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**AFNOR validation of PremiTest, a microbiological-based screening tube test for the detection of antimicrobial residues, in muscles from different animal origins**

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3 1 **AFNOR validation of Premi®Test, a microbiological-based screening tube test**  
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6 2 **for the detection of antimicrobial residues, in muscles from different animal**  
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8 3 **origins**  
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1  
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3 18 **Abstract**  
4

5 19 Premi®Test contains viable spores of a strain of *Bacillus stearothermophilus* which is  
6  
7 20 sensitive to antimicrobial residues such as beta-lactams, tetracyclines, macrolides, and  
8  
9 21 sulphonamides etc. The growth of the strain is inhibited by the presence of antimicrobial  
10  
11 22 residues in muscle samples. Premi®Test was validated according to AFNOR rules (French  
12  
13 Association for Normalisation). The AFNOR validation was based on the comparison of a  
14  
15 23 reference method [French Official method (Four Plate Test) and the STAR protocol (5 plate  
16  
17 24 test)] with the alternative method (Premi®Test). A preliminary study was conducted in an  
18  
19 25 expert laboratory (Community Reference Laboratory CRL) on both spiked and incurred  
20  
21 26 samples (field samples). Several method performance criteria (sensitivity, specificity, relative  
22  
23 27 accuracy) were estimated and are discussed, in addition to detection capabilities. Adequate  
24  
25 28 agreement was found between alternative methods and the reference method. However,  
26  
27 29 Premi®Test was more sensitive to beta-lactams and sulphonamides than the FPT.  
28  
29 30 Subsequently, a collaborative study with 11 laboratories was organised by the CRL. Blank  
30  
31 31 and spiked meat juice samples were sent to participants. The expert laboratory (CRL)  
32  
33 32 statistically analysed the results. It was concluded that Premi® Test could be used for the  
34  
35 33 routine determination of antimicrobial residues in muscle of different animal origin with  
36  
37 34 acceptable analytical performance. The detection capabilities of Premi®Test for beta-lactams  
38  
39 35 (amoxicillin, ceftiofur), one macrolide (tylosin) and tetracycline were at the level of the  
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41 36 respective Maximum Residue Limits (MRL) in muscle samples or even lower.  
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39 **Keywords:** *Validation, Premi®test, screening, antimicrobial residues, muscle, collaborative*  
40 *study, routine analysis*

## 41 Introduction

42 Animal treatment can lead to the presence of residues in food of animal origin. The presence  
43 of antimicrobial residues could lead to human safety problems such as allergies or toxicity  
44 when foodstuffs containing residues enter the food chain. Different families of antimicrobial  
45 residues are concerned: beta-lactams, sulfonamides, macrolides, tetracyclines, quinolones,  
46 etc. For these reasons, Maximum Residue Limits (MRL) were set for many antimicrobial  
47 residues to protect consumer safety (EEC Directive 2377/90 and amendments). Different  
48 methods were developed for the detection of antimicrobial residues in food of animal origin.  
49 These screening methods are usually microbiological methods which are based on the  
50 inhibition of bacterial growth by antimicrobial residues. Microbiological plate tests generally  
51 give results in 18 to 24 hours. However, these home-made methods require skilled  
52 technicians, with a specific training. Microbiological based tests are interesting at the  
53 screening step because they are able to detect a wide range of antimicrobial residues  
54 (several families of antimicrobials are detectable with one single method). The levels of  
55 detection of these methods are generally satisfactory with respect to European regulations  
56 (EEC, 1990) for beta-lactams, tetracyclines and macrolides (detection capabilities lower or  
57 equal to the MRL). The best detected class of compound is generally the beta-lactam family.  
58 The detection capabilities of sulphonamides are very variable depending on the  
59 sulphonamide structure. The least detected antibiotics are generally some sulphonamides,  
60 aminoglycosides and quinolones by microbiological based tests. None of the existing  
61 microbiological screening methods are able to detect all the MRL substances in animal  
62 matrices. Therefore, some countries implemented other kind of methods in parallel to look for  
63 these antimicrobials specifically. In France, the official method for the control of muscle  
64 samples is the Four Plate Test (FPT) (Bogaerts and Wolf 1980). In the past, some  
65 commercial tube tests already exist but only for the analysis of milk, like Delvotest® and  
66 since a few years COPAN® test. These tests are based on the detection of growth by  
67 production of acid, visible by a colour change of the test medium (pH indicator). More  
68 recently, the Premi®Test, a commercial growth inhibitor test, was developed for the detection

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3 69 of antimicrobial residues in muscle by DSM (DSM Food Specialities R&D, Delft, the  
4  
5 70 Netherlands). This test is based on the inhibition of the strain *Bacillus stearothermophilus*.  
6  
7 71 Premi®Test allows to detect antimicrobial residues in muscle in less than 4 hours (Fabre  
8  
9 72 2003; Stead. 2004; Fabre *et al.* 2004)).  
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12 73

13  
14 74 Over the last few years, several evaluations of the Premi®Test have already been published,  
15  
16 75 based on spiked meat juice samples (Reybroeck 2000a) or incurred poultry muscles  
17  
18 76 (Reybroeck 2000b). Premi®Test has generally been compared to a reference method. Some  
19  
20 77 studies were focused on a family of antimicrobials like beta-lactams, using spiked meat juice  
21  
22 78 samples and incurred poultry muscle samples (Popelka *et al.* 2005) or tetracyclines  
23  
24 79 (Okerman *et al.* 2004). The detection limits of different antimicrobials, calculated in spiked  
25  
26 80 juice samples, were very near in the different studies. The global conclusions of the different  
27  
28 81 studies were that Premi®Test is not suited for the detection of tetracyclines at MRL level in  
29  
30 82 spiked and incurred samples. Moreover, Premi®Test was very sensitive for the screening of  
31  
32 83 beta-lactams, more than usual microbiological plate tests. Premi®Test is suited for the  
33  
34 84 detection of beta-lactams and sulphonamides, at MRL level or even below MRL.  
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40 86 The AFNOR validation is based on the comparison between a reference method and an  
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42 87 alternative method. In the present study, the Premi®Test, the alternative method, was  
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44 88 compared with 2 other microbiological methods: the French Official method (Four Plate Test)  
45  
46 89 which was the reference method and the STAR protocol (the CRL protocol) (Gaudin *et al.*  
47  
48 90 2004). Afterwards, an interlaboratory study was organised, where only the Premi®Test has  
49  
50 91 been used to analyse spiked meat juice samples. This paper presents the results of  
51  
52 92 preliminary and collaborative studies.  
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## 57 94 **Materials and methods**

