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PROFICIENCY TEST « RAEMA »

animal société aliment

SCHEME N° 70 (9th MARCH 2020) GENERAL REPORT



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1. GENERAL DATA

1.1.PARTICIPATING LABORATORIES

343 laboratories participated to the 70th scheme. The sending was made on Monday 9th March 2020. We received **305** answers (88.9%). Due to the confinement period, this percentage is less important than usual. Considering the number of replies, this has no impact on the robustness of the statistical analysis.

1.2.DELIVERY TIME OF THE PARCEL

Reception	J0	J0+1	J0+2	J0+3	J0+4	J0+5	J0+7	J0+8	J0+15
Nb of laboratories	4	206	46	23	16	2	5	1	1

1.3.INFORMATIONS ABOUT SAMPLE

1.3.1. NATURE

The sample included:

- one strain of *Enterococcus sp.* at a concentration level of 10⁵ cfu/g in 5 units;
- one strain of Citrobacter sp. at a concentration level of 10³ cfu/g in 5 units;
- one strain of Serratia liquefaciens at a concentration level of 5.10² cfu/g in 5 units;
- one strain of Escherichia coli at a concentration level of 5.10² cfu/g in 5 units;
- one strain of Clostridium perfringens at a concentration level of 2.102 cfu/g in 2 units;
- one strain of Staphylococcus aureus at a concentration level of 2.10³ cfu/g in 5 units;
- one strain of Salmonella Anatum at a concentration level of 25 cfu/g in 2 units;
- one strain of Listeria monocytogenes at a concentration level of 2.10³ cfu/g in 4 units.

1.3.2. SIZE

180 kilogrammes of powder were produced, and distributed after contamination in bottles containing 75 grammes at least. Bottles were covered by a label with a 6 digit identification number.

⁽¹⁾Coordinator of the proficiency test « RAEMA »

1.3.3. HOMOGENEITY AND STABILITY TEST OF THE CONTAMINATION

Homogeneity and stability of samples are checked during the statistical analysis of participants results. A supplementary check of the contamination's homogeneity was realized on 10 samples for each unit by a double enumeration of aerobic microorganisms at 30°C.

The contamination's stability was also checked by enumeration / detection of all flora on 16, 23 and 30 March 2020. These checks were realized by a subcontractor accreditated by Cofrac.

1.3.4. FLORA FOR ENUMERATION / DETECTION

Enumeration of the following flora was proposed: microorganisms at 30°C, Enterobacteriaceae, total and thermotolerant coliforms, beta-glucuronidase positive *Escherichia coli*, anaerobic sulfite-reducing bacteria, *Clostridium perfringens*, coagulase positive staphylococci, *Listeria monocytogenes*, as well as detection of *Salmonella* and *Listeria monocytogenes*.

1.4.EXECUTION OF ANALYZES

1.4.1. DELIVERY TIME OF SAMPLES / BEGINNING OF ANALYZES

305 laboratories (100%) specified it.

Analysis time	J0+1	J0+2	J0+3	J0+4	J0+5	J0+6	J0+7	J0+8	J0+9	J0+10	J0+14	J0+15	J0+16	J0+17	J0+22
Nb of laboratories	32	57	28	7	1	1	116	34	9	2	10	5	1	1	1

1.4.2. PRESERVATION TEMPERATURE OF SAMPLES BEFORE ANALYSIS

301 laboratories (98.7%) specified it. The average temperature is **4.0°C** with a standard deviation of 1.2°C. The given data 20 and 21°C given by 3 laboratories were not taken into account for this calculation.

2. EXPLOITATION OF ANALYSIS REPORT

2.1.PREPARATION OF THE INITIAL SUSPENSION

For **305** answers (100%):

191 laboratories (62.6%) prepare the initial suspension with adding diluent to powder.

114 laboratories (37.4%) prepare the initial suspension with adding powder to diluent.

2.2.DILUENT USED FOR THE INITIAL SUSPENSION

This data has been added to have all needed elements for this stage.

For **304** answers (99.7%):

271 laboratories (88.9%) use Buffered Peptone Water for the initial suspension.

33 laboratories (10.8%) used another diluent for the initial suspension.

