



**European Union Reference Laboratory for
Parasites**



Istituto Superiore di Sanità

**Reports on the Proficiency Tests for *Trichinella* spp.
organised by the EURLP for the NRLs**

March, 2012

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A Report of the NRL Proficiency Test to detect *Trichinella spiralis* larvae in pork and/or horse meat samples according to the EU directive 2075/2005

1 Introduction

Nematode worms of the genus *Trichinella* are zoonotic parasites circulating in most of the European countries in both wild and domestic animals (Pozio et al., 2009). Humans acquire the infection by the consumption of raw or undercooked meat from pigs, horses, wild boars and other game animals (Pozio et al., 2003). According to the Commission Regulation (EC) No 2075/2005, all animals which are potential carrier of *Trichinella* spp. infective larvae should be tested at the slaughterhouse according to one of the approved test.

2 Scope

One of the core duties of the EURL for Parasites is to organise proficiency tests, as it is stated in the Regulation (EC) No 882/2004 of the European Parliament and of the Council (Commission Regulation No. 882/2004). The scope of this comparison is to test the competence of the appointed NRLs to detect *Trichinella* larvae in pork and/or horse meat samples according to one of the approved methods reported in the Annex 1 of the Commission Regulation EC No 2075/2005.

3 Proficiency test organization

For the first time, the proficiency test (PT) registration, result submission, and reporting, have been organized online on a dedicated section of the EURLP web site (<http://www.iss.it/crlp/test/index.php?lang=2&tipo=28>).

4 Time frame

The PT was announced to NRLs by email on 25 January, 2012 and the dead line to make the online registration was 17 February, 2012. On March 19, 2012, the samples were dispatched to participants by an international courier. Reporting deadline was 30 March, 2012.

5 Test material

The test material forwarded to each laboratory, consisted of 5 meat balls made with 100 ± 5 grams (or 35 ± 2 grams) of minced pork or 5 meat balls made with 100 ± 5 grams (or 35 ± 2 grams) of minced horse meat. Four meat balls contained in the core a known number of viable *Trichinella spiralis* larvae; whereas, one meat ball, which did not contain any larva, was the negative control. To evaluate the NRL competence and skill and the sensitivity of the digestion method in each participating laboratory, the number of larvae spiked into the meat balls was of three different sizes: 3, 5 or 8.

Larvae were obtained by a partial artificial digestion of *T. spiralis*-infected mice (isolate code ISS003), according to the EURLP Guidelines. Larvae were counted under a stereo-microscope using a watch glass of 2 cm of diameter and transferred to the meatball rinsing the watch glass with PBS. To ensure that no larva remained on the glass, it was examined twice under the stereo-microscope and rinsed with PBS allowing the PBS to reach the core of the ball (see Annex 2).

Each meat ball was close in a plastic bag under vacuum, a code was added on the envelop and the same code with the number of larvae was reported in an Excel file. Each envelope containing the meat ball was then stored at $+4^{\circ}\text{C}$ until the forwarding.

The 5 meat balls under vacuum were forwarded in a polystyrene box containing ice boxes. In the package, ice boxes were separated from the meat samples by a cardboard separator to avoid a direct contact between meat samples and ice boxes. The packages were forwarded according to the international forwarding regulations by an international courier.

To check the sample stability over time, and to estimate the suitability of the packing and forwarding conditions under which the meat balls were sent, two groups of 5 meat balls each, were packaged as those that were forwarded, stored at room temperature, and tested at the EURLP three and five days after packaging.

Each participant lab was invited to add the information about package content and its condition of preservation in the online “shipment form”.

6 Instructions to participants

The instructions were available to all the participants at the “instruction” page of the web site.

7 Statistical analysis

The analysis of data was performed using the STATA software; the statistical significance of the results was confirmed by the Kruskal-Wallis test.

8 Z-score

A Z-score was tentatively established to evaluate the laboratory performance. The evaluation parameters were: 1) at least 70% of the larvae present in the sample should be recovered for samples containing more than 9 larvae; 2) 50% of larvae present in the sample should be recovered for samples containing from 4 to 9 larvae; and 3) at least one larva should be detected in samples containing 1-3 larvae, but no Z-score value can be applied in this case. For a Z-score value ≤ 3.0 , the result is positive, for a Z-score value $2.0 < Z \leq 3.0$, there is a warning signal for the technician who performed the test; and for Z value >3.0 , the result is negative and there is an action signal for the technician who performed the test.

