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# Inter laboratory comparison (ILC) report\* ISTA PT22-SH 7-007

ISTA Proficiency test: Detection of *Phomopsis complex* on *Glycine max* (soybean) seeds

\*Original report signed and archived

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#### **PROFICIENCY TEST ORGANIZATION**

The aim of this Proficiency Test was to verify the ability of laboratories to detect and identify *Phomopsis complex* in soybean (*Glycine max*) seeds.

Table 1 : Schedule of the Proficiency Test

Sending of samples	7 <sup>th</sup> of November 2022
Deadline to begin analysis	21 <sup>st</sup> of November 2022
Deadline to send results	23 <sup>rd</sup> of December 2022
Sending by GEVES of report and individual sheet	28 <sup>th</sup> of February 2023

<sup>17</sup> laboratories participated to this test and were randomly allocated a number, so that results remained anonymous.

#### Notation of results

The laboratories indicated: qualitative and quantitative results for each sample and information about the method used.

## Composition of the sample panel

10 samples of 400 soybean seeds have been sent to each laboratory with different replicates per level of contamination, see table n°2.

Table 2: Characteristics of samples

Level of contamination	Number of samples	Expected value
Healthy	3	Negative
Medium	4	Low positive
High	3	High positive

## Validation of samples

The samples have been validated through homogeneity and stability tests.

The results of participating laboratories were compared to the expected results determined by the homogeneity and stability tests results.

The pretest results were confirmed by homogeneity and stability test.

#### Pre-test

We had four seed lots:

- A: Healthy lot
- C: Highly infected lot (already tested: high natural infection 18.3%)
- D: Healthy lot
- E: Highly infected lot (already tested: high natural infection 20%)

We have tested 10 subsamples of 400 seeds for the A lot, the 28<sup>th</sup> of April 2022 and 10 subsamples of 400 seeds for the C lot, the 2<sup>nd</sup> of May 2022. The method used is the ISTA 7-016 method. The results are given in table n°3.

Lots D and E were tested by the same method on 28<sup>th</sup> April 2022. With knowledge of the results on D and E lots, we created a medium infected lot:

• B: Medium infected lot: with 300 seeds of D lot and 100 seeds of E lot

For the pretest on the lot B, we tested 5 samples with 340 seeds of D lot and 60 seeds of the E lot, the average result was  $3.55 \% \pm 0.4$  of infection. In order to determine the number of infected seeds to mix with healthy seeds, to have a medium level of infection, we extrapolated the results of the pre-tests to obtain an infection level of 5.8 %. 300 healthy seeds mixed with 100 highly infected seeds would give a contamination rate for the medium level of 5.8 %. The results are given in table n°3.

Table 3: Re	sults of	f pre-test
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Codification of lot	Level of infection	Obtained results	Decision	Test execution date
А	Healthy	0 %	Accept	28/04/2022
В	Medium	5.8 % (±0.4)	Accept	02/05/2022
С	High	18.3 % (±1.2)	Accept	02/05/2022

#### Homogeneity Test

Homogeneity test was done after packaging and just before shipping of the seed samples to the participating laboratories. The method used is the ISTA 7-016 method. 10 extra samples of 400 seeds representing each infection level were tested. The samples were tested in October 2022.

The qualitative results, the minimum, the maximum and the average values are given in the table n°4. The quantitative results were analyzed by Hampel's method (Table n°5), by repartition against the mean (figure 1) and by box plot method (figure 2).

Table 4: Results of homogeneity test

Codification	Level of	Expected	Quantitative results		Qualitative	Conformity
of lot	infection	result	min - max	Average	result	Comornity
А	Healthy	Not detected	0 %	0 %	0+/10	Conform
В	Medium	Detected	4.5 % – 7.29 %	5.82 %	10 <sup>+</sup> /10	Conform
С	High	Detected	7.5 % - 20.25 %	15.97 %	10 <sup>+</sup> /10	Conform

Table 5: Homogeneity test results for medium and high levels according to the Hampel's method

Medium				High			
Rep	Values (Xi)	Xi - M	Status	Rep	Values (Xi)	Xi - M	Status
Homog.1	6.35	0.345	OK	Homog.1	17 <mark>.50</mark>	0.125	ОК
Homog.2	5.00	1.005	OK	Homog.2	12.75	4.625	ОК
Homog.3	7.2 <mark>9</mark>	1.285	OK	Homog.3	17 <mark>.25</mark>	0.125	ОК
Homog.4	6.50	0.495	OK	Homog.4	18 <mark>.25</mark>	0.875	ОК
Homog.5	6.25	0.245	ОК	Homog.5	18. <mark>50</mark>	1.125	ОК
Homog.6	4.82	1.185	ОК	Homog.6	7.50	9.875	ОК
Homog.7	4.50	1.505	ОК	Homog.7	13.50	3.875	ОК
Homog.8	6.58	0.575	OK	Homog.8	20.2 <mark>5</mark>	2.875	ОК
Homog.9	5.13	0.875	OK	Homog.9	<mark>14.75</mark>	2.625	ОК
Homog.10	5.76	0.245	ОК	Homog.10	19. <mark>50</mark>	2.125	ОК
Median (M): 6.005 MAD: 0.725 5.2 X MAD 3.770					Median (M): MAD: 5.2 X MAD	17.375 2.375 12.350	