2.3.HOMOGENEIZATION TECHNIQUE

For **304** answers (99.7%):

290 laboratories (95.1%) homogeneize their sampling with a Stomacher ND.

14 laboratories (4.6%) used another technique (manual, magnétic or other).

2.4.RESUSCITATION'S CONDITIONS

2.4.1. DURATION

297 laboratories (97.4%) specified it.

The average duration is **27.2 min** with a standard deviation of 15.3 min. The data 120, 180 and 1440 min given by 6 laboratories was not taken into account for this calculation.

70th scheme (edition 07/05/20)



2.4.2. TEMPERATURE

297 laboratories (97.4%) specified it.

The average temperature is **21.2°C** with a standard deviation of 3.4°C.

2.5.MICROORGANISM AT 30°C

Parameters	Mode	Nb laboratories
Method	NF EN ISO 4833-1 AFNOR 3M-01/1-09/89 NM ISO 4833-1 NF EN ISO 4833-2 AFNOR BIO-12/35-05/13 XP V08-034 Other	180 45 18 15 11 4
	+ V08-100 (spiral)	14
Culture medium	Plate Count Agar Petrifilms Plate Count Agar + Milk Tempo AC Other	215 47 16 11 1
Preparation	Home made Ready to use not pre-poured Ready to use, plate, film, card	106 118 66
Plating method	Surface Pour Culture medium for card	61 214 12
1 st dilution retained	- 1 - 2 - 3 - 4 - 5 1/400 1/4000	13 14 239 10 1 8 1
Incubation temperature	30°C 37°C	287 2
Incubation duration	69-73 h 40-48 h 24-26 h	243 42 4



2.6.ENTEROBACTERIACEA

Parameters	Mode	Nb laboratories
Method	NF V08-054 → NM 08.0.109 ⁽¹⁾ NF EN ISO 21528-2 AFNOR 3M-01/6-09/97 AFNOR BIO-12/21-12/06 AFNOR AES-10/07-01/08 AFNOR BRD-07/24-11/13 Other + V08-100 (spiral)	107 22 63 41 9 8 3 3
Culture medium	VRBG Petrifilms Tempo EB Rebecca Rapid'Enterobacteriaceae Other	189 43 9 9 3 2
Preparation	Home made Ready to use not pre-poured Ready to use, plate, film, card	84 117 55
1 st dilution retained	- 1 - 2 1/40 1/400	188 57 1 7
Incubation temperature	37±1°C 30°C 35°C	155 92 9
Incubation duration	18-24 h 48 h	251 5
Confirmatory test (New technical data requested to cover all stages of the standard)	Yes No	51 199

⁽¹⁾ Similar method to NF V08-054 according to ONSSA (Office National de Sécurité Sanitaire des produits Alimentaires).



2.7.TOTAL COLIFORMS

216 laboratories performed the enumeration.

Parameters	Mode	Nb laboratories
Method	NF V08-050 → NM 08.0.142 ⁽²⁾ NF ISO 4832 AFNOR 3M NM ISO 4832 AFNOR BIO-12/17-12/05 AFNOR BRD-07/08-12/04 Other + V08-100 (spiral)	109 10 53 20 13 4 3 4
Culture medium	VRBL Petrifilms Tempo TC Rapid Ecoli Other	186 22 4 3 1
Preparation	Home made Ready to use not pre-poured Ready to use, plate, film, card	83 107 25
1 st dilution retained	-1 -2 1/40 1/400	185 24 1 3
Incubation temperature	30°C 37±1°C	202 14
Incubation duration	20-27 h 48 h	214 1

AFNOR 3M method including:

³ laboratories specified utilization of AFNOR 3M-01/02-09/89 A method.

¹ laboratory specified utilization of AFNOR 3M Petrifilm CC method.

 $^{^{(2)}}$ Similar method to NF V 08-050 according to ONSSA.



2.8.THERMOTOLERANT COLIFORMS

197 laboratories performed the enumeration.