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3 95 *Presentation of the methods*

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5 96 *Principle of the kit.* The Premi®Test allows to detect antimicrobial residues in fresh meat,  
6  
7 97 kidneys, fish and eggs (Arts et al. 2000). The Premi®Test is based on the growth inhibition of  
8  
9  
10 98 *Bacillus stearothermophilus*. Standardized spores are included in a medium, with selected  
11  
12 99 nutrients. The meat juice was put on ready-to-use tubes. After 20 min of pre-diffusion at room  
13  
14 100 temperature, the meat juice was removed by three washing steps. Finally, the ampoule was  
15  
16 101 incubated during approximately 3 hours at 64°C. The reading of the "yes/no" result was  
17  
18 102 based on a colour comparison. Without antimicrobials, the spores germinated and  
19  
20 103 developed, involving the acidification of the medium and a change of colour (yellow).  
21  
22 104 Conversely, in the presence of antimicrobials, the bacterial growth was inhibited. A purple  
23  
24 105 colour indicated the presence of antibiotics, at or above the detection limit of the test.  
25  
26 106 Doubtful and positive samples were confirmed by the multi-residue LC/MS-MS method  
27  
28 107 described below. This broad spectrum test makes it possible to detect a great number of  
29  
30 108 antimicrobials usually used, in less than 4 hours, on muscle juice samples (extracted by  
31  
32 109 pressing a piece of meat).  
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37  
38 111 *Principle of the Four Plate Test (FPT).* The Four Plate Test is the French Official method for  
39  
40 112 the control of muscle samples (Bogaerts and Wolf 1980). A microorganism sensitive to  
41  
42 113 antibacterial substances is inoculated into an agar medium in a Petri dish. The following test  
43  
44 114 organisms are used: *Bacillus subtilis* BGA (reference 10649, Merck) (in test agar pH 6,  
45  
46 115 Merck), *Bacillus subtilis* BGA (reference 10649, Merck) (in test agar pH 8, Merck), *Bacillus*  
47  
48 116 *subtilis* BGA (reference 10649, Merck) (in agar ASS pH 7.4, Merck) and *Kocuria varians*  
49  
50 117 (ATCC 9341, Pasteur Institute) (in test agar pH 8, Merck). Slices of frozen muscle were  
51  
52 118 placed on the surface of the inoculated medium, and then incubated at the optimal  
53  
54 119 temperature for growth of the test organism. After diffusion, the presence of antibacterial  
55  
56 120 substance should produce an inhibition zone around the sample by inhibiting the growth of  
57  
58 121 the test organism.  
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3 123 *Principle of the STAR protocol and Bacillus cereus.* The STAR protocol is the CRL method  
4  
5 124 (Gaudin et al. 2004) which was developed to improve the performance of the Four Plate  
6  
7 125 Test. The detection principle is the same as Four Plate Test. The following test organisms  
8  
9  
10 126 are used: *Bacillus subtilis* BGA (*reference 10649, Merck*) (Antibiotic medium II at pH 8.0,  
11  
12 127 Difco), *Kocuria varians* ex. *Micrococcus luteus* (ATCC 9341, *Pasteur Institute*) (in test agar at  
13  
14 128 pH 8, Merck), *Bacillus cereus* Bc6 (ATCC 11778, *Pasteur Institute*) (in test agar at pH 6,  
15  
16 129 Merck), *Escherichia coli* (ATCC 11303, *Pasteur Institute*) (in test agar at pH 8, Merck),  
17  
18 130 *Bacillus stearothermophilus* (ATCC 10149, *Merck*) (in DST (Diagnostic Sensitive Test)  
19  
20 131 medium, Oxoid). The plate *Bacillus cereus* which was used by the field laboratories at the 3<sup>rd</sup>  
21  
22 132 step of the preliminary study is the plate Bc6 of the STAR protocol, which is selective for the  
23  
24 133 detection of the tetracyclines' family. Slices of muscle samples of 2 mm in thickness are cut  
25  
26 134 in frozen muscle and put on the plates.

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29 135  
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31 136 *LC/MS-MS method.* A multi-residue LC/MS-MS method was developed for the screening of  
32  
33 137 antimicrobials in meat. The principle is based on two different extractions : one with  
34  
35 138 trichloroacetic acid (TCA) allowing the detection of tetracyclines, aminoglycosides and  
36  
37 139 quinolones, the second with acetonitrile (ACN) allowing the detection of penicillins,  
38  
39 140 cephalosporins, macrolides and sulfonamides. TCA extracts are directly injected after  
41  
42 141 ultracentrifugation and filtration. ACN extracts are evaporated and the residue is then  
43  
44 142 dissolved in 0.6 ml of ammonium acetate before filtration and injection. Two different  
45  
46 143 gradients with pentafluoropropionic acid and ACN are used for the LC analyses. LC/MS-MS  
47  
48 144 is used with a Multi Reaction Monitoring Mode (MRM) and two MRM transitions are  
49  
50 145 monitored for each compound. The identification of the detected compounds is based on the  
51  
52 146 retention time and the presence of the two specific transitions. The quantitative determination  
53  
54 147 was carried out by using calibration curves obtained with spiked samples at 0.5 MRL, 1 MRL  
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56 148 and 1.5 MRL levels. Internal standard were used for the quantification. All the 50 monitored  
57  
58 149 compounds except some aminoglycosides were detected at a level below the MRL.