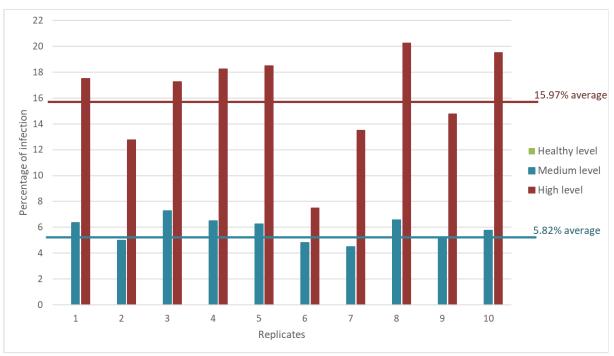


Figure 1: Homogeneity test results, repartition against the mean

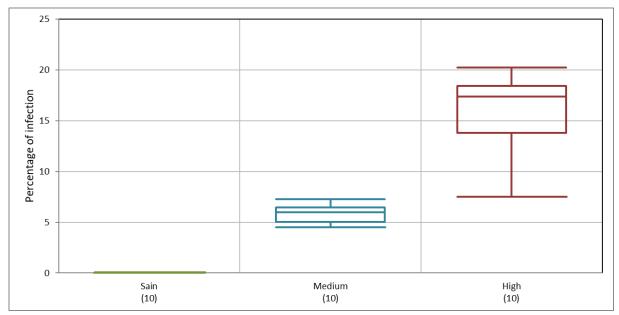


Figure 2: Boxplot representing results of the homogeneity test

## **Conclusion of homogeneity test:**

#### Results of healthy level

All samples were negative, there were no false positive results.

#### Results of medium level

Samples are homogeneous, there was no outlier. The average obtained was very close to the expected percentage based on the pre-test.

#### Results of high level

Samples are homogeneous, there was no outlier. The average obtained was close to the expected percentage based on the pre-test.

## **Stability Test**

Stability test has been started the 5<sup>th</sup> of December 2022. The method used is the ISTA 7-016 method. 5 samples representing the healthy level, 10 samples representing the medium level and 5 samples representing the high levels were tested.

The qualitative results, the minimum, the maximum and the average values are given in the table  $n^{\circ}6$ . The quantitative results were analyzed by Hampel's method (table  $n^{\circ}7$ ), by repartition against the mean (figure 3) and by box plot method (figure 4).

Table 6: Results of stability test

Codification of lot	Level of infection	Expected result	Value min - max	Average	Qualitative result	Conformity
А	Healthy	Not detected	0%	0%	0+/5	Conform
В	Medium	Detected	4% – 7.05%	5.11%	10+/10	Conform
С	High	Detected	13.38% - 18.25%	15.43%	5 <sup>+</sup> /5	Conform

dium				High			
Rep	Values (Xi)	Xi - M	Status				
Stab.1	4.50	0.500	ОК				
Stab.2	4.26	0.740	ОК	Don	\/alaa /\/:\	LV: NAI	Chahira
Stab.3	5.53	0.530	ОК	Rep	Values (Xi)		
Stab.4	5.00	0.000	ОК	Stab.1	16.00	0.750	ОК
Stab.5	6.06	1.060	ОК	Stab.2	15.25	0.000	OK
Stab.6	7.0 <mark>5</mark>	2.050	ОК	Stab.3	13.38	1.870	ОК
Stab.7	6.17	1.170	ОК	Stab.4	14.25	1.000	OK
Stab.8	4.50	0.500	ОК	Stab.5	18.25		
Stab.9	4.04	0.500	ОК	Stab.5	10.2	3.000	OK
Stab.10	4.00	1.000	OK	IV	ledian (M):	15.250	
				N	AD:	1.000	
	` '	5.000		5.	2 X MAD	5.200	
MAI		0.740					
5.2	X MAD	3.848					
20 —							
20							

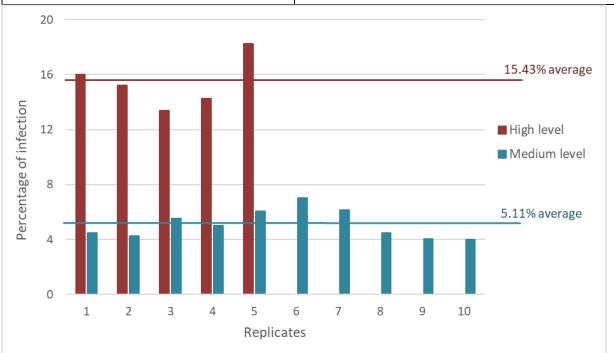


Figure 3: Stability test results, repartition against the mean

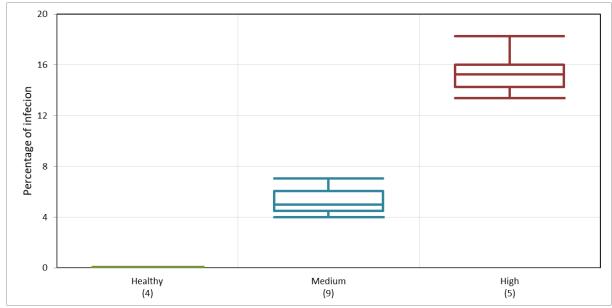


Figure 4: Boxplot representing results of the stability test

## **Conclusion of stability test:**

## Results of healthy level

All samples were negative, there were no false positive results.