9 Participating laboratories

Of the 27 MS, Luxembourg appointed the Belgium NRL for parasites, all the other NRLs participated at the PT. In addition, NRLs from Norway, Serbia and Switzerland agreed to participate. Consequently, a total of 29 labs were involved.

10 Results

10.1 Delivery of the package to NRLs

Out of the 29 packages, 10 were delivered within 24 h, 15 within 48 h, three within 72 h, and one was the NRL of Italy which is located in the same Institute of the EURLP. At the delivery, the internal temperature of 26 packages was less than 10°C, whereas the internal temperature was higher in three packages. The time elapsed from the arrival of the package at the NRL and the control of its content was within 1 h for 25 NRLs, of 4 h for 1 NRLs, and between 12 and 18 h for 3 NRLs; in these three last labs, the parcel was stored in a refrigerator at +4°C.

10.2 Digestion methods

The magnetic stirred method for the pooled sample digestion was the most used (21 NRLs, 72.4%); the mechanically assisted pooled sample digestion method/sedimentation technique was

used in 5 NRLs (17.2%) and the automatic digestion method for pooled samples up to 35 g was used in 3 NRLs (10.3%).

10.3 Type of meat samples

Out of the 29 participating laboratories, 24 labs (82.7%) requested pork samples of 100 ± 5 g, 3 labs (10.7%) pork samples of 35 ± 2 g and 2 labs (6.9%) horse meat samples of 100 ± 5 g.

10.4 Descriptive statistic of the results

The descriptive statistic of the difference between expected and reported counts by laboratory, shows a statistically significant variation among laboratories on the difference between expected and reported count according to the Kruskal-Wallis test ($p=0.0073$) (Annex 4). The descriptive statistic of the difference between the expected and reported larval count by laboratory and the comparison among the 2007, 2008, 2009, 2010, 2011 and 2012 results are shown in the Annex 5. The graphical representation by histograms of the absolute difference between expected and reported count by laboratory is shown in the Annex 6. All labs show an improvement in the digestion test from 2007 to 2008, while labs show fluctuating results from 2008 to 2012. The comparison of the overall mean difference (i.e. the mean of relative difference values considering all samples and all laboratories) confirm that a strong improvement was made from the first (2007) to the second (2008) PT, while the improvement is less evident in the further years (Table 1). In 2012, the overall mean difference value is a little bit higher than that of the last previous three years.

Table 1 - Overall mean difference comparison for the 2007-2012 period

Year	2007	2008	2009	2010	2011	2012
Relative difference values	0.41	0.25	0.23	0.22	0.20	0.26

10.5 Not consistent results

Two NRLs (6.9%) found one false negative each (Table 2). One lab found one false positive. The two samples which were stated as negative, had been spiked with 3 and 5 larvae (Table 2). Four laboratories showed an overestimation of the larvae in the sample: three labs an overestimation of one larva; and one lab an overestimation of three larvae.

Table 2 - Number of larvae in the sample considered to be negative by NRLs

Laboratory code	15	16
No. of larvae in the samples considered negative	5	3

10.6 Result evaluation

The result of each sample was evaluated by applying the Z-score when relevant. Out of the 29 labs participating to the PT, 10 labs (34.5%) obtained a positive evaluation for all the samples; 8 labs (27.6%) obtained one or more warning signals, but no negative results; and 11 labs (38%) obtained a negative evaluation for one or two samples.

11 Conclusions

The comparison of the 2007-2012 results shows that there has been an improvement in the laboratory performances from the first (2007) to the second (2008) PT, while during the following years the improvement was less evident. The presence of laboratories reporting a number of larvae greater than the number of larvae spiked into the sample could suggest: 1) a problem in counting the larvae



present in the sediment; and 2) cross contamination from one to the next digestion test. One false positive was detected, whereas 2 (6.9%) laboratories found false negatives (last year the value was 14.3%). The use of the Z-score seems to be a suitable method to evaluate the lab performance in the course of PTs. However according to the EC Regulation 2075/2005, technicians performing the test must be able to identify positive samples as positive, independently to the amount of larvae contained in the sample.

