinated cod	Ø	-LE	-MB	+MB	<i>L.innocua</i>	+	-	-MD	/	<i>L.innocua</i>	-	ND	G	5	c				
IPL	F1		Frozen calamari precooked	+MA	+MB	+MB	+HB	<i>L.monocytogenes</i>	+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	5	c				
IPL	F3		Ready-to-eat fish	-LE	-LE	-HE	-LE	/	-	-	-LE	/	/	-	NA	G	5	c				
IPL	I2		Cooked mussels	Ø	Ø	Ø	Ø	/	-	-	-LE	/	/	-	NA	G	5	c				
IPL	J4		Marinated herring fillet	Ø	-LE	Ø	-ME	/	-	-	-ME	/	/	-	NA	G	5	c				
IPL	J6		Salmon Carpaccio	Ø	-LE	Ø	-LE	/	-	+	-MA	-MA	<i>L.innocua</i>	+	PD	G	5	c				
IPL	J14		Salmon paupiettes with Saint Jacques	+LA	+LB	+MA	+MA	<i>L.monocytogenes</i> <i>L.welshimeri</i>	+	+	-LD	-LD	<i>L.welshimeri</i>	+	PA	G	5	c				
IPL	L13		Fish with bordelaise sauce	+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MA	+MB	<i>L.monocytogenes</i>	+	PA	G	5	c				
IPL	L14		Frozen breaded fish	+MB	+MB	+MB	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	c				
IPL	L15		Frozen breaded fish	+MB	+MB	+MA	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	c				
IPL	L17		Salmon with hollandaise sauce	+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	c				
IPL	L20		Salmon dish	+LA	+LA	+MA	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	+	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	5	c				
IPL	O7		Raw marinated cod	-LE	-LE	Ø	Ø	/	-	+	-MB	-MB	<i>L.innocua</i>	+	PD	G	5	c				
IPL	P6		Shrimps accrass	-ME	-LE	-LE	-ME	/	-	-	-LE	/	/	-	NA	G	5	c				
IPL	AB1		Ready-to-eat fish	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	5	c				
IPL	AB2		Cooked calamari	+MB	+MB	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	+	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	5	c				
IPL	AB7		Sushis	-LE	Ø	-LE	Ø	/	-	+	+MA	+MA	<i>L.monocytogenes</i>	+	PD	G	5	c				
IPL	AB13		Brochette fish	+LA	+LA	+MB	+MA	<i>L.monocytogenes</i> <i>L.innocua</i>	+	+	+LA	+MA	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	5	c				
IPL	AC3		Salmon with vegetables	Ø	-LE	Ø	Ø	/	-	-	-ME	/	/	-	NA	G	5	c				
IPL	AC6		Salmon fillet sauce Bellevue	+MB	+MA	+MA	+MB	<i>L.monocytogenes</i>	+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	5	c				
IPL	AD11		Ready-to-eat fish	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	5	c				
IPL	AD12		Salmon fillet with sorrel sauce	Ø	-LE	-LE	-LE	/	-	-	Ø	/	/	-	NA	G	5	c				
IPL	AD13		Cod fillet with florentine sauce	Ø	Ø	Ø	Ø	/	-	-	Ø	/	/	-	NA	G	5	c				