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3 151 *Principle of AFNOR (French Association for Normalisation) validation*

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5 152 An alternative method should be compared with a reference method. A reference method  
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7 153 could be a standardized method when it exists, an official method, or a widely known and  
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9  
10 154 used method, taken in reference. An alternative method is a commercial test allowing to  
11  
12 155 analyze, for a category of products given, the same analytes as that measured by the  
13  
14 156 reference method, but which presents moreover, one or more criteria following: speed of  
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16 157 analysis, easiness of execution and/or automation, analytical performances (limit of  
17  
18 158 detection, specificity, etc). A specific guide for validation of alternative methods in the field of  
19  
20 159 detection of antimicrobial residues in foodstuffs of animal origin was edited (Anon. 2005a). It  
21  
22 160 defined the requirements relating to the organisation of preliminary and collaborative studies,  
23  
24 161 carried out by one expert laboratory. This document established the general principle as well  
25  
26 162 as the technical protocol for the validation of alternative methods in the field of detection of  
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28 163 antimicrobial residues in foodstuffs of animal origin.  
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33 165 *Preliminary study*

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35 166 The preliminary step has been divided in 3 parts.

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37 167 *Step 1. Analysis of blank and spiked meat juice samples.* Porcine muscles came out of  
38  
39 168 practice . The absence of antimicrobial substances in these pig samples, assumed as blank  
40  
41 169 materials, was checked with the FPT and the STAR protocol. The negative results confirmed  
42  
43 170 that these pig samples could be used to prepare blank meat juices.

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45 171 Muscle samples were pressed with garlic press. Afterwards, meat juice samples were spiked  
46  
47 172 with known concentrations of 6 different antimicrobials: sulfamethazine (sulphonamides),  
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49 173 oxytetracycline (tetracyclines), tylosin (macrolides), amoxicillin (penicillins), ceftiofur  
50  
51 174 (cephalosporins) and gentamycine (aminoglycosides). Each antimicrobial was spiked at 3  
52  
53 175 concentrations: sulfamethazine (50, 100, 200  $\mu\text{g kg}^{-1}$ ), oxytetracycline (50, 100, 200  $\mu\text{g kg}^{-1}$ ),  
54  
55 176 tylosin (50, 100, 200  $\mu\text{g kg}^{-1}$ ), amoxicillin (25, 50, 100  $\mu\text{g kg}^{-1}$ ), ceftiofur (100, 200, 400  $\mu\text{g kg}^{-1}$ )  
56  
57 177 and gentamycine (50, 100, 200  $\mu\text{g kg}^{-1}$ ).

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3 178 5 aliquots (5 combinations sample/antimicrobial/concentration) were prepared “blind to the  
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5 179 technician” and analysed in duplicate only with the Premi®Test. The aliquots were stored in  
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7 180 freezer (approximately -20°C) before analyses. Blank meat juice and spiked meat juice  
8  
9 181 samples were analysed in duplicate only with the Premi®Test.  
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14 183 *Step 2. Treatment of animals and analysis of incurred porcine muscle samples.* Two  
15  
16 184 untreated pigs were slaughtered before the treatments of 3 other pigs. These pigs were  
17  
18 185 supplied by a farm guaranteeing that the pigs did not receive antimicrobial treatments before.  
19  
20 186 Moreover, the absence of antimicrobials in muscle samples was checked with the FPT and  
21  
22 187 the STAR protocol.  
23  
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25 188 Three pigs were treated: one with tylosin, one with amoxicillin and one with a mXture of  
26  
27 189 oxytetracycline and sulfadimethoxine and slaughtered. Each material was analysed by a  
28  
29 190 multi-residue LC/MS-MS method to quantify antimicrobials. The following concentrations  
30  
31 191 were obtained: tylosin  $750.9 \pm 76.1 \mu\text{gkg}^{-1}$  (material 1), amoxicillin  $269.5 \pm 18.0 \mu\text{gkg}^{-1}$   
32  
33 192 (material 2), oxytetracycline  $764.9 \pm 44.8 \mu\text{gkg}^{-1}$  and sulfadimethoxine  $151.1 \pm 13.2 \mu\text{gkg}^{-1}$   
34  
35 193 (material 3).  
36  
37

38 194 Afterwards, 5 pieces of each material (blank and treated animals) of approximately 20g were  
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40 195 (5 combinations sample/antimicrobial/concentration) prepared “blind to the technician” and  
41  
42 196 analysed in duplicate with 3 methods: Premi®Test, STAR protocol and FPT. These samples  
43  
44 197 were stored in freezer at approximately -20°C before analyses.  
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47 198

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49 199 *Step 3. Analysis of field samples.* This step allowed to compare Premi®Test, STAR protocol  
50  
51 200 and FPT, on “naturally” incurred samples, for a wide number of antimicrobials and matrices  
52  
53 201 of different origins (e.g. different species). The samples came from a pilot study concerning  
54  
55 202 the implementation of a new screening method of the antimicrobial residues in meat (Anon.  
56  
57 203 2005b). The samples were sent regularly to laboratories from the French veterinary services  
58  
59 204 (official control). SX field veterinary laboratories (LVD or “Laboratoire Vétérinaire  
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205 Départemental” in French) were trained to the implementation of the Premi®Test and one

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3 206 plate *Bacillus cereus* (detection of tetracyclines) (STAR protocol) and analyzed 1427 field  
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5 207 samples, over a period of 4 months. All their positive samples (with Premi®Test and/or  
6  
7 208 *Bacillus cereus*) were sent frozen to the CRL, every 15 days. All the samples characteristics  
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9  
10 209 (species, reception date, analysis date, sending date, ...etc...), as well as the results  
11  
12 210 obtained with Premi®Test and *Bacillus cereus*, were also sent to the CRL. All these samples  
13  
14 211 coming from the field laboratories where stored in the freezer at their arrival in our laboratory.  
15  
16 212 Then, blind analyses were carried out at the CRL with the Premi®Test, the FPT, and the  
17  
18 213 STAR protocol including the *Bacillus cereus* plate. Finally, the positive samples with at least  
19  
20 214 one of these methods were analyzed by the multi-residue LC/MS-MS method (systematic  
21  
22 215 screening, then identification and quantification). The doubtful samples with Premi®Test  
23  
24 216 were also tested by LC/MS-MS.  
25  
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27 217  
28