Parameters	Mode	Nb laboratories
Method	NF V08-060 → <i>NM 08.0.124</i> ⁽³⁾ AFNOR 3M NF ISO 4832 Other + V08-100 (spiral)	140 26 20 8 3
Culture medium	VRBL Petrifilms Other	175 21 1
Preparation	Home made Ready to use not pre-poured Ready to use, plate, film, card	77 99 21
1 st dilution retained	-1 -2	176 19
Incubation temperature	42-45°C 37°C 30°C	194 2 1
Incubation duration	20-24 h 48 h	196 1

AFNOR 3M method including:

- 4 laboratories specified utilization of AFNOR 3M-01/02-09/89 C method.
- 1 laboratory specified utilization of AFNOR 3M-01/05-03/97 B method.
- 1 laboratory specified utilization of AFNOR 3M Petrifilm CC method.

 $^{^{(3)}}$ Similar method to NF V08-060 according to ONSSA.

2.9.ESCHERICHIA COLI

271 laboratories performed the enumeration.

Parameters	Mode	Nb laboratories
Method	NF ISO 16649-2 AFNOR 3M NM ISO 16649-2 AFNOR BRD-07/01-07/93 AFNOR BIO-12/13-02/05 AFNOR AES-10/06-01/08 NF EN ISO 16649-3 AFNOR BIO-12/05-01/99 Other + V08-100 (spiral)	155 39 21 14 10 9 5 3 15
Culture medium	TBX Petrifilms Rapid E. coli Rebecca Tempo EC Coli ID Other	183 41 19 11 10 6
Preparation	Home made Ready to use not pre-poured Ready to use, plate, film, card	87 132 52
Plating method	Surface Pour Culture medium for card	39 217 13
1 st dilution retained	-1 -2 1/40 1/400	242 16 3 6
Incubation temperature	41-46°C 37±1°C 30°C	236 34 1
Incubation duration	16-25 h 44-48 h	267 4

AFNOR 3M method including:

9 laboratories specified utilization of AFNOR 3M-01/08-06/01 (SELECT'E. COLI) method.



2.10. ANAEROBIC SULFITE-REDUCING BACTERIA

Parameters	Mode	Nb laboratories
Method	NF V08-061 \rightarrow NM 08.0.154 ⁽⁴⁾ \rightarrow NM 08.0.125 ⁽⁴⁾ NF ISO 15213 NM ISO 15213 Other	140 3 11 39 15 9
Culture medium	TSC Iron Sulfite agar TSN Other	203 6 5 3
Preparation	Home made Ready to use not pre-poured Ready to use, plate, film, card	87 109 21
Seeding way (New technical data requested to take into account different seeding ways)	Plates Tubes	135 80
1 st dilution retained	-1 -2	182 33
Incubation temperature	44-49°C 37°C 30°C	151 65 1
Incubation duration	18-24 h 48 h 72 h 12-16 h	180 31 4 2

⁽⁴⁾ Similar method to NF V08-061 according to ONSSA.



2.11. CLOSTRIDIUM PERFRINGENS

Parameters	Mode	Nb laboratories
Method	NF EN ISO 7937	136
	NM ISO 7937	19
	Other	14
Culture medium	TSC	167
	Other	2
Preparation	Home made	59
. roparation	Ready to use not pre-poured	105
	Ready to use, plate, film, card	5
	, , , ,	
1 st dilution retained	-1	158
	-2	8
Incubation temperature	37°C	158
	44-46°C	11
	10 04 h	162
Incubation duration	18-24 h 48 h	163 5
	72 h	1
	12 11	ı
Confirmation test	None	30
	Lactose-sulfite	126
	Strip	5
	Other	4



2.12. COAGULASE POSITIVE STAPHYLOCOCCI

Parameters	Mode	Nb laboratories
Method	NF EN ISO 6888-2 NF V 08-057-1 → NM 08.0.112 ⁽⁵⁾ NF EN ISO 6888-1 NM ISO 6888-1 AFNOR 3M-01/9-04/03 AFNOR BKR-23/10-12/15 AFNOR BIO-12/28-04/10 NordVal No :049	120 40 5 36 18 15 11
	NM ISO 6888-2 Other	4 7
	+ V08-100 (spiral)	4
Culture medium	RPF BP+egg yolk tellurite BP+egg yolk tellurite+ sulfamethazine Petrifilm Easy Staph Tempo STA Rapid Staph Other	124 81 16 16 16 10 5
Preparation	Home made Ready to use not pre-poured Ready to use, plate, film, cards	63 110 97
Plating method	Surface Pour Culture medium for card	138 119 11
1 st dilution retained	-1 -2 -3 1/40 1/400	105 149 4 5 3
Incubation temperature	37±1°C 30°C	266 4
Incubation duration	42-48 h 18-25 h 72 h	191 78 1
Confirmation test	None Staphylo-coagulase Clumping factor DNase Other	161 82 7 11 7

⁽⁵⁾ Similar method to NF V 08-057-1 according to ONSSA.