ENVIRONMENTAL SAMPLES																								
Laboratory	Sample N°	Products (French name)	Products	Reference method: ISO 11290-1/A1						RESULT Listeria spp	BAX® Listeria	Alternative method: BAX® System PCR Assay for Genus Listeria 24E			FINAL RESULT	Agreement	Protocol	Category	Type					
				FRASER 1/2		FRASER		CONFIRMATION				CONFIRMATION 24 LEB												
				O&A1	P1	O&A2	P2	IDENTIFICATION				Brilliance 24h (IPL)	Brilliance 48h (IPL)	IDENTIFICATION										
IPL	K9		Water of wash lines pea	+MA	+MA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+HA	+HA	<i>L.monocytogenes</i>	+	PA	G	6	a					
IPL	R4		Arrival new water	Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	6	a					
IPL	R5		Water washbasin reception	Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	6	a					
IPL	R6		Washing 1	Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	6	a					
IPL	R9		Washing 2	Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	6	a					
IPL	R11		Water pipe 2	Ø	Ø	+MA	+MA	<i>L.ivanovii</i>		+	-	Ø	Ø	/	-	ND	G	6	a					
IPL	R19		Water of wash lines spinach	Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	6	a					
IPL	S10		Process water delicatessen retail	Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	6	a					
IPL	S13		Process water	Ø	Ø	Ø	Ø	/		-	-	Ø	Ø	/	-	NA	G	6	a					
IPL	S14		Process water	Ø	Ø	Ø	Ø	/		-	+	Ø	Fraser : -LE	/	-	PPNA	G	6						
IPL	U5		Process water delicatessen retail	Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	6	a					
IPL	U6		Rinsing water	Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	6	a					
IPL	V1		Water cheese dairy	Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	6	a					
IPL	V2		Process water	Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	6	a					
IPL	V3		Salted water	Ø	Ø	Ø	Ø	/		-	+	+MA	+MA	<i>L.ivanovii</i>	+	PD	G	6	a					
IPL	W1		Rinsing water	-LA	+MA	-MA	+MA	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	6	a					
IPL	W2		Rinsing water	-MA	+MA	-MA	+MA	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	6	a					
IPL	W3		Rinsing water	-MA	+MA	-MA	+MA	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	6	a					
IPL	W4		Clean	-MA	+MA	-MA	+MA	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	6	a					
IPL	W19		Rinsing water	Ø	-LE	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	6	a					
IPL	W20		Process water pastry retail	Ø	Ø	-ME	-LE	/		-	+	-MB	-HB	<i>L.innocua</i>	+	PD	G	6	a					
IPL	X1		Process water fish retail	+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	6	a					
IPL	X2		Water process	+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>		+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	6	a					
IPL	X5		Water process	+MA	+MA	+MB	+MB	<i>L.monocytogenes</i>		+	+	+LB	+MA	<i>L.monocytogenes</i>	+	PA	G	6	a					
ADRIA	4879	Eau de process (Industrie de poissons)	Process water (Fish industry)	st	st	st	st	/		-	-	st	st	/	-	NA	G	6	a					
ADRIA	4880	Eau de process (Industrie de poissons)	Process water (Fish industry)	st	st	st	st	/		-	-	st	st	/	-	NA	G	6	a					
ADRIA	4881	Eau de process (Industrie de poissons)	Process water (Fish industry)	st	st	st	st	/		-	-	st	st	/	-	NA	G	6	a					
ADRIA	4882	Eau de process (Industrie de poissons)	Process water (Fish industry)	st	st	st	st	/		-	-	st	st	/	-	NA	G	6	a					
ADRIA	4883	Eau mélange poissons (Industrie de poissons)	Rinse water (Fish industry)	st	st	-	-	/		-	-	-	-	/	-	NA	G	6	a					
IPL	K16		Sponge cheese dairy	-LE	-ME	-LE	-ME	/		-	-	-ME	/	/	-	NA	G	6	b					
IPL	K21		Surface tank	+MB	+MB	+MB	+MB	<i>L.innocua</i> <i>L.monocytogenes</i>		+	+	+HB	+HB	<i>L.innocua</i> <i>L.monocytogenes</i>	+	PA	G	6	b					
IPL	Q2		Surface	+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	6	b					
IPL	T20		Meat tray	Ø	Ø	-LE	-LE	/		-	-	-ME	/	/	-	NA	G	6	b					
IPL	U17		Stainless steel preparation room	+LB	+LB	+MA	+MB	<i>L.monocytogenes</i>		+	+	+MD	+LB	<i>L.monocytogenes</i>	+	PA	G	6	b					
IPL	U18		Cutting table for fish	-LE	-LE	-LE	-LE	/		-	-	-ME	/	/	-	NA	G	6	b					
IPL	U19		Stainless steel preparation room for vegetables	Ø	Ø	-ME	-ME	/		-	-	-ME	/	/	-	NA	G	6	b					
IPL	W18		Surface pastry retail	-LE	-LE	-LE	-LE	/		-	-	-ME	/	/	-	NA	G	6	b					

Laboratory	Sample N°	Products (French name)	Products	ENVIRONMENTAL SAMPLES													Agreement	Protocol	Category	Type							
				Reference method: ISO 11290-1/A1				Alternative method: BAX® System PCR Assay for Genus Listeria 24E				CONFIRMATION 24 LEB			FINAL RESULT												
				FRASER 1/2		FRASER		CONFIRMATION		RESULT Listeria spp	BAX® Listeria	Brilliance 24h (IPL)	Brilliance 48h (IPL)	IDENTIFICATION													
				O&A1	P1	O&A2	P2	IDENTIFICATION				O&A 24h (Adria)	O&A 48h (Adria)														
IPL	X3	Sponge		+LA	+MB	+MB	+MB	<i>L.monocytogenes</i>		+	+	+LB	+MB	<i>L.monocytogenes</i>	+	PA	G	6	b								
IPL	Y4	Surface pastry retail		-MA	+MA	-MA	+MD	<i>L.welshimeri</i>		+	-	-ME	-LE Fraser : -LE	/	-	ND	G	6	b								
IPL	Y12	Surface of delicatessen knife		-LA	-LA	-MA	+MB	<i>L.welshimeri</i>		+	+	-MA	-MB	<i>L.welshimeri</i>	+	PA	G	6	b								
IPL	Z3	Cutting table for fish		-LE	-LE	-LE	-ME	/		-	-	-LE	/	/	-	NA	G	6	b								
IPL	Z13	Floor in packaging room		-ME	-ME	-ME	-ME	/		-	-	-ME	/	/	-	NA	G	6	b								
ADRIA	5763	Chiffonnette caniveau haut filetage (Industrie poissons)	Surface sample (fish industry)	st	st	st	st	/		-	-	st	st	/	-	NA	G	6	b								
ADRIA	5764	Chiffonnette tapis parage avant nettoyage (Industrie poissons)	Surface sample (fish industry)	st	st	st	st	/		-	-	st	st	/	-	NA	G	6	b								
ADRIA	5765	Chiffonnette tapis déchets peleuse (Industrie poissons)	Surface sample (fish industry)	st	st	st	-	/		-	-	st	st	/	-	NA	G	6	b								
ADRIA	5766	Chiffonnette sol haut filetage (Industrie poissons)	Surface sample (fish industry)	H+/H-	+	H+/H-	+	<i>L.monocytogenes</i> / <i>L.welshimeri</i>		+	+	H+	H+ / H-	<i>L.monocytogenes</i> (RLM) / <i>L.welshimeri</i>	+	PA	G	6	b								
ADRIA	5767	Chiffonnette sol couloir co-produits (Industrie poissons)	Surface sample (fish industry)	H-	+	H-	+	<i>L.welshimeri</i>		+	+	H-	H-	<i>L.innocua</i>	+	PA	G	6	b								
ADRIA	5768	Chiffonnette pousoir réception matières premières (Industrie poissons)	Surface sample (fish industry)	st	st	-	-	/		-	-	st	-	/	-	NA	G	6	b								
ADRIA	5950	Chiffonnette table	Surface sample	H+ (7)	+	H+	+	<i>L.monocytogenes</i>		+	+	H+	H+	<i>L.monocytogenes</i>	+	PA	G	6	b								
IPL	N7	Scraps from filter machine		Ø	Ø	-LE	-LE	/		-	-	-ME	/	/	-	NA	G	6	c								
IPL	S11	Scraps from cheese dairy		+LA(2)	+LA(8)	+MA	+MA	<i>L.monocytogenes</i>		+	-	-LE	-LE Fraser : -LE	/	-	ND	G	6	c								
IPL	T1	Scraps from cheese dairy		Ø	Ø	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	6	c								
IPL	T2	Vegetables scraps		Ø	-LE	Ø	-LE	/		-	-	Ø	/	/	-	NA	G	6	c								
IPL	T11	Scraps from fish retail outlet		Ø	-LE	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	6	c								
IPL	T13	Cheese scraps		Ø	Ø	Ø	Ø	/		-	-	Ø	/	/	-	NA	G	6	c								
IPL	U4	Scraps from delicatessen retail outlet		Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	6	c								
IPL	U7	Scraps from delicatessen retail outlet		+LA	+LA	+MA	+MA	<i>L.monocytogenes</i>		+	+	+LA	+MA	<i>L.monocytogenes</i>	+	PA	G	6	c								
IPL	U8	Scraps from packaging line		Ø	Ø	Ø	Ø	/		-	-	-ME	-ME	/	-	NA	G	6	c								
IPL	U9	Scraps from meat cutting table		+LA(5)	+LB	+MB	+MB	<i>L.monocytogenes</i> <i>L.welshimeri</i>		+	+	+MB	+MB	<i>L.monocytogenes</i> <i>L.welshimeri</i>	+	PA	G	6	c								
IPL	U14	Scraps from fish retail outlet		-LE	-LE	-LE	-LE	/		-	-	-ME	-ME	/	-	NA	G	6	c								
IPL	U15	Scraps		Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	6	c								
IPL	U16	Scraps production line		Ø	Ø	Ø	Ø	/		-	+	-ME	-ME / FR Ø	/	-	PPNA	G	6	c								
IPL	U20	Scraps from butchery cutting table		-MA	+MB	-MA	+MB	<i>L.welshimeri</i>		+	+	-HB	-MB	<i>L.welshimeri</i>	+	PA	G	6	c								