#### 29 218 *Interlaboratory study*

30  
31 219 11 laboratories have been contacted to participate to this inter-laboratory study, including the  
32  
33 220 6 field laboratories which had already participated to the step 3 of the preliminary study.  
34  
35 221 When an AFNOR validation is implemented, an expert laboratory is designated by the  
36  
37 222 AFNOR Technical Office. The expert laboratory performed the intra-laboratory validation and  
38  
39 223 organised the inter-laboratory study. Moreover, during the inter-laboratory study, the results  
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41 224 of the expert laboratory are not included with all the participants' results, but these results are  
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43 225 considered as the reference results (to be obtained by the participants). In this case, the  
44  
45 226 Community Reference Laboratory was considered as the expert laboratory.  
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47  
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49 227

50  
51 228 Spiked muscle juice samples were used for the interlaboratory study, instead of incurred  
52  
53 229 samples, for several reasons: Firstly, it is very difficult to prepare raw muscle with a target  
54  
55 230 concentration of each antimicrobial because of the individual variability of animals'  
56  
57 231 pharmacokinetic (depletion of the antibiotic in tissues). Moreover, the use of raw pieces of  
58  
59 232 meat did not allow to mince meat to adjust the concentration. Secondly, the study was based  
60  
233 on several animal species. Therefore, the treatment should have been performed in each

1  
2  
3 234 specie. This would have been very expensive, time consuming and impossible to be  
4  
5 235 implemented in our laboratory. Thirdly, when producing incurred muscles and sending raw  
6  
7 236 muscle samples (pieces of meat), it is very difficult to ensure the homogeneity of the  
8  
9  
10 237 samples, which is a basic condition when organising a collaborative study.  
11

12 238  
13  
14 239 *Preparation of the materials.* Porcine, bovine and chicken muscles were pressed. Blank meat  
15  
16 240 juice samples (negative controls and blank unknown samples) were prepared. Furthermore,  
17  
18 241 spiked samples were prepared with 4 different antibiotics, at 3 concentrations  
19  
20 242 Oxytetracycline and ceftiofur were added to porcine meat juice samples, sulfamethazine to  
21  
22 243 bovine meat juice samples and finally tylosin to chicken meat juice samples Table I presents  
23  
24 244 the content of the 16 combinations.  
25

26  
27 245 “[Insert Table I about here]”  
28

29 246 A stability study was carried out on materials before sending of the samples and over the  
30  
31 247 period of analyses by the participants. The samples were analysed with the Premi®Test at  
32  
33 248 the CRL. Stability of materials was proved over the period of analyses. A random codification  
34  
35 249 of the materials was performed. Each laboratory was identified by a code (from A to P).  
36

37 250  
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40 251 *Sending of the materials.* One negative control from each species was sent and one positive  
41  
42 252 control containing penicillin G at  $10 \mu\text{g kg}^{-1}$ . 32 different frozen meat juice samples (16  
43  
44 253 materials in blind duplicate: spiked and blank samples) were sent under frozen conditions  
45  
46 254 (dried ice) to the participants, in order to ensure stability of matrix and analyte. The samples  
47  
48 255 were stored in freezer at their arrival. Eleven laboratories received with their parcel  
49  
50 256 instructions and a results form. The participants had no information about the antibiotics  
51  
52 257 contained in the materials.  
53

54 258  
55  
56  
57 259 Because it is recommended to test one negative control from each analysed species, each  
58  
59 260 laboratory received a table indicating the species of origin of each sample, according to  
60  
261 sample code (1 to 32), in order to compare the results of each sample with the negative

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2  
3 262 control of the corresponding species. All analyses were carried out in blind duplicate (2  
4  
5 263 different series of analyses), with Premi®Test only. The analyses were performed within one  
6  
7 264 week maximum after receiving the samples. A negative control for each species, a positive  
8  
9 265 sample (spiked with penicillin G at 10 µg kg<sup>-1</sup>) and coded materials were analysed. All the  
10  
11 266 results were returned rapidly and compiled at the CRL.  
12  
13  
14 267

## 16 268 **Results and discussion**

### 18 269 *Preliminary study*

20 270 *Step 1. Analysis of blank and spiked meat juice samples.* Among the 20 blank meat juice  
21  
22 271 samples analysed in duplicate, only one meat juice was positive twice and 4 other meat  
23  
24 272 juices were "doubtful" on one of the two repetitions and negative on the other. 15 meat juices  
25  
26 273 were negative twice. The false positive rate is the number of positive results for blank  
27  
28 274 samples (free of antibiotic substances) divided by the total number of positive samples (the  
29  
30 275 same number plus the number of contaminated samples showing positive results) and  
31  
32 276 multiplied by 100. 18% ( $= 6/(6+27)*100$ ) of false positive results was observed. The result  
33  
34 277 was satisfactory because the false positive rate should be minimal for a screening method,  
35  
36 278 since the samples declared positive should be confirmed by a physicochemical method for  
37  
38 279 identification and quantification.  
39  
40  
41  
42 280

44 281 The detection limit corresponded to the lowest concentration which gave a positive or a  
45  
46 282 doubtful result for each of the 5 replicates. Table II presents the determination of limits of  
47  
48 283 detection of Premi®Test for the 6 antimicrobials tested.  
49

51 284 "[Insert Table II about here]"  
52  
53 285

55 286 The study of spiked pig meat juice samples showed that, for 5 molecules belonging to 4  
56  
57 287 different classes of antibiotics, the detection limit of Premi®Test was at the level of one or  
58  
59 288 two MRL maximum. For only one antibiotic (gentamicin), the detection limit was higher than 2  
60  
289 times the MRL (40% of positive results at 2xMRL). The false negative rate corresponds to

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3 290 the number of negative results obtained for contaminated samples (spiked samples) divided  
4  
5 291 by the total number of negative samples (the same number plus the number of blank  
6  
7 292 samples giving a negative result) and multiplied by 100. The false negative rate calculated at  
8  
9 293 1\*MRL was equal to 22 % ( = 10/(10+34)). However, the false negative rate calculated at  
10  
11 294 twice the MRL was equal to 8 % ( = 3/(3+34)).  
12  
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14 295

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16 296 *Step 2. Treatment of animals and analysis of incurred porcine muscle samples.* The  
17  
18 297 concentrations of naturally incurred samples were much higher than the respective MRL of  
19  
20 298 the 4 antimicrobials (from 1.5 to 7.6 times the MRL). However, the analyses of incurred  
21  
22 299 materials with known antimicrobial concentrations was of great interest. Table III summarises  
23  
24  
25 300 the results of step 2.