2.13. LISTERIA MONOCYTOGENES - ENUMERATION

212 laboratories performed the enumeration.

RESUSCITATION

77 laboratories announce the realization of a resuscitation step.

The average duration for these laboratories is **46.3 min** with a standard deviation of 24.9 min. The average temperature for these laboratories is **21.2°C** with a standard deviation of 3.6°C.

Parameters	Mode	Nb laboratories
Method	AFNOR AES-10/05-09/06 NF EN ISO 11290-2 AFNOR BKR-23/05-12/07 NM ISO 11290-2 AFNOR BRD-07/05-09/01 AFNOR BRD-07/17-01/09	61 57 45 18 17 6
	Other	8
Resuscitation step	Yes No	77 121
Resuscitation medium	Buffered Peptone Water Fraser base Other	60 12 3
Enumeration medium	ALOA Count Compass Listeria Rapid Lmono AL Agar OCLA Palcam Other	101 66 21 12 5 4
Preparation	Home made Ready to use not pre-poured Ready to use, plate, film, card	30 46 135
Plating method	Surface Pour Culture medium for card	173 37 0



Parameters	Mode	Nb laboratories
1 st dilution retained	-1 -2 -6	139 69 1
Incubation temperature	37°C 30°C	210 2
Incubation duration	46-51 h 22-24 h	176 36
Confirmation test	None Biochemical Biochemical + CAMP Other	45 128 29 8
Nb of colonies tested per plate	1 2-3 5 10 18 100 150	48 16 87 1 1 1 2

2.14. SALMONELLA - DETECTION

275 laboratories performed the detection.

Methods used by laboratories are clarified in the following table :

Parameter	Mode	Nb laboratories
Method	NF EN ISO 6579-1	73
	AFNOR BKR 23/07-10/11 (IRIS Salmonella)	64
	AFNOR BRD 07/11-12/05 (Rapid Salmonella)	33
	NM ISO 6579-1	28
	AFNOR BIO 12/16-09/05 (VIDAS Easy Salmonella)	21
	AFNOR BIO 12/32-10/11 (VIDAS SPT)	20
	AFNOR BIO 12/41-03/17 (SALMA One day)	18
	Other	18

No detail of methodology was asked to laboratories using other methods than NF EN ISO 6579-1 and NM ISO 6579-1 methods and proposed in the questionnaire.

You will find below a short description of these methods :

Method	Pre-enrichment	Enrichment	Isolation
AFNOR BIO 12/16-09/05 VIDAS Easy Salmonella	BPW / 37°C - 16/20h	SX2 / 41,5°C - 22/26h	Chrom ID / 37°C - 24h
AFNOR BIO 12/32-10/11 VIDAS SPT		BPW + Salmonella supplement / 41,5°C - 18/24h	Chrom ID / 37°C - 24h
AFNOR BKR 23/07-10/11 IRIS Salmonella		IRIS Salmonella Enrichment / 41,5°C - 18±2h	IRIS / 37°C - 24±3h
AFNOR BRD 07/11-12/05 Rapid Salmonella		BPW + Salmonella capsule / 41,5°C - 18±2h	Rapid Salmonella / 37°C - 24±2h
AFNOR BIO 12/41-03/17 SALMA One day		BPW + Salmonella supplement / 41.5°C – 16/24h	SALMA / 37°C - 24±3h



The detail of the methodology followed by 101 laboratories using NF EN ISO 6579-1 and NM ISO 6579-1 method, and the 18 laboratories using another method, is clarified in the following table :

Parameter	Mode	Nb laboratories
Method	NF EN ISO 6579-1 NM ISO 6579-1 Other	73 28 18
Pre-enrichment medium	Buffered Peptone Water Other	113 5
Pre-enrichment temperature	36-37°C 41-42.5°C 20-22°C	111 5 2
Pre-enrichment duration	16-20 h 22-26 h	81 37
Enrichment medium	RVS MKTTn Selenite-cystine broth Other	103 95 19 5
Isolation medium	XLD Hektoen Bismuth Sulfate ASAP Brilliance Salmonella IRIS Salmonella agar GVB SS Compass Salmonella Rambach Rapid Salmonella Other	94 34 18 11 10 10 10 7 5 5 3
Confirmation test	Biochemical Biochemical + serological agglutination	49 60
	Other	7

2.15. LISTERIA MONOCYTOGENES - DETECTION

237 laboratories performed the detection.