11 References

Pozio E., Rinaldi L., Marucci G., Musella V., Galati F., Cringoli G., Boireau P., La Rosa G. (2009) Hosts and habitats of *Trichinella spiralis* and *Trichinella britovi* in Europe. *Int. J. Parasitol.* 39:71-79.

Pozio E., Gomez Morales M.A., Dupouy Camet J. (2003). Clinical aspects, diagnosis and treatment of trichinellosis. *Expert Review of Anti-infective Therapy* 1:471-482.

Commission Regulation (EC) No 2075/2005 of 5 December 2005 laying down specific rules on official controls for *Trichinella* in meat. OJ L338/60-82.

Commission Regulation (EC) No. 882/2004 of the European Parliament and of the Council of 29 April 2004 on official controls performed to ensure the verification of compliance with feed and food law, animal health and animal welfare rules. OJ L191/1-59.

Community Reference Laboratory for Parasites (2007). Guideline for the detection of *Trichinella* larvae at the slaughterhouse or connected laboratory in a Quality Assurance System, pp. 1-14, www.iss.it/crlp/docu/.

B Report of the NRL Proficiency Tests on *Trichinella* spp. larva identification at the species level by a molecular method

1 Introduction

In the MS of the European Union, four species of the genus *Trichinella* are circulating among domestic and/or wild animals: *Trichinella spiralis*, *Trichinella nativa*, *Trichinella britovi* and *Trichinella pseudospiralis*. *Trichinella britovi* is the most widespread species infecting prevalently carnivore mammals but also wild boars; whereas, it is seldom detected in domestic pigs. This species is circulating in most of MS excluding Cyprus, Denmark, Ireland, Luxembourg, Malta and UK (Pozio et al., 2009). *Trichinella spiralis* infecting prevalently domestic pigs and wild boars is more scattered in single foci present in Austria, Bulgaria, Finland, Germany, Hungary, Lithuania, Poland, Romania, Ireland and North Ireland (UK), Slovak Rep., Spain and Sweden (Pozio et al., 2009). *Trichinella nativa* is circulating among carnivore mammals living in cold regions (Estonia, Finland, Latvia, Lithuania and Sweden) but recently it has been detected in one red fox of Poland and in two red foxes of Germany. *Trichinella pseudospiralis* is the only species infecting both mammals and birds (Pozio et al., 2009). The main reservoir host is the wild boars; seldom this species has been detected in carnivore mammals (foxes and raccoon dogs). This species has been detected in 12 MS of the EU (Bulgaria, Czech Rep., Denmark, Estonia, Finland, France, Germany, Hungary, Italy, Netherlands, Slovak Rep. and Sweden). The identification of the *Trichinella* species is extremely important for the epidemiological investigation and to trace back the origin of infections in animals and humans. These parasites can be distinguished among them only by molecular methods (Pozio and La Rosa, 2010). Several protocols are today available in the literature, but only few of them allow to identify single organisms (www.iss.it/binary/crlp/cont/PCR_method_WEB_SITE.pdf).

2 Scope

According to of the Commission Regulation (EC) No 2075/2005 (Annex 1, Chapter I), when *Trichinella* spp. larvae are detected by digestion of animal muscles, the larvae are to be kept in 90% ethyl alcohol for preservation and identification at the species level at the EURLP or NRLs. During the NRL Workshop of 2010, participants requested the EURLP to organise a PT on the identification of *Trichinella* spp. larvae at the species level. The scope of this PT is to test the competence of NRLs to correctly identify the species circulating in Europe.

3 Time frame

The proficiency test (PT) was announced to NRLs by email on 19 January, 2012 and the dead line to send the participation form was 25 February, 2012. On March 19, 2012, the samples were dispatched to participants by an international courier. Reporting deadline was 6 April, 2012.