26  
27 301 “[Insert Table III about here]”  
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29 302

30  
31 303 The data exploitation was carried out according to the reference document of AFNOR  
32  
33 304 validation (Anon. 2005a). The exploitation is based on the comparison of 2 methods: the  
34  
35 305 reference method and the alternative method. The FPT was set as the reference method and  
36  
37 306 the Premi®Test is the alternative method.  
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42 308 Three different parameters were calculated which allowed to compare reference and  
43  
44 309 alternative method: relative accuracy ( $AC = (PA + NA)/N \cdot 100\%$ ), relative specificity ( $SP =$   
45  
46 310  $(NA/N_-) \cdot 100\%$ ), relative sensitivity ( $SE = (PA/N_+) \cdot 100\%$ ), where: NA is the negative  
47  
48 311 agreement (negative result obtained with both methods), PA: the positive agreement  
49  
50 312 (positive result obtained with both methods), ND: the negative discrepancy (positive result  
51  
52 313 obtained with reference method and negative result with alternative test), PD: the positive  
53  
54 314 discrepancy (negative result obtained with reference method and positive result with  
55  
56 315 alternative test); N = NA + PA + PD + ND: total number of samples; N- is the total number of  
57  
58 316 negative samples obtained with the reference method (NA + PD); N+ is the total number of  
59  
60 317 positive samples obtained with the reference method (PA + ND).

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3 318 The relative accuracy, the relative specificity and the relative sensitivity were calculated as  
4  
5 319 70%, 42.9% and 84.6% respectively. Therefore, the relative accuracy and specificity were  
6  
7 320 quite satisfactory. The low value of relative specificity could be explained because the  
8  
9 321 sensitivity of the FPT was sometimes insufficient for certain antibiotics.  
10  
11  
12 322

13  
14 323 In conclusion, FPT and Premi®Test gave concordant results when looking at the  
15  
16 324 performance characteristics (relative accuracy, the relative specificity and the relative  
17  
18 325 sensitivity). However, at this step, the false negative and the false positive rates were lower  
19  
20 326 for the Premi®Test (both 0 %) than for the FPT (33 and 40 % respectively).  
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23 327

24  
25 328 *Step 3. Analysis of field samples.* 1427 field incurred samples, originated from 6 French field  
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27 329 veterinary laboratories, were analyzed implementing Premi®Test and *Bacillus cereus* plate.  
28  
29 330 1325 samples were detected negative and 102 samples doubtful (36) and positive (66) with  
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31 331 the Premi®Test.  
32  
33 332

34  
35 333 Furthermore, among the 1325 negative results, 10 samples were detected positive with the  
36  
37 334 *Bacillus cereus* plate. Therefore, 112 muscle samples were sent to the CRL to be confirmed.  
38  
39 335 The correlation between the Premi®Test results of LVD and AFSSA was studied (Table IV).  
40  
41 336 76% of the samples (78/102) found positive or doubtful with Premi®Test in the LVD, were  
42  
43 337 found positive or doubtful with Premi®Test at the CRL. 24 samples were positive or doubtful  
44  
45 338 at the LVD and negative at AFSSA. It could be due either to false positive results of the field  
46  
47 339 laboratories, or to antibiotic instability between the 2 analyses, although the best storage and  
48  
49 340 transport conditions have been strictly respected. Among the 10 samples negative with  
50  
51 341 Premi®Test and positive with *Bacillus cereus* in the field laboratories, 6 were found positive  
52  
53 342 or doubtful with Premi®Test at the CRL and confirmed positive with *Bacillus cereus*. The 4  
54  
55 343 other samples were found positive only on plate Bc6. Finally, the presence of a tetracycline  
56  
57 344 was confirmed by LC/MS-MS in 9 of these samples. Therefore, *Bacillus cereus* plate (one  
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3 345 plate of the STAR protocol, which is selective for the detection of tetracyclines) is more  
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5 346 sensitive for tetracyclines than Premi®Test.

7 347 “[Insert Table IV about here]”  
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10 348  
11 349 The false positive rate in field laboratories was sometimes high, mainly at the beginning of  
12  
13 350 the study. Afterwards, with the reading experience, most of the positive results with  
14  
15 351 Premi®Test at the field laboratories were confirmed positive with Premi®Test at the CRL.  
16  
17 352 Many animal species were studied in step 3 because samples were from field laboratories  
18  
19 353 origin, while the 2 first steps were based only on pig muscle. The distribution of the samples  
20  
21 354 in the different species is presented (Table IV).  
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25 355  
26  
27 356 Confirmatory rate of LC/MS-MS method (number of samples really containing antimicrobial  
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29 357 residues divided by the number of tested samples (positive screening) multiplied by 100) was  
30  
31 358 equal to 41% (25/61). The confirmatory rate varied between species. There were more false  
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33 359 positive results in some species (37 % for bovine samples instead of 75 % for poultry  
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35 360 samples).  
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38 361  
39  
40 362 The results of the comparative analysis between the alternative method Premi®Test and the  
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42 363 reference method FPT are presented in Table V.  
43