#### Results of medium level

Samples are homogeneous, there was no outlier. The average obtained was very close to the expected percentage based on the homogeneity test.

## Results of high level

Samples are homogeneous, there was no outlier. The average obtained was close to the expected percentage based on the homogeneity test.

## Comparison of homogeneity and stability tests results

The comparison between homogeneity test results and stability test results are illustrated in the figure 5.

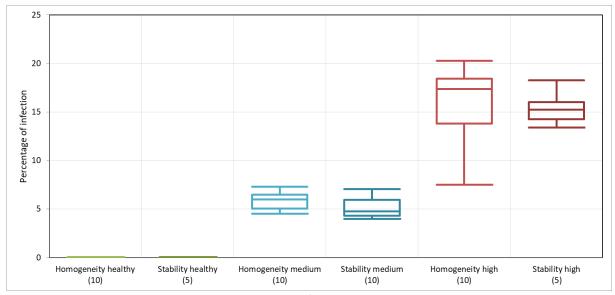


Figure 5: Boxplot representing results of the homogeneity and the stability tests

## **Conclusion on homogeneity and stability tests results:**

#### Results of healthy level

All samples were negative, there were no false positive results.

#### **Results of medium level**

Samples are homogeneous, there was no outlier. The values are between 4% and 7.29%.

#### Results of high level

Samples are homogeneous, there was no outlier. The values are between 7.5% and 20.25%.

#### **PROFICIENCY TEST RESULTS**

#### Qualitative results

#### Statistical tools

## □ Diagnostic sensitivity – specificity and accuracy

For homogeneous samples, the analysis was done by addition of the results of the 3 lots (healthy, medium and high level) according to the Standard NF EN ISO 16140 for qualitative results.

This norm gives us performance assessment criteria on diagnostic sensitivity, diagnostic specificity and accuracy calculated as in table n°8 and table n°9.

Table 8: Evaluation criteria for sensitivity, specificity and accuracy

	Expected result + (infected sample)	Expected result – (healthy sample)
Obtained result +	Positive agreement +/+ (PA)	Positive deviation -/+ (PD)
Obtained result -	Negative deviation +/- (ND)	Negative agreement -/- (NA)

Sensitivity: Percentage of samples correctly identified as positives.  $\Sigma PA/(\Sigma PA + \Sigma ND)x100$ .

Specificity: Percentage of samples correctly identified as being negative.  $\Sigma NA/(\Sigma NA + \Sigma PD) \times DA/(\Sigma NA + \Sigma PD) \times DA/(\Sigma NA + \Sigma PD)$ 

100.

Accuracy:  $(\Sigma NA + \Sigma PA)/(\Sigma PA + \Sigma NA + \Sigma PD + \Sigma ND) \times 100$ .

PA = positive agreement ND = negative deviation NA = negative agreement PD = positive deviation

Table 9: Conformity of results

, , , ,				
Performance criteria	Level to obtain			
Sensitivity	100%: all infected samples are positive; no false negative results have been obtained			
Specificity	100%: all healthy samples are negative; no false positive results have been obtained			
Accuracy	Synthesis of the two performance criteria. So, no false positive or negative results have been obtained			

The analysis of the results for a participating laboratory led to a declaration of conformity or non-conformity of the results in an individual sheet:

- "conform": obtained results correspond to expected results.
- "not conform": obtained results do not correspond to expected results.

## Statistical analysis of data

Raw data of all laboratories are given in appendix. All laboratories used a medium identification method (similar to the ISTA method 7-016).

## **Specificity and sensibility**

Analysis of results of healthy and high levels has been carried out according to the Norm NF EN ISO 16140 suitable to results expressed as positive / negative.

Results given in table n°10 and table n°11 present the criteria of performance.