Laboratory	Sample N°	Products (French name)	Products	ENVIRONMENTAL SAMPLES												FINAL RESULT	Agreement	Protocol	Category	Type				
				Reference method: ISO 11290-1/A1						Alternative method: BAX® System PCR Assay for Genus Listeria 24E														
				FRASER 1/2		FRASER		CONFIRMATION		RESULT Listeria spp	BAX® Listeria	CONFIRMATION 24 LEB			IDENTIFICATION									
				O&A1	P1	O&A2	P2	IDENTIFICATION				Brilliance 24h (IPL)	Brilliance 48h (IPL)											
IPL	V4		Scraps from cheese production "parmigiano"	Ø	Ø	-MA	+MB	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	6	c					
IPL	V5		Scraps from spices line	-LE	-LE	-LE	-LE	/		-	-	Ø	/	/	-	NA	G	6	c					
IPL	W10		Scraps from packaging line	+LA	+LA	+MB	+MB	<i>L.ivanovii</i>		+	+	+MB	+LB	<i>L.ivanovii</i>	+	PA	G	6	c					
IPL	W11		Scraps from storage tanks	-LA	+LA	-MA	+MA	<i>L.innocua</i>		+	+	-MA	-MA	<i>L.innocua</i>	+	PA	G	6	c					
IPL	W12		Scraps from rinsing tanks	-LE	Ø	-LE	-LE	/		-	-	-LE	/	/	-	NA	G	6	c					
IPL	W13		Scraps from delicatessen retail outlet	Ø	Ø	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	6	c					
IPL	W15		Scraps from pastry fabrication line	Ø	Ø	Ø	Ø	/		-	-	-LE	/	/	-	NA	G	6	c					
IPL	W16		Scraps from delicatessen retail outlet	-LE	+LD	-MB	+MB	<i>L.welshimeri</i>		+	+	-MB	-LB	<i>L.welshimeri</i>	+	PA	G	6	c					
IPL	W17		Scraps from pastry retail outlet	-LE	-LE	-LE	-LE	/		-	+	-MA	-MA	<i>L.innocua</i>	+	PD	G	6	c					
IPL	X4		Scraps from fish retail outlet	+LA	+MA	+MA	+MB	<i>L.monocytogenes</i>		+	+	+MA	+MA	<i>L.monocytogenes</i>	+	PA	G	6	c					
IPL	Y5		Vegetables scraps	-MA	+MA	-MA	+MB	<i>L.innocua</i>		+	-	-ME	-LE Fraser : -LE	/	-	ND	G	6	c					
IPL	Y7		Fish scraps	-LE	-LE	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	6	c					
IPL	Y9		Scraps from fish retail outlet	Ø	Ø	Ø	Ø	/		-	-	-ME	/	/	-	NA	G	6	c					
IPL	Y11		Scraps from butchery retail outlet	-LA	+LA	-MA	+MA	<i>L.welshimeri</i>		+	+	-MB	-MB	<i>L.innocua</i>	+	PA	G	6	c					
IPL	Y13		Scraps from delicatessen preparation line	+LA	+LB	+MA	+MA	<i>L.monocytogenes</i>		+	-	-LE	-LE Fraser : -LE	/	-	ND	G	6	c					
IPL	Y14		Scraps from mincer	Ø	Ø	Ø	Ø	/		-	+	+MB	+MB	<i>L.monocytogenes</i>	+	PD	G	6	c					
IPL	Y15		Scraps from cream production line	+LB	+LB	+MB	+MB	<i>L.monocytogenes</i>		+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	6	c					
IPL	Z1		Scraps from delicatessen retail outlet	-MA	+MA	-MA	+MB	<i>L.monocytogenes</i> <i>L.welshimeri</i>		+	+	+MB	-MA	<i>L.monocytogenes</i> <i>L.welshimeri</i>	+	PA	G	6	c					
IPL	Z2		Scraps from mincer	Ø	-LE	-LE	-LE	/		-	+	+MB	+MD	<i>L.monocytogenes</i>	+	PD	G	6	c					
IPL	Z4		Scraps production line	+MB	+MB	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>		+	+	+MB	+MB	<i>L.monocytogenes</i> <i>L.innocua</i>	+	PA	G	6	c					
IPL	Z5		Scraps from packaging line	+MB	+MA	+MA	+MD	<i>L.monocytogenes</i>		+	-	+LD	+LD	<i>L.monocytogenes</i>	-	ND	G	6	c					
IPL	Z6		Scraps from transferring line	-LE	-LE	-ME	-ME	/		-	-	-LE	/	/	-	NA	G	6	c					
IPL	Z7		Scraps from cutting line	+LA(4)	+LD(2)	+MA	+MA	<i>L.monocytogenes</i>		+	+	+MB	+MB	<i>L.monocytogenes</i>	+	PA	G	6	c					