44 364 “[Insert Table V about here]”  
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49 366 When the data analysis was based only on the 112 samples reanalysed in the expert  
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51 367 laboratory, relative accuracy (33.9%) and relative specificity (33.4%) were very low. Only  
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53 368 relative sensitivity (70 %) was satisfactory, as in the first step of the validation study.  
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55 369 However, in this case, Premi®Test negative samples (1315 negative samples / 1427  
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57 370 analyzed samples) obtained in the field laboratories were not taken into account. Only the  
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59 371 positive samples in the field laboratories were reanalysed at the CRL with Premi®Test and  
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372 FPT. None of the 1315 negative samples was analysed with the FPT. By experience (a



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3 373 previous study in 2005) (data not shown), all the samples which were declared negative with  
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5 374 Premi®Test and negative with *Bacillus cereus* were also all negative with the FPT (28  
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7 375 samples). Therefore it was assumed that the 1315 samples, found negative with both  
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9  
10 376 Premi®Test and *Bacillus cereus*, would have also been negative with the FPT. In this case,  
11  
12 377 the agreement (relative accuracy 94.8%, relative specificity 95%) between the results of  
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14 378 Premi®Test and FPT was very satisfactory. The 2 methods were declared different by a  
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16 379 statistical test, because of a high rate of positive deviation (Premi®Test + / FPT -) which may  
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18 380 be caused partially by false positive results of the Premi®Test and/or by the lack of sensitivity  
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20 381 of the reference method towards some antimicrobial residues.  
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23 382

24  
25 383 The agreement (relative accuracy) between the results of Premi®Test and STAR was equal  
26  
27 384 to 42.99%, the relative specificity 31.3% and the relative sensitivity 71.9%. Therefore, it is  
28  
29 385 quite similar to the comparison of FPT with Premi®Test. The agreement between the FPT  
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31 386 and the STAR protocol was higher (70 %) (data not shown). Only one sample was negative  
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33 387 with the STAR protocol and positive with the FPT. The absence of antimicrobial residues in  
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35 388 this sample was confirmed by LC/MS-MS.  
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40 390 The results obtained in the field laboratories, and then at the CRL, with the Premi®Test, the  
41  
42 391 FPT, the STAR protocol and the confirmatory method by LC/MS-MS are compared (Table  
43  
44 392 VI). The number of positive results with the Premi®Test (78) was much higher than the  
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46 393 number of positive with the multi-plate tests (FPT 10; STAR 31 samples). Among the 112  
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48 394 samples analysed at the CRL, only the positive or doubtful samples with one of the screening  
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50 395 methods (Premi®Test, FPT, STAR) were confirmed by LC/MS-MS (88 samples).  
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52

53 396 “[Insert Table VI about here]”  
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56  
57 398 After the confirmatory analyses, 31 samples really contained antimicrobial residues (35% of  
58  
59 399 the confirmed samples) at different levels (26 samples at concentrations lower than the  
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400 respective MRLs and 5 only at concentrations higher than the respective MRLs).

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3 401 The false positive rates of the FPT, the Premi®Test and the STAR protocol were 3%  
4  
5 402  $((1/(1+31))*100)$ , 62%  $((52/(52+31))*100)$  and 31 %  $((14/(14+31))*100)$  respectively. The false  
6  
7 403 positive (rate of each method was calculated as the number of positive results for blank  
8  
9 404 samples (free of antibiotic substances) divided by the total number of positive samples (the  
10  
11 405 same number plus the number of contaminated samples showing positive results) and  
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13 406 multiplied by 100.

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16 407  
17  
18 408 52 samples were Premi®Test positive or doubtful, but no antibiotic residue could be  
19  
20 409 identified when analyzed by LC/MS-MS (). However, a false-positive result could be due to  
21  
22 410 the degradation of the antimicrobials initially contained in the sample (too long delay before  
23  
24 411 the analysis by LC/MS-MS, very unstable molecules...). Furthermore, it could be false  
25  
26 412 compliant results of the confirmatory method. The antimicrobial present in the sample could  
27  
28 413 not be detected and identified by the multi-residue LC/MS-MS method if the molecule is not  
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30 414 present in the spectrum of detection of the method (metabolites, ...). or the sensitivity could  
31  
32 415 be insufficient.

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35 416  
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38 417 After a positive result with Premi®Test, the presence of antimicrobial residues was  
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40 418 confirmed in 26 samples (33 %), from which 4 samples contained antimicrobials at  
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42 419 concentrations upper than the MRL. Three of the 10 positive FPT samples and 4 of the  
43  
44 420 STAR positive samples were at concentrations upper than the MRL. Premi®Test and STAR  
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46 421 screened positive 22 and 13 samples respectively which really contained antimicrobial  
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48 422 compounds at concentrations lower than MRL. The FPT detected only 6 of these samples.  
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50  
51 423 Therefore, the sensitivity of Premi®Test and STAR protocol was lower than those of the FPT.  
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53 424 The highest confirmatory rate was obtained for the FPT (90 %). The false negative rates of  
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55 425 the FPT, the Premi®Test and the STAR protocol were equal to 8 %, 28 % and 20 %  
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57 426 respectively. The false negative rate was calculated as the number of false negative results  
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59 427 divided by the total of the true negative samples plus the number of false negative results for  
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428 each method.