Table 10: Overview of qualitative results for each laboratory on the three levels

Lab number	Healthy	Medium	High
01	0+/3	4+/4	3+/3
02	0+/3	4+/4	3+/3
03	0+/3	4+/4	3+/3
04	0+/3	4+/4	3+/3
05	0+/3	4+/4	3+/3
06	0+/3	4+/4	3+/3
07	0+/3	4+/4	3+/3
08	0+/3	4+/4	3+/3
09	0+/3	4+/4	3+/3
10	0+/3	4+/4	3+/3
11	0+/3	4+/4	3+/3
12	1+/3	4+/4	3+/3
13	1+/3	4+/4	3+/3
14	0+/3	4+/4	3+/3
15	0+/3	4+/4	3+/3
16	0+/3	4+/4	3+/3
17	0+/3	4+/4	3+/3

(Cells in grey correspond to lab results different from expected ones)

Table 11: Criteria of performance for each laboratory

Lab number	Specificity	Sensitivity	Accuracy
01	100%	100%	100%
02	100%	100%	100%
03	100%	100%	100%
04	100%	100%	100%
05	100%	100%	100%
06	100%	100%	100%
07	100%	100%	100%
08	100%	100%	100%
09	100%	100%	100%
10	100%	100%	100%
11	100%	100%	100%
12	67%	100%	83%
13	67%	100%	83%
14	100%	100%	100%
15	100%	100%	100%
16	100%	100%	100%
17	100%	100%	100%

(Cells in grey correspond to lab results different from expected ones)

15 out of 17 laboratories obtained 88.24% of specificity (no false positive) and 100% of sensitivity (no false negative).

#### Rating system

The calculation of the rating is done with the Excel file developed in collaboration with the Statistical committee of ISTA. It is based on an A, B, C and BMP rating. In this case:

- A corresponds to no false positive in healthy level and no false negative in medium and/or high level.
- BMP (Below Minimum Performance) corresponds to a not expected result. A false positive in healthy level or false negative in medium and/or high level, lead to BMP.

The calculation of the rating for each laboratory is presented in table n°12 and the distribution of the rating is presenting in figure 6.

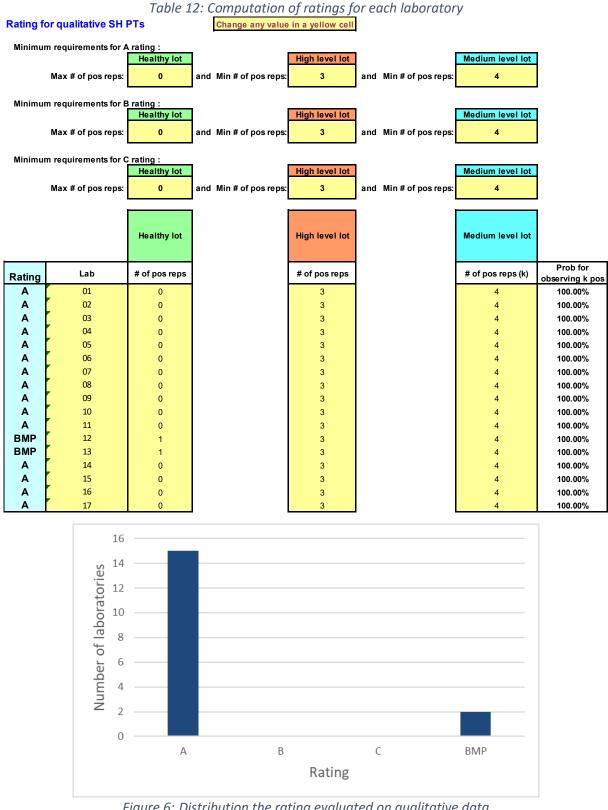


Figure 6: Distribution the rating evaluated on qualitative data

The distribution of the rating evaluated on qualitative data demonstrates that the note A represents 88.24% of the laboratories and the BMP rating represents 11.76%.

#### Quantitative results

#### Statistical tools

#### **⇒** Boxplot

Statistical analysis of results has been realized with the Boxplot tool. "Box plot" are graphical tools for visualizing key statistical measures. The aims are to give an idea of the center using to median, of variability and to identify the aberrant values. Values given by participants have been compared to values obtained during homogeneity and stability tests for medium and high levels.

## Statistical analysis of data

Raw data of all laboratories are given in appendix. All laboratories used a medium identification method (similar to the ISTA method 7-016).

The figure 7 indicates the mean of the 3 or 4 replicates obtained by the laboratories for the healthy, the medium and the high levels.

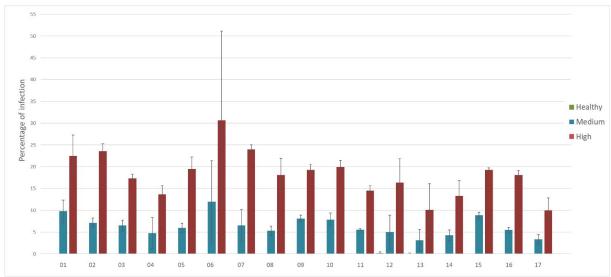


Figure 7: Graphic representing the average percentage of infection per sample obtained by the laboratories and in the stability and homogeneity tests.

Bars represent the mean and error bars represent the standard deviation.

#### **Conclusion of the laboratories:**

#### Results of healthy level

2 laboratories did not obtain the expected value (lab 12 and lab 13). They found false positive results.

#### Results of medium level

The results of the 4 replicates were compared to the results of the homogeneity and stability tests as presented in figure 8.