4 Participating laboratories

Of the 29 MS, only 13 MS agreed to participate at the PT (Austria, Belgium, Bulgaria, Estonia, Finland, France, Germany, Greece, Italy, Netherland, Slovak Rep., Slovenia and Spain). In addition, NRLs from Norway, Serbia and Switzerland agreed to participate (see Annex 1).

5 Test material

The test material forwarded to each laboratory consisted of *Trichinella* muscle larvae in ethanol. According to the lab requests, two different type of samples were forwarded: 1. pool of larvae and 2. single larvae. The pool of larva panel consisted of four vials, each vial containing 20 larvae of the species: *T. spiralis*, *T. britovi*, *T. papuae* and *T. pseudospiralis*. The single larva panel consisted of



12 vials containing one single larva each (i.e., three vials for each of the four species). Larvae were obtained by artificial digestion of mouse carcasses infected with *T. spiralis* (isolate code ISS003), *T. papuae* (isolate code ISS572), *T. britovi* (isolate code ISS002), and *T. pseudospiralis* (isolate code ISS013), according to the EURLP Guidelines. Participating labs were invited to use one of the published methods available in the international literature. Even if *T. papuae* is not circulating in Europe, larvae of this species were included in the PT to test the skill of the NRLs to identify an imported *Trichinella* species.

6 Results

6.1 Type of samples

Out of 16 participating laboratories, 10 labs (62.5%) requested pool of larvae, 6 labs (37.5%) requested single larvae.

6.2 Methods of analysis

Out of 16 labs which received the PT samples, only 14 (87.5%) sent the results. Of them, five labs followed the EURLP protocol, four labs followed the protocol published by Pozio and La Rosa (2010), one lab followed the protocol published by Zarlenga et al. (1999), two labs followed the protocol published by De Bruyne et al. (2005), one lab used a in house method based on Zarlenga et al. (1999) and three labs did not communicate this information.

6.3 Molecular identification

6.3.1 PCR products

All the ten labs which analyzed pool of larvae, were able to obtain PCR products. Out of the six labs which analyzed single larvae, only two failed to obtain PCR products from a number of samples between 1 and 3. Two labs did not send their results.

6.3.2 Species identification

Independently of the type of sample (single or pool of larvae), three labs failed to correctly identify some samples at the species level due to the lack of experience and/or the use of a non-appropriate protocol (Annex 8).

7 Conclusions

Out of the 26 NRLs for parasites of EU, 16 (62%) agreed to participate at the PT, 14 (54%) sent the results, but only 7 (27%) were able to correctly identify all the larvae at the species level; of them, 4 labs analyzed pools of larvae and three labs single larvae. Among the three non-EU labs, only one correctly identified all the samples. The failure to obtain PCR products and the mistakes in *Trichinella* spp. larva identification were mainly due to the use of not suitable diagnostic tools or protocols and to the lack of an appropriate training (mainly for the single larva analysis).

8 References

De Bruyne A, Yera H, Le Guerhier F, Boireau P, Dupouy-Camet J, 2005. Simple species identification of *Trichinella* isolates by amplification and sequencing of the 5S ribosomal DNA intergenic spacer region. *Vet Parasitol.* 132, 57-61.

Pozio E, Hoberg E, La Rosa G, Zarlenga DS, 2009. Molecular taxonomy, phylogeny and biogeography of nematodes belonging to the *Trichinella* genus. *Infect Genet Evol.* 9, 606-616.



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Pozio E, La Rosa G, 2010. *Trichinella*. In: Dongyou, L. (ed.), Molecular detection of foodborne pathogens. Taylor and Francis, CRC Press, Boca Raton, London, New York, pp. 851-863.

Zarlenga DS, Chute MB, Martin A, Kapel CM, 1999. A multiplex PCR for unequivocal differentiation of all encapsulated and non-encapsulated genotypes of *Trichinella*. *Int J Parasitol.* 29, 1859-1867.