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3 429 The most commonly detected antimicrobial classes detected during this pilot study were  
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5 430 beta-lactams (penicillin and amoxicillin) (15 samples) and tetracyclines (doxycycline,  
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7 431 tetracycline, oxytetracycline and chlortetracycline) (15 samples). Premi®Test (12 positive  
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9 432 samples) was much more sensitive for the detection of beta-lactams than the FPT (1 positive  
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11 433 sample) and the STAR protocol (5 positive samples), even too sensitive in some cases (10  
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13 434 positive samples confirmed at concentrations lower than MRL). Premi®Test was also more  
14  
15 435 sensitive for the screening of sulphonamides than the 2 other methods (FPT and STAR failed  
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17 436 to detect 1 sample at a concentration higher than the MRL). Finally, Premi®Test was less  
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19 437 sensitive than the FPT and especially than the STAR protocol for the detection of  
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21 438 tetracyclines. Therefore, this study demonstrated the ability of Premi®Test to detect samples  
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23 439 at the MRL level for sulphonamides and beta-lactams, but not for tetracyclines. These  
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25 440 conclusions confirmed the results obtained by different teams during the past years  
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27 441 ((Reybroeck 2000a, (Reybroeck 2000b, Okerman *et al.* 2004, Popelka *et al.* 2005).  
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34 443 Table VII presented the combined results of a previous study realized in 2003 and 2004 (64  
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36 444 samples really containing antimicrobial residues) with the pilot study.  
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38 445 “[Insert Table VII about here]”  
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43 447 After the confirmatory analyses of positive samples, 31 samples contained antimicrobial  
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45 448 residues at concentrations lower than the respective MRLs and 33 samples at concentrations  
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47 449 higher than the respective MRLs. The total number of positive samples reported with  
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49 450 Premi®Test (52) was higher than with FPT (38). However, the number of samples detected  
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51 451 positive with Premi®Test (26) or FPT (25) which really contained antimicrobial compounds at  
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53 452 concentrations higher than MRL was identical. Moreover, 6 false negative results were  
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55 453 obtained with Premi®Test and 7 with FPT.  
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3 455 The presence of beta-lactams was confirmed in 17 samples by the LC/MS-MS method. All of  
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5 456 these 17 samples were detected positive or doubtful samples with Premi®Test, while only 7  
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7 457 samples were detected positive with the FPT (1<MRL and 6> MRL). Therefore, the detection  
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9 458 capability of Premi®Test for beta-lactams was better than FPT (). The sensitivity was also  
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11 459 lower for sulphonamides (4 positive samples Premi®Test, 4 negative samples FPT). Two of  
12  
13 460 these samples really contained sulphonamides at concentrations higher than the MRL and  
14  
15 461 the FPT failed to detect them while the Premi®Test succeeded to detect. However, the  
16  
17 462 sensitivity of FPT for tetracyclines was better (15 doubtful or positive samples with  
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19 463 Premi®Test instead of 19 positive samples with FPT).  
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25 465 The most detected antimicrobials were tetracyclines (25 samples) and beta-lactams (17  
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27 466 samples). Then, macrolides (6), sulphonamides (4), quinolones (3) or mix of antimicrobials  
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29 467 (8) were found.  
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#### 33 469 *Collaborative study*

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35 470 *Results of the expert laboratory.* The expert laboratory obtained only one positive false result  
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37 471 for the 2 series of analyses for the white chicken sample (code 10), whereas the same  
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39 472 sample (code 27) was found negative for the 2 series. The L1 concentration was selected to  
40  
41 473 give negative results (Table I). The expert laboratory did not obtain any false negative result  
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43 474 because the L3 concentration was that designed to give positive results, whereas the L2  
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45 475 concentration was to be at the limit of sensitivity of the test.  
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51 477 It should be noted that the later analysis of the results was carried out while taking as  
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53 478 principle that the doubtful results are positive results, this which is applied at the time for  
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55 479 routine analyses.  
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57 480 “[Insert Table VIII about here]”  
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3 482 The results of the expert laboratory were very satisfactory (Table VIII): 14 blank samples out  
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5 483 of 16 were negative (only 2 positive results for chicken samples) (L0). Below the assumed  
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7 484 limit of detection, 14 samples out of 16 were negative (L1). At the assumed limit of detection,  
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10 485 all the samples were positive (L2). Above the assumed limit of detection, all the samples  
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12 486 were positive (L3).

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16 488 *Results of the participants.* Before the study, the expert laboratory stated that the results of  
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18 489 one laboratory would be removed when: the negative control was detected positive, the  
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20 490 positive control was detected negative, if samples were in bad condition at the reception, if a  
21  
22 491 problem of storage was established. Two laboratories were finally eliminated from the  
23  
24 492 analysis of the results: Laboratory L for several reasons: the parcel delivery was delayed,  
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26 493 samples were defrosted and chicken negative controls were detected positive for the 2 series  
27  
28 494 of analyses; Laboratory H because chicken negative controls were detected positive for 1st  
29  
30 495 series of analyses. The raw data of the participants for the 16 combinations are presented in  
31  
32 496 Table IX. Each participant received 32 samples (16 materials in blind duplicate) that they  
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34 497 have analysed in 2 different series.

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38 498 “[Insert Table IX about here]”  
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42 500 The results were homogeneous between laboratories. The results were analysed and  
43  
44 501 summarised in Table X. Concerning negative samples (L0) (16 by laboratories corresponding  
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46 502 to 3 different species), 4 laboratories found 100% of negative results, 5 laboratories found 1  
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48 503 or 2 samples doubtful or positive, sometimes in only one series of analyses (laboratories M  
49  
50 504 and N).

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53 505 “[Insert Table X about here]”  
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57 507 For the samples containing antibiotics at a concentration below the detection limit (L1), 6  
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59 508 laboratories found 100% of negative results; 3 laboratories found 1 or 2 samples which were  
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509 doubtful or positive. When the concentration was considered near the detection limits (L2),