**Appendix 5 - Relative level of detection: raw data****IPL Legend****Total bacteria growth**

Ø : no growth

L = low

M = medium

H = high

**Distribution of flora**

A = pure culture of suspicious colonies

B = mix with a majority of suspicious colonies

C = mix with a minority of suspicious colonies

D = mix with rare suspicious colonies

E = absence of suspicious colonies

**ADRIA Legend**

H-: characteristic *Listeria* colonies without halo

H+: characteristic *Listeria* colonies with halo

-: no typical colonies but presence of background microflora

st: plate without any colony

Study performed by IPL in 2008

Matrix : Raw milk

Strain: *Listeria ivanovii* L150

Aerobic mesophilic flora: 4 000 CFU/ml

Level	Inoculation level (b/25 g)	Reference method: ISO 11290-1/A1						Alternative method: BAX "Automate" Genus <i>Listeria</i>			
		Fraser 1/2 (10 µl)		Fraser		Result	Conclusion	BAX L.spp test	Streaking BAX 24 LEB onto OCLA	Final result	Conclusion
		O&A1	P1	O&A2	P2						
1	0	Ø	Ø	Ø	Ø	-	0/6	-	/	-	0/6
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
2	0,32	Ø	Ø	Ø	Ø	-	1/6	-	/	-	1/6
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		+	Ø / +MA (RLM)	+	
		Ø	Ø	+MA	+MA	+		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
3	0,64	+LA	+LA(1)	+MA	+MA	+	3/6	+	Ø / +MA (RLM)	+	3/6
		Ø	Ø	Ø	Ø	-		-	/	-	
		+LA(2)	Ø	+MA	+MA	+		-	/	-	
		+LA(5)	+LA(3)	+MA	+MA	+		+	Ø / +MA (RLM)	+	
		Ø	Ø	Ø	Ø	-		+	Ø / +MA (RLM)	+	
		Ø	Ø	Ø	Ø	-		-	/	-	
4	1,28	+LA	+LA(1)	+MA	+MA	+	4/6	-	/	-	4/6
		Ø	Ø	Ø	Ø	-		+	Ø / +MA (RLM)	+	
		Ø	Ø	Ø	Ø	-		-	/	-	
		+LA(1)	Ø	+MA	+MA	+		+	Ø / +MA (RLM)	+	
		+LA(4)	Ø	+MA	+MA	+		+	Ø / +MA (RLM)	+	
		+LA(1)	Ø	+MA	+MA	+		+	Ø / +LA (5b) / +MA (RLM)	+	
5	2,24	Ø	Ø	+MA	+MA	+	4/6	+	Ø / +MA (RLM)	+	5/6
		Ø	Ø	+MA	+MA	+		+	Ø / +LA (5b) / +MA (RLM)	+	
		+LA(4)	Ø	+MA	+MA	+		+	Ø / +MA (RLM)	+	
		+LA	Ø	+MA	+MA	+		-	/	-	
		Ø	Ø	Ø	Ø	-		+	Ø / +LA (5b) / +MA (RLM)	+	
		Ø	Ø	Ø	Ø	-		+	Ø / +MA (RLM)	+	
6	3,68	Ø	Ø	+MA	+MA	+	6/6	+	Ø / +MA (RLM)	+	6/6
		+LA(1)	Ø	+MA	+MA	+		+	Ø / +MA (RLM)	+	
		+LA(1)	Ø	+MA	+MA	+		+	Ø / +MA (RLM)	+	
		+LA(1)	Ø	+MA	+MA	+		+	Ø / +MA (RLM)	+	
		+LA(4)	+LA(1)	+MA	+MA	+		+	Ø / +MA (RLM)	+	
		+LA(4)	+LA(4)	+MA	+MA	+		+	Ø / +MA (RLM)	+	