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Annex 1

National Reference Laboratories (NRL) participating at the proficiency tests for *Trichinella*

NRL for parasites	Country	PT digestion	PT PCR
Institut für Veterinärmedizin, Innsbruck	Austria	yes	yes
Institute of Tropical Medicine, Antwerp	Belgium	yes	yes
National Diagnostic and Research Veterinary Institute, Sofia	Bulgaria	yes	yes
State Veterinary Laboratory, Nicosia	Cyprus	yes	no
State Veterinary Institute, Olomouc	Czech Rep	yes	no
National Veterinary Institute, Copenhagen	Denmark	yes	no
Estonian Veterinary and Food Laboratory, Tartu	Estonia	yes	yes
Research Unit Finnish Food Safety Authority, Evira, Oulu	Finland	yes	yes
Food borne parasite NRL & UMR BIPAR ANSES, ENVA, UPEC, ANSES-Laboratoire de Santé Animale, Maison Alfort	France	yes	yes
Federal Institute for Risk Assessment, BfR, Berlin	Germany	yes	yes
Centre of Athens Veterinary Institutions, Athens	Greece	yes	yes
Central Veterinary Institute, Budapest	Hungary	yes	no
Central Meat Control Laboratory, Celbridge, County Kildare, Istituto Superiore di Sanità, Rome	Ireland Italy	yes yes	no yes
Laboratory of Food and Environmental Investigations, Riga	Latvia	yes	no
National Food and Veterinary Risk Assessment Institute, Vilnius	Lithuania	yes	no
National Veterinary Laboratory, Alberttown, Marsa	Malta	yes	no
National Institute of Public Health and the Environment, Bilthoven	Netherlands	yes	yes
National Veterinary Research Institute, Pulawy	Poland	yes	no
Laboratório Nacional de Investigação Veterinária, Lisboa	Portugal	yes	no
Hygiene and Public Veterinary Health Institute, Bucharest	Romania	yes	no
State Veterinary and Food Institute, Bratislava	Slovak Rep.	yes	yes
National Veterinary Institute, Ljubljana	Slovenia	yes	yes
Centro Nacional de Alimentación, Agencia Española de Seguridad Alimentaria y Nutrición, Majadahonda	Spain	yes	yes
Statens Veterinärmedicinska Anstalt, Uppsala	Sweden	yes	no
Veterinary Laboratories Agency, Weybridge	UK	yes	no
Norwegian Veterinary institute, Oslo	Norway	yes	yes
Institute for Laboratory Diagnostic INEP, Belgrade	Serbia	yes	yes
Institute of Parasitology, University of Bern, Bern	Switzerland	yes	yes

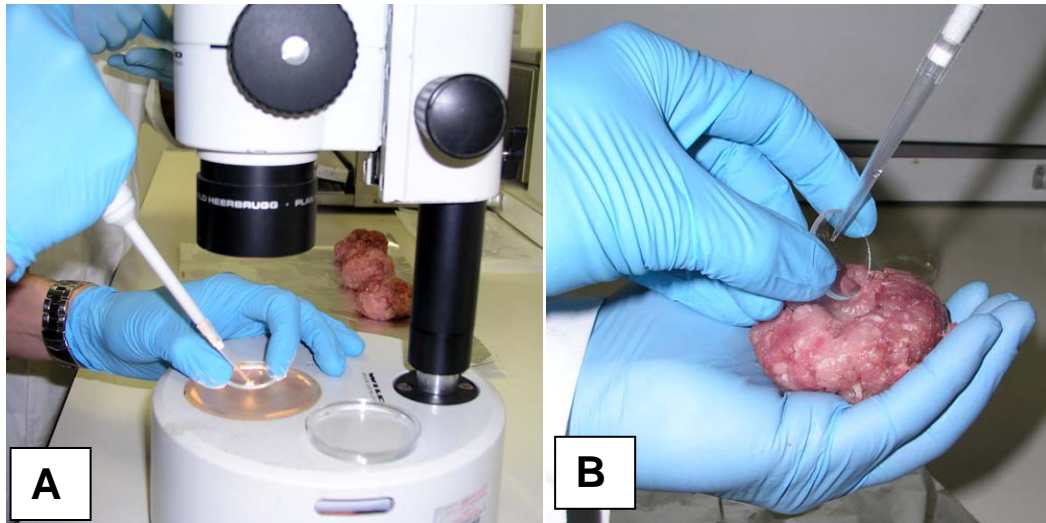
Annex 2



Meatball samples (35g and 100g)