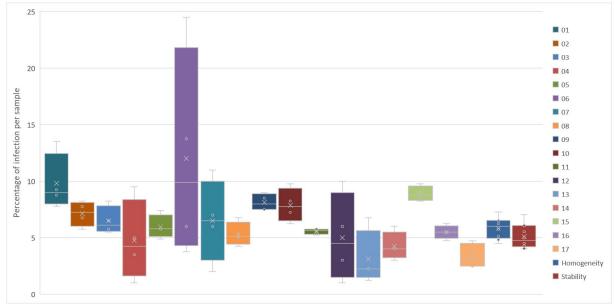


Figure 8: Comparison of results between laboratories, homogeneity and stability tests for the medium level

The figure 8 shows that the laboratory 06 has a value much higher (24.5%) than the homogeneity and stability test results and that the results of the laboratory 04 and the laboratory 12, with an 1% value, are lower than the results of the homogeneity and stability tests. The other laboratories have homogeneous results.

The Hampel's method was used to determine the outlier results. The result is given in table n°13.

Table 13: Participants' results for medium level evaluated by the Hampel's method

Lab | Values (Xi) | | Xi - M| | Status |

Lab	Values (Xi)	Xi - M	Status
Lab 01. 1 Lab 01. 2	8.75 9.25	2.625 3.125	OK OK
Lab 01. 2 Lab 01. 3	7.75	1.625	OK OK
Lab 01. 4	13.50	7.375	OK
Lab 02. 1	8.25	2.125	OK
Lab 02. 2	6.75	0.625	OK
Lab 02. 3	7.75	1.625	OK
Lab 02. 4	5.75	0.375	OK
Lab 03. 1 Lab 03. 2	5.50 5.75	0.625 0.375	OK OK
Lab 03. 2	6.50	0.375	OK OK
Lab 03. 4	8.25	2.125	OK
Lab 04. 1	5.00	1.125	OK
Lab 04. 2	9.50	3.375	OK
Lab 04. 3	3.50	2.625	OK
Lab 04. 4	1.00	5.125	OK
LAb 05. 1 LAb 05. 2	7.40 5.82	1.273 0.302	OK OK
LAb 05. 3	5.75	0.375	OK OK
LAb 05. 4	4.90	1.228	OK
Lab 06. 1	6.00	0.125	OK
Lab 06. 2	24.50	18.375	Outlier
Lab 06. 3	13.75	7.625	OK
Lab 06. 4	3.75	2.375 0.125	OK
Lab 07. 1	6.00	0.125	OK
Lab 07. 2 Lab 07. 3	7.00 11.00	4.875	OK OK
Lab 07. 3	2.00	4.125	OK OK
Lab 08. 1	5.25	0.875	OK
Lab 08. 2	5.00	1.125	OK
Lab 08. 3	6.75	0.625	OK
Lab 08. 4	4.25 9.00	1.875 2.875	OK OK
Lab 09. 1 Lab 09. 2	7.50	2.875 1.375	OK OK
Lab 09. 2	7.50	1.375	OK OK
Lab 09. 4	8.50	2.375	OK
Lab 10. 1	6.25	0.125	OK
Lab 10. 2	7.25	1.125	OK
Lab 10. 3	8.25	2.125	OK
Lab 10. 4	9.75	3.625	OK OK
Lab 11. 1 Lab 11. 2	5.25 5.50	0.875 0.625	OK OK
Lab 11. 2	5.75	0.375	OK OK
Lab 11. 4	5.75	0.375	OK
Lab 12. 1	6.00	0.125	OK
Lab 12. 2	10.00	3.875	OK
Lab 12. 3	3.00	3.125	OK
Lab 12. 4 Lab 13. 1	1.00 2.25	5.125	OK OK
Lab 13. 1	2.25	3.875 3.875	OK OK
Lab 13. 3	6.75	0.625	OK OK
Lab 13. 4	1.25	4.875	OK
Lab 14. 1	6.00	0.125	OK
Lab 14. 2	3.00	3.125	OK
Lab 14. 3	4.00	2.125	OK
Lab 14. 4 Lab 15. 1	4.00	2.125 3.625	OK OK
Lab 15. 1	9.75 8.25	2.125	OK OK
Lab 15. 2	9.00	2.875	OK OK
Lab 15. 4	8.50	2.375	OK
Lab 16. 1	5.50	0.625	OK
Lab 16. 2	6.25	0.125	OK
Lab 16. 3	4.75	1.375	OK
Lab 16. 4 Lab 17. 1	5.50 2.50	0.625 3.625	OK OK
Lab 17. 1	2.49	3.625	OK OK
Lab 17. 2	4.74	1.387	OK OK
Lab 17. 4	3.73	2.394	OK
Homog.1	6.35	0.225	OK
Homog.2	5.00	1.125	OK
Homog.3	7.29	1.165	OK
Homog.4 Homog.5	6.50 6.25	0.375 0.125	OK OK
Homog.6	4.82	1.305	OK OK
Homog.7	4.50	1.625	OK OK
Homog.8	6.58	0.455	OK
Homog.9	5.13	0.995	OK
Homog.10	5.76	0.365	OK
Stab.1	4.50	1.625	OK
Stab.2 Stab.3	4.26 5.53	1.865 0.595	OK OK
Stab.3 Stab.4	5.53	1.125	OK OK
Stab.5	6.06	0.065	OK
Stab.6	7.05	0.925	OK
Stab.7	6.17	0.045	OK
Stab.8	4.50	1.625	OK
Stab.9	4.04	2.085	OK
Stab.10	4.00	2.125	OK

Median (M): 6.125 MAD: 1.500 5.2 X MAD 7.800 The table n°13 shows an outlier value for the 2<sup>nd</sup> replicate of laboratory 06. The percentage obtained by this laboratory is much higher than the percentage obtained by the other laboratories for the medium level.