**Matrix: Raw vegetable mix**Strain: *Listeria monocytogenes* 4b L58

Aerobic mesophilic flora: 420 000 CFU/g

Level	Inoculation level (b/25 g)	Reference method: ISO 11290-1/A1						Alternative method: BAX "Automate" Genus <i>Listeria</i>			
		Fraser 1/2 (10 µl)		Fraser		Result	Conclusion	BAX L.spp test	Streaking BAX 24 LEB onto OCLA	Final result	Conclusion
		O&A1	P1	O&A2	P2			-	-		
1	0	Ø	Ø	Ø	Ø	-	0/6	-	/	-	0/6
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
2	0,38	Ø	Ø	Ø	Ø	-	0/6	-	/	-	1/6
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		+	+MA	+	
3	0,60	Ø	Ø	Ø	Ø	-	2/6	+	+MB	+	3/6
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	-LE	-LE	-LE	-		+	+MB	+	
		Ø	Ø	Ø	Ø	-		-	/	-	
		+LA	+LA	+MA	+MA	+		-	/	-	
		+LA	+LA	+MA	+MA	+		+	+MB	+	
4*	1,34	Ø	Ø	+MA	+MA	+	5/6	+	+MA	+	4/6
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		-	/	-	
		Ø	Ø	Ø	Ø	-		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		-	/	-	
5*	2,68	+LA	+LA	+MA	+MA	+	6/6	+	+MA	+	6/6
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	

**Matrix : Process water**Strain: *Listeria innocua* L175

Aerobic mesophilic flora: 300 000 CFU/m and \* 4 500 CFU/ml

Level	Inoculation level (b/25 g)	Reference method: ISO 11290-1/A1						Alternative method: BAX "Automate" Genus <i>Listeria</i>			
		Fraser 1/2 (10µl)		Fraser		Result	Conclusion	BAX L.spp test	Streaking BAX 24 LEB onto OCLA	Final result	Conclusion
		O&A1	P1	O&A2	P2						
1	0,00	Ø	Ø	Ø	Ø	-	0/6	-	/	-	0/6
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
2	0,49	Ø	+LA	+MB	+MB	+	2/6	+	+MA	+	2/6
		Ø	Ø	Ø	Ø	-		+	+MA	+	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	+LA	+LA	+		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
3*	0,85	Ø	Ø	Ø	Ø	-	3/6	+	-MA	+	3/6
		Ø	Ø	+LA	+MA	+		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		+LA	Ø	+MA	+MA	+		-	/	-	
		Ø	Ø	Ø	Ø	-		+	-MA	+	
		+LA	+LA	+MA	+MA	+		+	-MA	+	
4*	2,56	+LA	+LA	+MA	+MA	+	5/6	+	-MA	+	6/6
		+LA	+LA	+MA	+MA	+		+	-MA	+	
		+LA	+LA	+MA	+MA	+		+	-MA	+	
		Ø	Ø	Ø	Ø	-		+	-MA	+	
		+LA	+LA	+MA	+MA	+		+	-MA	+	
		+LA	+LA	+MA	+MA	+		+	-MA	+	
5	4,86	+LB	+LB	+MA	+MA	+	6/6	+	+MA	+	6/6
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+LB	+MA	+MA	+		+	+MA	+	
		+LB	+LA	+MA	+MA	+		+	+MB	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+MA	+MA	+MA	+		+	+MB	+	

**Matrix : Rillettes**Strain: *Listeria monocytogenes* L49

Aerobic mesophilic flora: 4 000 000 CFU/g

Sample	Inoculation level (b/25 g)	Reference method: ISO 11290-1/A1						Alternative method: BAX "Automate" Genus <i>Listeria</i>			
		Fraser 1/2 (10 µl)		Fraser		Result	Conclusion	BAX L.spp test	Streaking BAX 24 LEB onto <i>Brilliance Listeria</i>	Final result	Conclusion
		O&A 1	P1	O&A2	P2						
1	0	Ø	Ø	Ø	Ø	-	0/6	-	/	-	0/6
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	-LE	-LE	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
		Ø	Ø	Ø	Ø	-		-	/	-	
2	0.35	-LE	-LE	-LE	-ME	-	1/6	-	-ME	-	1/6
		+MA	+MA	+MA	+HA	+		-	-ME	-	
		-LE	-LE	-ME	-ME	-		-	-ME	-	
		-LE	-LE	-ME	-ME	-		+	+MA	+	
		-LE	-ME	-ME	-ME	-		-	-ME	-	
		-LE	-ME	-ME	-ME	-		-	-ME	-	
3	0.62	Ø	Ø	Ø	Ø	-	4/6	-	/	-	3/6
		Ø	Ø	Ø	Ø	-		+	+MA	+	
		+LA	+LA	+MA	+MA	+		-	/	-	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		-	/	-	
4	1.06	Ø	Ø	Ø	Ø	-	3/6	+	+MA	+	4/6
		+LA	+LA	+MA	+MA	+		-	/	-	
		Ø	Ø	Ø	Ø	-		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		Ø	Ø	Ø	Ø	-		-	/	-	
5	1.58	+MA	+MA	+MA	+MA	+	6/6	+	+MA	+	6/6
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	