**An hollow is made in the center of each meatball to house the
larvae**



Larvae are counted under a stereo-microscope using a watch glass (A) and transferred to the meatball rinsing the watch glass with 200 μ l of PBS (B)



The watch glass is examined under the stereo-microscope to ensure that no larva remains on it



Each sample is under vacuum sealed (A) and labeled with a numeric code (B)

Annex 3

Descriptive statistics of the absolute difference between expected and reported count by laboratory for all samples (horse meat and pork) in 2012

Laboratory code	No. of samples	Mean	SD
1	4	2.25	2.21
2	4	0.5	0.57
3	4	1.75	0.5
4	4	0.5	0.57
5	4	1.25	1.5
6	4	1.25	1.89
7	4	0	0
8	4	0.25	0.5
9	4	1.25	1.2
10	4	0.75	0.95
11	4	1.75	0.95
12	4	3	2
13	4	0.5	0.57
14	4	0.75	0.5
15	4	3	1.41
16	4	3	0.81
17	4	0.75	1.5
18	4	1.75	1.7
19	4	0.25	0.5
20	4	0.25	0.5
21	4	0.25	0.5
22	4	0.75	0.95
23	4	2.25	0.95
24	4	1.25	0.95
25	4	2.25	1.5
26	4	1.75	1.5
34	4	1.75	0.95
35	4	3	1.15
40	4	0.75	0.95
All	116	1.33	1.37

Kruskal-Wallis test, $p = 0.0073$

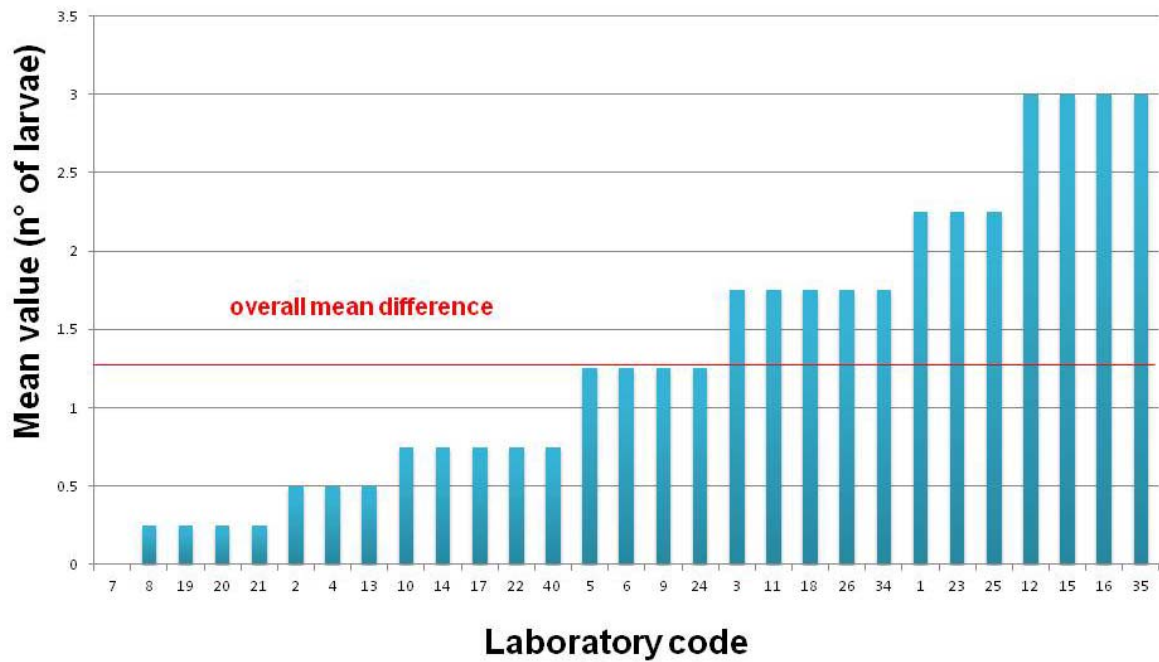
Annex 4

Descriptive statistic of the difference between the expected and reported larval count by laboratory and comparison among the years 2007, 2008, 2009, 2010, 2011 and 2012.