## Results of high level

The results of the 3 replicates were compared to the homogeneity and stability test results. The result of this comparison is given in figure 9.

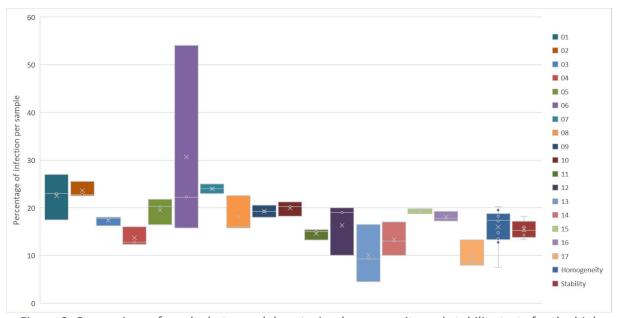


Figure 9: Comparison of results between laboratories, homogeneity and stability tests for the high level

The figure 9 shows that the laboratory 06 has a value much higher (54%) than the homogeneity and stability test results. The other laboratories have homogeneous results.

The Hampel's method was used to determine the outlier. The result is given in table n°14.

Table 14: Participants' results for high levels evaluated by the Hampel's method

		is jor nign iev	
Lab	Values (Xi)	Xi - M	Status
Lab 01. 1	27.00	8.875	OK
Lab 01. 2	17.50	0.625	OK
Lab 01. 3	23.00	4.875	OK .
Lab 02. 1	25.50	7.375	OK
		4.375	
Lab 02. 2	22.50		OK
Lab 02. 3	2 <mark>2.75</mark>	4.625	OK
Lab 03. 1	17.75	0.375	OK
Lab 03. 2	16.25	1.875	OK
Lab 03. 3	18.00	0.125	OK
Lab 04. 1	12.80	5.325	OK
Lab 04. 1	16.00	2.125	OK OK
Lab 04. 3	12.30	5.825	OK
LAb 05. 1	21.75	3.625	OK
LAb 05. 2	16.50	1.625	OK
LAb 05. 3	20.31	2.180	OK
Lab 06. 1	54.00	35.875	Outlier
Lab 06. 2	22.25	4.125	OK
Lab 06. 3	15.75	2.375	OK
Lab 07. 1	24.00	5.875	OK
Lab 07. 2	25.00	6.875	OK
Lab 07. 3	23.00	4.875	OK
Lab 08. 1	22.50	4.375	OK
Lab 08. 2	16.00	2.125	OK
Lab 08. 2	15.75	2.375	OK
		2.3/5	
Lab 09. 1	18.00	0.125	OK
Lab 09. 2	20.50	2.375	OK
Lab 09. 3	19.25	1.125	OK
Lab 10. 1	18.25	0.125	OK
Lab 10. 2	20.25	2.125	OK
Lab 10. 3	21.25	3.125	OK
		3.125	
Lab 11. 1	15.00		OK
Lab 11. 2	15.39	2.740	OK
Lab 11. 3	13.25	4.875	OK
Lab 12. 1	10.00	8.125	OK
Lab 12. 2	20.00	1.875	OK
Lab 12. 3	19.00	0.875	OK
Lab 13. 1	16.50	1.625	OK OK
Lab 13. 2	4.50	13.625	OK
Lab 13. 3	9.25	8.875	OK
Lab 14. 1	17.00	1.125	OK
Lab 14. 2	13.00	5.125	OK
Lab 14. 3	10.00	8.125	OK
Lab 15. 1	19.25	1.125	OK OK
Lab 15. 1	19.75	1.625	_
			OK
Lab 15. 3	18.75	0.625	OK
Lab 16. 1	17.75	0.375	OK
Lab 16. 2	19.25	1.125	OK
Lab 16. 3	17.25	0.875	OK
Lab 17. 1	13.22	4.908	OK
Lab 17. 2	7.96	10.165	OK
Lab 17. 2	8.73	9.397	OK
Homog.1	17.50	0.625	OK
Homog.2	12.75	5.375	OK
Homog.3	17.25	0.875	OK
Homog.4	18.25	0.125	OK
Homog.5	18.50	0.375	OK
Homog.6	7.50	10.625	OK
Homas 7	13.50	4.625	OK
Homog.7			
Homog.8	20.25	2.125	OK
Homog.9	14.75	3.375	OK
Homog.10	19.50	1.375	OK
Stab.1	16.00	2.125	OK
Stab.2	15.25	2.875	OK
Stab.3	13.38	4.745	OK OK
Stab.4	14.25	3.875	OK
Stab.5	18.25	0.125	OK

Median (M): MAD: 5.2 X MAD

18.125	
2.933	
15.249	

The table n°14 shows an outlier value for the 1<sup>st</sup> replicate of laboratory 06. The percentage obtained by this laboratory is much higher than the percentage obtained by the other laboratories for the high level.