**Matrix : Smoked salmon**Strain: *Listeria monocytogenes* 1/2a L5

Aerobic mesophilic flora: 4 000 000 CFU/g

Level	Inoculation level (b/25 g)	Reference method: ISO 11290-1/A1						Alternative method: BAX "Automate" Genus <i>Listeria</i>			
		Fraser 1/2 (10 µl)		Fraser		Result	Conclusion	BAX L.spp test	Streaking BAX 24 LEB onto Brilliance <i>Listeria</i>	Final result	Conclusion
		O&A 1	P1	O&A2	P2			-	-		
1	0	Ø	Ø	Ø	Ø	-	0/6	-	-LE	-	0/6
		Ø	Ø	Ø	Ø	-		-	-LE	-	
		Ø	Ø	Ø	Ø	-		-	-ME	-	
		Ø	Ø	Ø	Ø	-		-	-LE	-	
		Ø	Ø	Ø	Ø	-		-	Ø	-	
		Ø	Ø	Ø	Ø	-		-	-LE	-	
2	0.28	Ø	Ø	Ø	Ø	-	3/6	+	+MA	+	1/6
		Ø	Ø	Ø	Ø	-		-	-ME	-	
		Ø	Ø	Ø	Ø	-		-	-ME	-	
		+MA	+MA	+MA	+MA	+		-	Ø	-	
		+MA	+MA	+MA	+MA	+		-	Ø	-	
		+MA	+MA	+MA	+MA	+		-	-ME	-	
3	0.55	Ø	Ø	Ø	Ø	-	2/6	+	+MA	+	5/6
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		Ø	Ø	Ø	Ø	-		+	+MA	+	
		Ø	Ø	Ø	Ø	-		+	+MA	+	
		Ø	Ø	Ø	Ø	-		-	-ME	-	
4	1.11	+MA	+MA	+MA	+MA	+	5/6	+	+MA	+	6/6
		+LA	+MA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
		Ø	Ø	Ø	Ø	-		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+LA	+LA	+MA	+MA	+		+	+MA	+	
5	2.21	+MA	+MA	+MA	+MA	+	6/6	+	+MA	+	6/6
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	
		+MA	+MA	+MA	+MA	+		+	+MA	+	

Study performed by ADRIA in 2017

Matrix : Deli-salad

Strain : *Listeria seeligeri* Ad1293Aerobic mesophilic flora:  $3 \cdot 10^3$  CFU/g

N° sample	Level	Inoculation level (cfu/sample)	Reference method : ISO 11290-1/A1*					BAX® System PCR Assay for Genus Listeria 24E 24LEB for 24h at 37°C						
			Half Fraser		Fraser		Final Result	Number positive samples/Total	PCR result	Confirmation			Final result	Number positive samples/Total
			O&A	Palcam	O&A	Palcam				O&A	Palcam	Identification		
5566	0	0	st	-	-	-	-	0/5	-	-	-	/	-	0/5
5567			st	-	-	-	-		-	-	-	/	-	
5568			st	st	st	st	-		-	-	-	/	-	
5569			st	st	-	st	-		-	-	-	/	-	
5570			st	st	st	-	-		-	-	-	/	-	
5571	Low	0.7	H-	+	H-	+	+	14/20	+	H-	+	+	+	9/20
5572			H-	+	H-	+	+		+	H-	+	+	+	
5573			st	st	st	st	-		-	-	-	/	-	
5574			H-	+	H-	+	+		+	H-	+	+	+	
5575			st	-	-	st	-		-	-	-	/	-	
5576			H-	+	H-	+	+		+	H-	+	+	+	
5577			H-	+	H-	+	+		+	H-	+	+	+	
5578			H-	+	H-	+	+		-	-	-	/	-	
5579			H-	+	H-	+	+		+	H-	+	+	+	
5580			st	-	st	-	-		-	-	-	/	-	
5581			-	st	st	st	-		-	-	-	/	-	9/20
5582			H-	+	H-	+	+		-	-	-	/	-	
5583			H-	+	H-	+	+		+	H-	+	+	+	
5584			H-	+	H-	+	+		-	-	-	/	-	
5585			H-	+	H-	+	+		+	H-	+	+	+	
5586	High	2.1	-	st	-	-	-	5/5	-	-	-	/	-	4/5
5587			-	-	st	st	-		+	H-	+	+	+	
5588			H-	+	H-	+	+		-	-	-	/	-	
5589			H-	+	H-	+	+		-	-	-	/	-	
5590			H-	+	H-	+	+		-	-	-	/	-	
5591			H-	+	H-	+	+		+	H-	+	+	+	
5592			H-	+	H-	+	+		-	-	-	/	-	
5593			H-	+	H-	+	+		+	H-	+	+	+	
5594			H-	+	H-	+	+		+	H-	+	+	+	
5595			H-	+	H-	+	+		+	H-	+	+	+	

\* Analyses performed according to the COFRAC accreditation (Accreditation Testing n°1-0144, scope available on [www.cofrac.fr](http://www.cofrac.fr))