Laboratory code	Mean					
	2007	2008	2009	2010	2011	2012
1	0.36	0.24	0.11	0.25	0.23	0.35
2	0.14	0.22	0.07	0.08	0.09	0.08
3	0.52	0.37	0.23	0.10	0.29	0.34
4	0.20	0.12	0.03	0.12	0.13	0.13
5	0.14	0.06	0.08	0.12	0.09	0.26
6	0.30	0.08	0.17	0.10	0.04	0.17
7	0.23	0.05	0.03	0.07	0.04	0
8	0.13	0.08	0.03	0.12	0.02	0.05
9	0.13	0.19	0.06	0.11	0.18	0.23
10	0.17	0.45	0.28	0.41	0.15	0.14
11	0.33	0.22	0.24	0.23	0.16	0.36
12	0.72	0.10	0.10	0.26	0.18	0.5
13	1	0.36	0.40	0.25	0.14	0.1
14	0.75	0.12	0.20	0.12	0.18	0.16
15	0.64	0.64	0.42	0.25	0.42	0.63
16	0.41	0.33	0.13	0.23	0.33	0.62
17	0.86	0.26	0.68	0.05	0.14	0.09
18	0.26	0.27	0.45	0.37	0.53	0.34
19	0.08	0.12	0.03	0.16	0.08	0.08
20	-	0.19	0.08	0.01	0.07	0.03
21	0.40	0.25	0.02	0.51	0.16	0.03
22	0.36	0.41	0.30	0.23	0.15	0.18
23	0.48	0.38	0.17	0.08	0.26	0.43
24	0.45	0.27	0.37	0.36	0.26	0.24
25	0.70	0.43	0.75	0.26	0.50	0.46
26	-	-	-	0.15	0.11	0.33
34	-	-	0.23	0.54	0.23	0.34
35	-	-	-	0.44	0.36	0.59
40	-	-	-	-	-	0.15

Annex 5

Graphical representation by histograms of the absolute difference between expected and reported count of all the samples (horse meat and pork), by laboratory





Annex 6

Form 6A Laboratory _____

Proficiency test to identify Trichinella larvae at the species level by a molecular method

Identification of pool of larvae

Results

Table with 5 columns: Sample code, Molecular Method, Date (dd/mm), Identified species, Notes

1) Explain briefly or add a reference for the molecular method used.

Date _____

Technician (name/sign): _____

Form 6B Laboratory _____

Proficiency test to identify Trichinella larvae at the species level by a molecular method

Identification of single larvae

Results

Table with 5 columns: Sample code, Molecular Method, Date (dd/mm), Identified species, Notes

1) Explain briefly or add a reference for the molecular method used.

Date _____

Technician (name/sign): _____

Annex 7

**PT on *Trichinella* spp. larva identification at the species level by a molecular method.
Labs coded 2 and 25 did not send any result.**

Lab code	Sample type	Correct identification	Incorrect identification	Not identified	Molecular method
1	single larvae	12	-	-	EURLP
2	pool	no result provided	no result provided	no result provided	no information provided
4	single larvae	12	-	-	EURLP
6	pool	4	-	-	De Brunye et al., 2005
7	pool	3	1	-	EURLP
8	pool	4	-	-	Pozio and La Rosa, 2010
10	single larvae	12	-	-	EURLP and Zarlenga et al., 1999
11	pool	3	1	-	no information provided
16	pool	4	-	-	Pozio and La Rosa, 2010
21	pool	3	-	1*	Pozio and La Rosa, 2010
22	single larvae	11	-	1**	EURLP
23	pool	4	-	-	EURLP
25	pool	no result provided	no result provided	no result provided	no information provided
34	single larvae	7	2	3**	Pozio and La Rosa, 2010
35	pool	2	-	2***	An in house method based on Zarlenga et al., 1999
40	single larvae	12	-	-	De Brunye et al., 2005

*the lab was unable to combine the PCR pattern with the *Trichinella* species.

**no PCR amplicons were obtained.

***the used method was not suitable to identify two of the species present in the panel.