## **Rating system**

The calculation of the rating is done with the Excel file developed in collaboration with the Statistical committee of ISTA. It is based on an A, B, C and BMP rating. This calculation uses median values of all labs without consider laboratories with outlier result. Outlier result, per level of contamination, are identified using Hampel's tool. Results of labs will be analyzed using Z scores and automatically calculated by an Excel sheet.

The results are presented in table n°15 and the distribution of the notes are presenting in figure 10.

Table 15: Computation of ratings for each laboratory

#### **Rating for quantitative SH PTs**

	Change any value in a yellow cell						
	Max abs(z-score) for the Healthy lot  Max abs(z-score) for the Medium lot  Max abs(z-score) for the High lot						
	0.67	and	0.67	and	0	Minimum requirements for A rating	
	1.5	and	1.5	and	0	Minimum requirements for B rating	
	2.33	and	2.33	and	0	Minimum requirements for C rating	
	abs(z-score)	ĺ	abs(z-score)		abs(z-score)		
z-score	for the High lot		for the Medium lot		for the Healthy lot	Lab	
С	0.90		2.10		0.00	01	
В	1.12		0.61		0.00	02	
Α	0.13		0.26		0.00	03	
В	0.85		0.71		0.00	04	
Α	0.31		0.04		0.00	05	
BMP	2.54		3.31		0.00	06	
В	1.20		0.26		0.00	07	
Α	0.02		0.40		0.00	08	
В	0.25		1.16		0.00	09	
В	0.39		1.02		0.00	10	
В	0.69		0.26		0.00	11	
ВМР	0.33		0.57		0.17	12	
ВМР	1.58		1.61		0.08	13	
В	0.93		0.99		0.00	14	
С	0.25		1.58		0.00	15	
Α	0.02		0.30		0.00	16	
С	1.60		1.48		0.00	17	

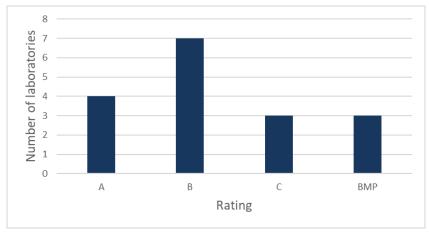


Figure 10: Distribution the rating evaluated on quantitative data

#### **CONCLUSIONS**

For this proficiency test, for the qualitative results, the A rating represents 88.24% of the laboratories and the BMP rating represents 11.76%. For the quantitative results, the A rating represents 23.52% (4 out of 17) of the laboratories, the B rating represents 41.18% (7 out of 17) of the laboratories, the C rating represents 17.65% (3 out of 17) of the laboratories and the BMP rating represents 17.65% (3 out of 17) of the laboratories.

The final rating was defined according to the qualitative and quantitative results obtained by each laboratory. The lowest rating between the two results was retained for the final rating. The number of laboratories according to the final ratings are shown in the figure 11. The A rating represents 23.52% (4 out of 17) of the laboratories, the B rating represents 41.18% (7 out of 17) of the laboratories, the C rating represents 17.65% (3 out of 17) of the laboratories and the BMP rating represents 17.65% (3 out of 17) of the laboratories.



Figure 11: Distribution of laboratories according to the final rating

- All lots used in this proficiency test were tested by a detection and identification method on media (similar to the ISTA method 7-016) by the participating laboratories.
- A point of vigilance should be considered for the laboratories that had false positive in healthy level (lab 12 and lab 13) and for the laboratory that had two outliers (lab 06).

## **Appendix:**

Lab number	Level of contamination	Number of samples	Quantitative result	Qualitative result
		144	0.000%	Neg
	Healthy	84	0.000%	Neg
		242	0.000%	Neg
		325	8.750%	Pos
01	N. d. a. dii	289	9.250%	Pos
01	Medium	248	7.750%	Pos
		78	13.500%	Pos
		58	27.000%	Pos
	High	326	17.500%	Pos
		273	23.000%	Pos
		4	0.000%	Neg
	Healthy	37	0.000%	Neg
		318	0.000%	Neg
		139	8.250%	Pos
02	Medium	147	6.750%	Pos
02	ivieurum	197	7.750%	Pos
		259	5.750%	Pos
	High	250	25.500%	Pos
		309	22.500%	Pos
		9	22.750%	Pos
		160	0.000%	Neg
	Healthy	255	0.000%	Neg
		287	0.000%	Neg
		143	5.500%	Pos
03	Medium	102	5.750%	Pos
03		161	6.500%	Pos
		45	8.250%	Pos
		151	17.750%	Pos
	High	182	16.250%	Pos
		261	18.000%	Pos
		40	0.000%	Neg
	Healthy	18	0.000%	Neg
		249	0.000%	Neg
		74	5.000%	Pos
04	Medium	11	9.500%	Pos
<del>-</del>	Wediam	131	3.500%	Pos
		170	1.000%	Pos
		96	12.800%	Pos
	High	133	16.000%	Pos
		140	12.300%	Pos