## Appendix 6 – Inclusivity and exclusivity study: raw data

INCLUSIVITY (Study realized by IPL, 2008)					
Reference	Strain	Origin	Inoculation level in 225 mL 24 LEB broth	BAX® Genus <i>Listeria</i> 24E Result	Brilliance <i>Listeria</i>
L5	<i>Listeria monocytogenes</i> 1/2a	Pieces of smoked salmon	8,0	+	+MA
L9	<i>Listeria monocytogenes</i> 1/2a	Munster cheese (rind)	8,0	+	+MA
L11	<i>Listeria monocytogenes</i> 1/2a	Munster cheese (rind)	7,0	+	+MA
L12	<i>Listeria monocytogenes</i> 1/2a	Smoked salmon	8,0	+	+HA
L14	<i>Listeria monocytogenes</i> 1/2b	Pork ear	8,0	+	+MA
L16	<i>Listeria monocytogenes</i> 1/2c	Ground meat	6,0	+	+MA
L17	<i>Listeria monocytogenes</i> 1/2c	Bacon	3,0	+	+MA
L18	<i>Listeria monocytogenes</i> 1/2c	Munster cheese (rind)	8,0	+	+MA
L25	<i>Listeria monocytogenes</i> 1/2	Smoked salmon	10,0	+	+MA
L28	<i>Listeria monocytogenes</i> 1/2c	Environment sample	5,0	+	+MA
L37	<i>Listeria monocytogenes</i> 1/2b	Maroille cheese	4,0	+	+HA
L39	<i>Listeria monocytogenes</i>	Saucisson	5,0	+	+MA
L43	<i>Listeria monocytogenes</i> 1/2a	Ground meat	9,0	+	+HA
L44	<i>Listeria monocytogenes</i> 1/2a	Saucisson	5,0	+	+HA
L47	<i>Listeria monocytogenes</i> 1/2a	Browed potatoes	5,0	+	+MA
L51	<i>Listeria monocytogenes</i> 1/2b	Germain cheese	10,0	+	+LA
L57	<i>Listeria monocytogenes</i> 4a	ATCC 19114	2,0	+	+MA
L58	<i>Listeria monocytogenes</i> 4b	Salad	4,0	+	+LA
L61	<i>Listeria monocytogenes</i> 4e	ATCC 19118	5,0	+	+MA
L62	<i>Listeria monocytogenes</i> 4e	Reblochon cheese	4,0	+	+HA
L70	<i>Listeria monocytogenes</i>	Salmon from Ireland	4,0	+	+MA
L116	<i>Listeria monocytogenes</i> 1/2a	Fish meal	5,0	+	+HA
L117	<i>Listeria monocytogenes</i> 1/2c	Montbeliard sausage	8,0	+	+MA
L119	<i>Listeria monocytogenes</i>	Spinaches	7,0	+	+MA
L121	<i>Listeria monocytogenes</i>	Neufchatel cheese	5,0	+	+MA
L123	<i>Listeria monocytogenes</i>	Mozzarella cheese	5,0	+	+MA
L124	<i>Listeria monocytogenes</i>	Perch fillet	6,0	+	+MA
L125	<i>Listeria monocytogenes</i>	Vegetables pan fry	6,0	+	+MA
L128	<i>Listeria monocytogenes</i> 1/2a	Soya cattle cake	4,0	+	+MA
L129	<i>Listeria monocytogenes</i> 1/2a	Browed potatoes	2,0	+	+MA
L137	<i>Listeria monocytogenes</i>	Ground meat	2,0	+	+MA
L141	<i>Listeria monocytogenes</i>	Environmental sample	4,0	+	+MA
L149	<i>Listeria monocytogenes</i>	Environmental sample	6,0	+	+MA
L152	<i>Listeria monocytogenes</i>	Environmental sample	5,0	+	+LA
L156	<i>Listeria monocytogenes</i>	French pies	2,0	+	+MA
L176	<i>Listeria monocytogenes</i>	Beef meat	9,0	+	+MA
L140	<i>Listeria seeligeri</i>	Frozen French pies	6,0	+	+LA
L142	<i>Listeria seeligeri</i>	Raw milk cheese	6,0	+	+LA
L 148	<i>Listeria seeligeri</i>	Environment	8,0	-	BAX LEB 24 : Ø BN : +LA
			44,0	-	
			1,0E+05	+	
L115	<i>Listeria seeligeri</i>	Waste water	12,0	+	+MA

INCLUSIVITY (Study realized by IPL, 2008)					
Reference	Strain	Origin	Inoculation level in 225 mL 24 LEB broth	BAX® Genus <i>Listeria</i> 24E Result	Brilliance <i>Listeria</i>
L83	<i>Listeria seeligeri</i> 1/2b	Beef tongue	9,0	+	+MA
L84	<i>Listeria seeligeri</i> 1/2b	Ground meat	8,0	+	+LA
L80	<i>Listeria ivanovii</i>	Collection	9,0	+	+LA
L 154	<i>Listeria ivanovii</i>	Sausage with herbs	9,0	+	+MA
L 157	<i>Listeria ivanovii</i> spp. <i>ivanovii</i>	Collection	9,0	+	+LA
L 158	<i>Listeria ivanovii</i>	NSB 79332	6,0	+	+LA
L 159	<i>Listeria ivanovii</i> spp. <i>ivanovii</i>	Collection	8,0	+	+MA
L 173	<i>Listeria ivanovii</i>	Collection	8,0	+	+MA
L179	<i>Listeria ivanovii</i>	Environment	3,0	+	+LA
L170	<i>Listeria ivanovii</i>	Collection	9,0	+	+MA
L185	<i>Listeria ivanovii</i>	Net	8,0	+	+MA
L153	<i>Listeria ivanovii</i>	Environmental sample	15,0	+	+MA
L133	<i>Listeria ivanovii</i>	Roquefort cheese	4,0	+	+MA
L151	<i>Listeria ivanovii</i>	Ground meat	1,0	+	+LA
L172	<i>Listeria ivanovii</i>	ATCC 19119	9,0	+	+LA
L150	<i>Listeria ivanovii</i>	Dairy product	4,0	+	+LA
L1	<i>Listeria innocua</i> 6a	Collection ATCC 33090	7,0	+	+MA
L64	<i>Listeria innocua</i>	Epoisses cheese	5,0	+	+MA
L72	<i>Listeria innocua</i>	Boulettes d'Avesnes cheese	7,0	+	+MA
L76	<i>Listeria innocua</i> 6b	Ground meat	10,0	+	+MA
L77	<i>Listeria innocua</i> 6a	Toulouse sausage	4,0	+	+MA
L108	<i>Listeria innocua</i>	Gorgonzola cheese	7,0	+	+MA
L90	<i>Listeria welshimeri</i>	Delicatessen	1,0	+	+LA
L91	<i>Listeria welshimeri</i>	Pure pork slicing sausage	5,0	+	+MA
L99	<i>Listeria welshimeri</i>	Sausages	3,0	+	+LA
L101	<i>Listeria welshimeri</i>	Ham	9,0	+	+MA
L 155	<i>Listeria welshimeri</i>	Salmon fillet	8,0	+	+MA