Parameter	Mode	Nb laboratories
Method	AFNOR AES 10/03-09/00 (ALOA one day)	61
	AFNOR BKR 23/02-11/02 (Compass L. mono)	50
	NF EN ISO 11290-1	44
	NM ISO 11290-1	24
	AFNOR BRD 07/04-09/98 (Rapid' L. mono)	21
	AFNOR BIO 12/11-03/04 (VIDAS LMO2-37°C)	8
	AFNOR BIO 12/27-02/10 (VIDAS LMX)	7
	AFNOR BRD 07/16-01/09 (Agar Listeria)	6
	AFNOR BIO 12/02-06/94 (VIDAS Listeria)	2
	AFNOR UNI 03/04-04/05 (Listeria Precis)	2
	Other	12

No detail of methodology was asked to laboratories using other method than NF EN ISO 11290-1 and NM ISO 11290-1 methods and proposed in the questionnaire.

You will find below a short description of these methods :

Méthod	Primar	Primary enrichment		ndary enrichment	Isolation
Wethod	Medium	Incubation	Medium	Incubation	isolation
AFNOR BRD 07/04-09/98 Rapid' L. mono	Fraser 1/2	30°C - 24±2h			Rapid L'mono 37°C – 24h
AFNOR BIO 12/02-06/94 VIDAS Listeria	Fraser 1/2	37°C - 26/30h	Fraser	30°C - 24/26h	Palcam et Oxford 37°C – 24h
AFNOR BIO 12/27-02/10 VIDAS LMX	LMX	37°C - 26/30h			ChromID 37°C – 24h
AFNOR BIO 12/11-03/04 VIDAS LMO2 (37°C)	Fraser 1/2	30°C - 24/26h	Fraser	37°C - 24/26h	ChromID 37°C – 24h
AFNOR AES 10/03-09/00 ALOA one day	Fraser 1/2	30°C - 24±2h			ALOA One Day 37°C – 24/48h
AFNOR BKR 23/02-11/02 Compass L. mono	Fraser 1/2	30°C - 24±2h			Compass Listeria Agar 37°C – 24h
AFNOR BRD 07/16-01/09 Agar Listeria	Fraser 1/2	30°C - 24±2h			Agar Listeria 37°C – 24h
AFNOR UNI 03/04-04/05 Listeria Precis	One Broth Listeria	30°C - 24±2h			Brilliance Listeria 37°C – 24h



The detail of the methodology followed by 68 laboratories using NF EN ISO 11290-1 and NM ISO 11290-1 methods, and the 12 laboratories using another method, is clarified in the following table :

Parameter	Mode	Nb laboratories
Method	NF EN ISO 11290-1 NM ISO 11290-1 Other	44 24 12
Primary enrichment medium	Half-Fraser One broth Listeria Other	66 3 10
Primary enrichment temperature	30°C 37°C	71 8
Primary enrichment duration	20-28 h 48 h	77 1
Secondary enrichment medium	Fraser Other	64 3
Secondary enrichment temperature	37±1°C 30°C 24°C	63 4 1
Secondary enrichment duration	22-25 h 46-48 h	54 14
Isolation medium	Palcam Ottaviani et Agosti Compass Listeria Oxford Rapid L'mono Brilliance Listeria Other	48 39 28 12 4 3
Isolation temperature	37°C 30°C	74 1
Isolation duration	46-48 h 24 h	50 25
Confirmation test	None Biochemical Biochemical + CAMP Other	5 47 24 1
Nb of colonies per plate	1 2-4 5 6	25 9 31 1



3. ASSESSMENT OF PERFORMANCE (INDIVIDUEL REPORTS)

3.1.PERFORMANCES IN ENUMERATION

Performance is assessed on two criteria : **precision and trueness**.