Lab number	Level of contamination	Number of samples	Quantitative result	Qualitative result
	Healthy	154	0.000%	Neg
		213	0.000%	Neg
		135	0.000%	Neg
		206	7.398%	Pos
		62	5.823%	Pos
05	Medium	302	5.750%	Pos
		71	4.897%	Pos
		103	21.750%	Pos
	High	235	16.500%	Pos
	o o	123	20.305%	Pos
		278	0.000%	Neg
	Healthy	340	0.000%	Neg
	,	2	0.000%	Neg
		61	6.000%	Pos
		53	24.500%	Pos
06	Medium	316	13.750%	Pos
		49	3.750%	Pos
		271	54.000%	Pos
	High	44	22.250%	Pos
	111611	190	15.750%	Pos
		223	0.000%	
	Healthy	240	0.000%	Neg
	пеанну	198		Neg
	Medium		0.000%	Neg
		301	6.000%	Pos
07		224	7.000%	Pos
		231	11.000%	Pos
		283	2.000%	Pos
	11: -1-	39	24.000%	Pos
	High	337	25.000%	Pos
		22	23.000%	Pos
	tr. dub	60	0.000%	Neg
	Healthy	8	0.000%	Neg
		64	0.000%	Neg
		13	5.250%	Pos
08	Medium	257	5.000%	Pos
		208	6.750%	Pos
		260	4.250%	Pos
		130	22.500%	Pos
	High	112	16.000%	Pos
		113	15.750%	Pos
		196	0.000%	Neg
	Healthy	313	0.000%	Neg
		251	0.000%	Neg
		138	9.000%	Pos
09	Medium	225	7.500%	Pos
		119	7.500%	Pos
		184	8.500%	Pos
		116	18.000%	Pos
	High	69	20.500%	Pos
	9	220	19.250%	Pos

Lab number	Level of contamination	Number of samples	Quantitative result	Qualitative result
	Healthy	200	0.000%	Neg
		219	0.000%	Neg
		67	0.000%	Neg
		155	6.250%	Pos
10	Madium	101	7.250%	Pos
10	Medium	127	8.250%	Pos
		89	9.750%	Pos
		38	18.250%	Pos
	High	48	20.250%	Pos
		134	21.250%	Pos
		152	0.000%	Neg
	Healthy	159	0.000%	Neg
		209	0.000%	Neg
		142	5.250%	Pos
11	Madium	70	5.500%	Pos
11	Medium	99	5.750%	Pos
		280	5.750%	Pos
	High	90	15.000%	Pos
		321	15.385%	Pos
		179	13.250%	Pos
		222	0.000%	Neg
	Healthy	188	0.500%	Pos
		239	0.000%	Neg
		108	6.000%	Pos
12	Medium	169	10.000%	Pos
12	iviedium	93	3.000%	Pos
		16	1.000%	Pos
		81	10.000%	Pos
	High	36	20.000%	Pos
		210	19.000%	Pos
		284	0.000%	Neg
	Healthy	178	0.250%	Pos
		106	0.000%	Neg
		253	2.250%	Pos
13	Medium	285	2.250%	Pos
15	Wicarum	94	6.750%	Pos
		339	1.250%	Pos
		105	16.500%	Pos
	High	336	4.500%	Pos
		176	9.250%	Pos

Lab number	Level of contamination	Number of samples	Quantitative result	Qualitative result
		33	0.000%	Neg
	Healthy	279	0.000%	Neg
		30	0.000%	Neg
		82	6.000%	Pos
14	Medium	236	3.000%	Pos
14	ivieurum	191	4.000%	Pos
		158	4.000%	Pos
		12	17.000%	Pos
	High	111	13.000%	Pos
		272	10.000%	Pos
		175	0.000%	Neg
	Healthy	332	0.000%	Neg
		334	0.000%	Neg
		293	9.750%	Pos
15	Madium	265	8.250%	Pos
15	Medium	51	9.000%	Pos
		328	8.500%	Pos
	High	117	19.250%	Pos
		204	19.750%	Pos
		55	18.750%	Pos
		501	0.000%	Neg
	Healthy	635	0.000%	Neg
		462	0.000%	Neg
		453	5.500%	Pos
16	Medium	547	6.250%	Pos
10	ivieurum	470	4.750%	Pos
		468	5.500%	Pos
		555	17.750%	Pos
	High	493	19.250%	Pos
		604	17.250%	Pos
		174	0.000%	Neg
	Healthy	241	0.000%	Neg
		54	0.000%	Neg
		304	2.500%	Pos
17	Medium	77	2.488%	Pos
1/	ivieututti	237	4.738%	Pos
		65	3.731%	Pos
		120	13.217%	Pos
	High	167	7.960%	Pos
		5	8.728%	Pos