EXCLUSIVITY (Study realized by IPL, 2008)				
Reference	Strain	Origin	Inoculation level in 10 mL non-selective nutrient broth	BAX® Genus <i>Listeria</i> 24E Result
BA2	<i>Bacillus cereus</i>	Beet	1.0E+05	-
BA4	<i>Bacillus stearothermophilus</i>	Dairy product	1.5E+05	-
BA23	<i>Bacillus sphaericus</i>	Collection	1.0E+05	-
BA5	<i>Bacillus sphaericus</i>	Meat product	1.0E+05	-
BA5	<i>Bacillus sphaericus</i>	Meat product	2.0E+05	-
BA7	<i>Bacillus coagulans</i>	Collection	5.0E+05	-
BA19	<i>Bacillus cereus</i>	Environment	3.0E+05	-
BA9	<i>Bacillus cereus</i>	Potatoes flakes	2.0E+05	-
BA14	<i>Bacillus cereus</i>	Egg	3.0E+05	-
BA15	<i>Bacillus cereus</i>	Custard	6.0E+05	-
BA21	<i>Bacillus cereus</i>	Tabbouleh	2.0E+05	-
E1	<i>Enterococcus faecalis</i>	Egg product	4.0E+05	-
E2	<i>Enterococcus faecium</i>	Collection ATCC 3286	1.7E+05	-
E3	<i>Streptococcus bovis</i>	Collection	1.5E+05	-
E13	<i>Streptococcus bovis</i>	Collection CIP 5623	1.5E+05	-
E8	<i>Enterococcus durans</i>	Meat product	6.0E+05	-
E9	<i>Enterococcus faecium</i>	Tarama	7.0E+05	-
E17	<i>Streptococcus equinus</i>	Collection	1.0E+05	-
E6	<i>Enterococcus faecalis</i>	Collection ATCC 19433	3.0E+04	-
E7	<i>Enterococcus faecium</i>	Collection CIP 5433	3.0E+05	-
E14	<i>Streptococcus anginosus</i>	Collection	1.5E+05	-
32	<i>Rhodococcus equi</i>	Meat product	3.0E+05	-
33	<i>Lactobacillus casei</i>	Dairy product	2.0E+03	-
L139	<i>Jonesia denitrificans</i>	Collection	3.0E+05	-
Le1	<i>Rhodotorula rubra</i>	Pastry	2.0E+04	-
Le3	<i>Candida albicans</i>	Collection	1.0E+05	-
ST26	<i>Staphylococcus intermedius</i>	Collection	3.0E+05	-
ST17	<i>Staphylococcus aureus</i>	Iced yoghurt	5.0E+05	-
ST3	<i>Staphylococcus epidermidis</i>	Yoghurt	4.0E+05	-
M1	<i>Micrococcus</i>	Environment	2.0E+05	-
L 147	<i>Listeria grayi</i>	ATCC 25 401	56.0	-

**Appendix 7 - Results obtained by the collaborators (study realized by IPL, 2008)****Laboratory A**

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		

Total flora of milk (UFC/ml): &lt;1

**Laboratory B**

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		

Total flora of milk (UFC/ml): NC

Laboratory C

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		
Total flora of milk (UFC/ml):				1700								

Laboratory D

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		
Total flora of milk (UFC/ml):				1200								

Laboratory E

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agreement		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		
Total flora of milk (UFC/ml):		<4										

Laboratory F

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agreement		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		
Total flora of milk (UFC/ml):		3600										

Laboratory G

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		

Total flora of milk (UFC/ml): 1300

Laboratory H

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		

Total flora of milk (UFC/ml): 5,8.10<sup>6</sup>

Laboratory K

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	-	+	-	ND		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		

Total flora of milk (UFC/ml): 54000

Laboratory L

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		

Total flora of milk (UFC/ml): 3,8.10<sup>4</sup>

Laboratory M

Sample N°	Reference method				Result	Comparison / expected results	Alternative method BAX® L. spp			Agree-ment		
	Fraser 1/2		Fraser				Test BAX® L. spp	Confirmation	Result			
	O&A	PALCAM	O&A	PALCAM								
3	-	-	-	-	-	=	-	-	-	NA		
4	-	-	-	-	-	=	-	-	-	NA		
9	-	-	-	-	-	=	-	-	-	NA		
10	-	-	-	-	-	=	-	-	-	NA		
11	-	-	-	-	-	=	-	-	-	NA		
12	-	-	-	-	-	=	-	-	-	NA		
19	-	-	-	-	-	=	-	-	-	NA		
20	-	-	-	-	-	=	-	-	-	NA		
1	+	+	+	+	+	=	+	+	+	PA		
2	+	+	+	+	+	=	+	+	+	PA		
5	+	+	+	+	+	=	+	+	+	PA		
6	+	+	+	+	+	=	+	+	+	PA		
13	+	+	+	+	+	=	+	+	+	PA		
14	+	+	+	+	+	=	+	+	+	PA		
21	+	+	+	+	+	=	+	+	+	PA		
22	+	+	+	+	+	=	+	+	+	PA		
7	+	+	+	+	+	=	+	+	+	PA		
8	+	+	+	+	+	=	+	+	+	PA		
15	+	+	+	+	+	=	+	+	+	PA		
16	+	+	+	+	+	=	+	+	+	PA		
17	+	+	+	+	+	=	+	+	+	PA		
18	+	+	+	+	+	=	+	+	+	PA		
23	+	+	+	+	+	=	+	+	+	PA		
24	+	+	+	+	+	=	+	+	+	PA		
Total flora of milk (UFC/ml): 4400												