The assigned value of the contamination used to assess the trueness and the reference value for the assessment of the precision are consensual values obtained with the results of all the participants. These values are obtained by robust estimation methods in order to eliminate influence of aberrant results. However some results are excluded of the statistical analysis. That is the case when laboratories do not give results for all contaminated units, when results are "less than CFU/g", when samples are analyzed after the deadline (time of receipt > 4 days after sending or time of analysis >15 days after sending) or when this information is not specified.

A statistical analysis has also be done to highlight potential relations between techniques used (delay of analysis, preservation temperature, preparation of the initial suspension, homogenization technique, resuscitation conditions, method used, media used, manufacturers of media, preparation mode, plating method, incubation conditions) and results obtained. We need to clarify that this statistical link is not involved in a cause - effect relationship. Indeed, this link may be due to a not documented factor.

When a significant statistical link is identified between use of a technique and the obtained results, the assessment of performance is done considering the influence of one or several factors involved if their effect translates into a contamination's difference higher than 0.15 log CFU/g for non-selective media or higher than 0.30 log CFU/g for selective media (these limits match with productivity limits of culture media usually recommended in the standard NF EN ISO 11133).

PRECISION

The precision reflects the repeatability (or reproducibility intra-laboratory) of your work.

The standard deviation of your results, s, is compared to the robust estimation of the standard deviation (reference standard deviation of precision), s^* , obtained with algorithm S from the standard ISO 13528 applied to all standard deviations obtained by laboratories included in the statistical analysis.

An index score is then calculated using the following formula: $i = (k-1) \cdot \frac{s^2}{s^{*2}}$ (with k, number of contaminated units and retained in the statistical analysis, usually 5).

The standard ISO 13528 do not provide warning and action limits for this score, so its interpretation is left to your discretion.

As an indicator, we suggest following values by analogy with those indicated for the evaluation of trueness.

For k=5, a score lower than 0.1 or higher than 18 may be considered as an action signal and a score lower than 0.45 or higher than 11.5 may be considered as a warning signal.

For k=4, a score lower than 0.03 or higher than 15.5 may be considered as an action signal and a score lower than 0.2 or higher than 9.5 may be considered as a warning signal.

For k=3, a score lower than 0.003 or higher than 13.2 may be considered as an action signal and a score lower than 0.05 or higher than 7.5 may be considered as a warning signal.

For k=2, a score lower than 0.000002 or higher than 10.3 may be considered as an action signal and a score lower than 0.0008 or higher than 5.2 may be considered as a warning signal.



TRUENESS

The trueness reflects the closeness of the mean of your results to the contamination's assigned value of samples. It has been evaluated for all enumerated flora.

The mean of your results in log CFU/g, m (on contaminated units and included in the statistical analysis), is compared to the contamination's assigned value, $m_{\rm pt}$, obtained with algorithm A from the standard ISO 13528 applied to all laboratories mean included in the statistical analysis. When groups are formed, each one is characterize by its own assigned value.

A z score is then calculated with the following formula: $z = \frac{m - m_{pt}}{\sigma_{pt}}$, where σ_{pt} is the standard

deviation for proficiency assessment (robust estimation of the standard deviation obtained by participants).

The standard ISO 13528 specifies that z score lower than -3 or higher than +3 must be considered as an action signal and that a z score lower than -2 or higher than +2 must be considered as a warning signal.

In this report, we specify, estimations of interlaboratories standard deviation for enumerations proposed as well as reproducibility standard deviation or global standard deviation for the test (parameters including interlaboratories variability and the variability of the precision).

INDIVIDUAL REPORTS - FOR EACH CRITERIA YOU FIND THE FOLLOWING INFORMATIONS

- your results in logarithm base 10 (-1 when the answer is < limit and NaN when there is no answer). Comment: the presentation order of your results does not necessarily correspond to the order you sent them, this order is the same for all the flora.
- histogram for the studied parameter (laboratories standard deviations for the precision and laboratorie's means for the trueness) with an asterisk indicating the location of your result,
- standard deviation (precision) or mean (trueness) of your results (on contaminated units and retained in the statistical analysis),
- the method declared in your results input,
- when necessary, your group in relation to the technique used,
- precision score or z score,
- number of laboratories which made analysis (and belonging to your group),
- number of laboratories included in the statistical analysis,
- reference standard deviation for the precision or assigned value of the contamination and standard deviation aptitude assessment (trueness),
- number of laboratories with a satisfactory signal,
- number of laboratories with a warning signal,
- number of laboratories with an action signal.

3.1.1. MICROORGANISMS AT 30°C

None significant effect of the analysis technique has been highlighted.

Microorganisms at 30°C			
Assigned value of the contamination (log CFU/g)	4.876		
Assigned value uncertainty (log CFU/g)	0.0066		
Standard deviation for proficiency assessment (log CFU/g)	0.0882		
Standard deviation for precision (log CFU/g)	0.0580		
Interlaboratory's standard deviation (log CFU/g)	0.0843		
Reproducibility standard deviation (log CFU/g)	0.1023		

3.1.2. ENTEROBACTERIACEAE

A significant "effect" of the culture media, manufacturer and the retained dilution has been highlighted. This effect results in a contamination's difference higher than 0.3 log CFU/g, then results have been gathered in three groups :

Enterobacteriaceae	Group 1	Group 2	Group 3
Assigned value of the contamination (log CFU/g)	2.829	2.996	3.169
Assigned value uncertainty (log CFU/g)	0.0184	0.0358	0.0327
Standard deviation for proficiency assessment (log CFU/g)	0.1968	0.1765	0.1382
Standard deviation for precision (log CFU/g)		0.0897	
Interlaboratory's standard deviation (log CFU/g)	0.1927	0.1719	0.1323
Reproducibility standard deviation (log CFU/g)	0.2125	0.1939	0.1598

3.1.3. TOTAL COLIFORMS

A significant "effect" of the culture media and manufacturer has been highlighted. This effect results in a contamination's difference lower than 0.3 log CFU/g, then results have been gathered in one group:

Total coliforms			
Assigned value of the contamination (log CFU/g)	2.790		
Assigned value uncertainty (log CFU/g)	0.0198		
Standard deviation for proficiency assessment (log CFU/g)	0.2270		
Standard deviation for precision (log CFU/g)	0.0857		
Interlaboratory's standard deviation (log CFU/g)	0.2238		
Reproducibility standard deviation (log CFU/g)	0.2396		

3.1.4. THERMOTOLERANT COLIFORMS

A significant "effect" of the retained dilution has been highlighted. This effect results in a contamination's difference lower than 0.3 log CFU/g, then results have been gathered in one group:

Thermotolerant coliforms	
Assigned value of the contamination (log CFU/g)	2.759
Assigned value uncertainty (log CFU/g)	0.0182
Standard deviation for proficiency assessment (log CFU/g)	0.1987
Standard deviation for precision (log CFU/g)	0.0851
Interlaboratory's standard deviation (log CFU/g)	0.1950
Reproducibility standard deviation (log CFU/g)	0.2127

3.1.5. ESCHERICHIA COLI

None significant effect of the analysis technique has been highlighted.

Escherichia coli	
Assigned value of the contamination (log CFU/g)	2.719
Assigned value uncertainty (log CFU/g)	0.0154
Standard deviation for proficiency assessment (log CFU/g)	0.1992
Standard deviation for precision (log CFU/g)	0.0920
Interlaboratory's standard deviation (log CFU/g)	0.1949
Reproducibility standard deviation (log CFU/g)	0.2155

3.1.6. ANAEROBIC SULFITE-REDUCING BACTERIA

Only units n°4 and 5 were artificially contaminated.

A significant "effect" of the culture media and the retained dilution has been highlighted. This effect results in a contamination's difference lower than 0.3 log CFU/g, then results have been gathered in one group:

Anaerobic sulfite-reducing bacteria	
Assigned value of the contamination (log CFU/g)	2.321
Assigned value uncertainty (log CFU/g)	0.0183
Standard deviation for proficiency assessment (log CFU/g)	0.2092
Standard deviation for precision (log CFU/g)	0.1037
Interlaboratory's standard deviation (log CFU/g)	0.1960
Reproducibility standard deviation (log CFU/g)	0.2217

Comment:

- 9 laboratories detected ASR in unit n°1 non-artificially contaminated by *C. perfringens* with a contamination level from 10 to 1900 CFU/g.
- 7 laboratories detected ASR in unit n°2 non-artificially contaminated by *C. perfringens* with a contamination level from 20 to 1600 CFU/g.
- 11 laboratories detected ASR in unit n°3 non-artificially contaminated by *C. perfringens* with a contamination level from 2 to 1500 CFU/g.

3.1.7. CLOSTRIDIUM PERFRINGENS

Only units n°4 and 5 were artificially contaminated.

A significant "effect" of the culture media has been highlighted. This effect results in a contamination's difference lower than 0.3 log CFU/g, then results have been gathered in one group:

Clostridium perfringens		
Assigned value of the contamination (log CFU/g)	2.300	
Assigned value uncertainty (log CFU/g)	0.0189	
Standard deviation for proficiency assessment (log CFU/g)	0.1902	
Standard deviation for precision (log CFU/g)	0.0985	
Interlaboratory's standard deviation (log CFU/g)	0.1770	
Reproducibility standard deviation (log CFU/g)	0.2026	

Comment:

- 4 laboratories detected *C. perfringens* in unit n°1 non-artificially contaminated by *C. perfringens* with a contamination level from 100 to 1500 CFU/g.
- 4 laboratories detected *C. perfringens* in unit n°2 non-artificially contaminated by *C. perfringens* with a contamination level from 10 to 1400 CFU/g.
- 5 laboratories detected *C. perfringens* in unit n°3 non-artificially contaminated by *C. perfringens* with a contamination level from 50 to 1400 CFU/g.

3.1.8. COAGULASE POSITIVE STAPHYLOCOCCI

None significant effect of the analysis technique has been highlighted.

Coagulase positive Staphylococci		
Assigned value of the contamination (log CFU/g)	3.441	
Assigned value uncertainty (log CFU/g)	0.0124	
Standard deviation for proficiency assessment (log CFU/g)	0.1594	
Standard deviation for precision (log CFU/g)	0.0823	
Interlaboratory's standard deviation (log CFU/g)	0.1551	
Reproducibility standard deviation (log CFU/g)	0.1756	

3.1.9. LISTERIA MONOCYTOGENES

Only units n°2, 3, 4 and 5 were artificially contaminated.

None significant effect of the analysis technique has been highlighted.

Listeria monocytogenes	
Assigned value of the contamination (log CFU/g)	3.367
Assigned value uncertainty (log CFU/g)	0.0097
Standard deviation for proficiency assessment (log CFU/g)	0.1098
Standard deviation for precision (log CFU/g)	0.0661
Interlaboratory's standard deviation (log CFU/g)	0.1047
Reproducibility standard deviation (log CFU/g)	0.1238

3.2.PERFORMANCES IN DETECTION

The performance is assessed by the capacity to detect only samples contaminated by *Salmonella* and *Listeria monocytogenes* (no false positive or false negative results).

3.2.1. DETECTION - SALMONELLA

Only units n°4 and 5 were artificially contaminated.

261 laboratories obtained correct results.

5 laboratories obtained false positive results (respectively 3, 2 and 3 false-positive for units n°1, 2 and 3).

11 laboratories obtained false negative results (respectively 4 and 9 false-negative for units n°4 and 5).

3.2.2. DETECTION - LISTERIA MONOCYTOGENES

Only units n°2, 3, 4 and 5 were artificially contaminated.

234 laboratories obtained correct results.

2 laboratories obtained false positive results for unit n°1.

3 laboratories obtained false negative results (respectively 1, 0, 0 and 2 false-negative for units $n^{\circ}2$, 3, 4 and 5).

3.3.EVOLUTION OF PERFORMANCE

You will find, on each page of your performance's assessment, a graph representing evolution of it on different tests since the 50^{th} scheme.

In order to interpret your control card with z scores, you can refer to the standard ISO 13528 $\S10.8.2.2$, explaining the 3 « out of control » situations :

- Just one overtaking of the action limit (z<-3 or z>3),
- 2 consecutives z scores out of 3 overtaking of the warning limit (2<z<3 or -3<z<-2),
- 6 consecutives z scores regularly increasing or decreasing.