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3 510 tylosin was generally detected (83 % of positive results), whereas the other antibiotics were  
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5 511 less detected (oxytetracycline 11%, ceftiofur 11% and sulfamethazine 17%).  
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9  
10 513 The average rate of positive samples to the L3 concentration was 81 %. The rate of detection  
11  
12 514 to the L3 concentration is 100% for the 4 antibiotics and for 3 laboratories (B, D, N). This rate  
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14 515 was 100% for the totality of the laboratories concerning tylosin and ceftiofur.  
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18 517 The results of the participating laboratories were analyzed in order to calculate different  
19  
20 518 validation parameters. A laboratory was removed from the analysis because due to a delay  
21  
22 519 in transport the samples arrived in defrosted conditions.  
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26  
27 521 Table X presents also the analysis of the results of all the participating laboratories in term of  
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29 522 reproducibility, by material (combination animal/antibiotic species) and in a global way. The  
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31 523 reproducibility, expressed as a percentage, is the ratio of the number of identical results, the  
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33 524 most common type (e.g. negative results for blank samples or positive result for samples  
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35 525 contaminated with an antimicrobial concentration exceeding the detection limit) on the total  
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37 526 number of analyses.  
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42 528 The results of the participating laboratories in term of reproducibility are very satisfactory with  
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44 529 an average percentage of 89,1%. The worse reproducibility was observed for the  
45  
46 530 combination bovine/sulfamethazine. A little higher concentration of sulfamethazine would  
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48 531 undoubtedly have given better results in term of reproducibility.  
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52  
53 533 The percentage of specificity SP for the levels L0 and L1 was calculated as:  $SP = [1 - (FP/N_-) * 100\%]$  where:  $N_-$ : total number of tests L0 and L1 and FP: number of false positive results.  
54  
55 534

56  
57 535 The percentage of sensitivity SE for each positive contamination level L2 and L3 was  
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59 536 calculated using the following equation:  $SE = (TP/N_+) * 100\%$  where:  $N_+$ : total number of  
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537 tests L2 or L3 respectively and TP: a number of true positive.

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3 538 Specificity (95,3 %) and sensitivity of the Premi®Test to the level L3 (72,5 %) were very  
4  
5 539 satisfactory.

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9 541 The repeatability was estimated in each laboratory: 1- by comparing the results of the 2 tests  
10 542 performed on each sample (2 different sets of analyses), knowing that the knowledge of the  
11 543 initial result can influence the reading at the second analysis, 2- by comparing the results  
12 544 obtained with the 2 samples of each pair. The repeatability expressed as a percentage, is the  
13 545 ratio of the number of identical results per couple of analyses on the total number of couples.  
14 546 The following table presents the total analysis of the results of all the participants obtained  
15 547 with the Premi®Test in term of repeatability. (Tables XI).

16 548 “[Insert Table XI about here]”  
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29 550 The results of the participating laboratories in term of repeatability are very satisfactory with  
30 551 an average percentage of 94,8% for the same sample and 92,7% per 2 identical samples  
31 552 (pair). The limits of detection of the Premi®Test during the preliminary study were confirmed  
32 553 by the collaborative study.  
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## 39 40 555 **Conclusion**

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42 556 This paper presents a significant dataset concerning the performance of the Premi®Test, for  
43 557 different kind of samples: spiked meat juice samples, incurred samples, routine field samples  
44 558 (confirmed by a LC/MS-MS method). The detection capabilities of Premi®Test for beta-  
45 559 lactams (amoxicillin, ceftiofur), one macrolide (tylosin) and tetracycline were at the level of  
46 560 the respective Maximum Residue Limits (MRL) in muscle samples or even lower. The  
47 561 applicability of the test to different animal species was proven. Moreover, the applicability of  
48 562 Premi®Test to routine analysis of samples was demonstrated. The Four Plate Test  
49 563 (reference method) and the Premi®Test showed comparable performances in term of  
50 564 sensitivity and specificity. Moreover, the false negative rate of Premi®Test was always lower  
51 565 than that of the FPT. This is the most important parameter to minimise for a screening

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3 566 method for antibiotic residues. On the contrary, the false positive rate of Premi®Test  
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5 567 appeared, in step 3, higher than that of the FPT. This means that the number of samples to  
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7 568 confirm by physico-chemical methods would be higher if the laboratories used the  
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9  
10 569 Premi®Test alone. However, it means also that more real positive samples would be  
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12 570 detected because Premi®Test was more sensitive than FPT for some antimicrobials (beta-  
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14 571 lactams and sulphonamides). Beta-lactams and some sulphonamides were satisfactorily  
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16 572 detected at the MRL level by the Premi®Test.  
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18 573  
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20 574 This is the first time to our knowledge that an interlaboratory study organised for the  
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22 575 detection of antibiotics with Premi®Test was described. The results of the collaborative study  
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24 576 were very satisfactory. The results of 9 laboratories were finally analyzed, plus the expert  
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26 577 laboratory. Specificity was estimated at 95.3 %, moreover the sensitivity of the test to the L3  
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28 578 level was calculated to 72,5%. These results were similar with those obtained at the time of  
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30 579 preceding validations of kits of detection of the antibiotic residues in milk. The results in term  
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32 580 of repeatability and reproducibility are very satisfactory, with average percentages of 94,8%  
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34 581 and 92,7 % for the repeatability and of 89,1 for reproducibility.  
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38 583 In conclusion, Premi®Test is easy to perform. It is ideal for an "on site" use  
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40 584 (slaughterhouses, test laboratories) as no special laboratory equipment is needed to perform  
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42 585 the test. The rapidly response "yes/no" result is simply read by colour comparison.  
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44 586 Premi®Test is applicable to the muscles of various species (porcine, bovine, ovine...), by  
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46 587 using as negative control a "blank" muscle of each analyzed specie to optimise the reading  
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48 588 time.  
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52 590 Finally, since 2006, the field laboratories in France are authorised to use the Premi®Test as  
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54 591 a pre-screening test. All positive samples with Premi®Test are then mandatory analysed by  
55  
56 592 the FPT (Anon. 2006). All positive samples with the FPT are sent to our laboratory for  
57  
58 593 confirmation as usual.  
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4

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8  
9  
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11  
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13  
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15  
16 600 participated to the preliminary study. Thanks finally to all the participants of the collaborative  
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18 601 study.  
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1 **Table I. Interlaboratory study: Content of the 16 materials sent in blind duplicate.**

		Porcine	Bovine	Porcine	Chicken
Antimicrobial	Level of contamination	Oxytetracycline	Sulfamethazine	Ceftiofur	Tylosin
MRL	/	100	100	1000	100
Spiked concentrations ( $\mu\text{g}/\text{kg}$ )	L0	'Blank'	'Blank'	'Blank'	'Blank'
	L1	20	20	40	10
	L2	200	200	400	100
	L3	400	400	800	200

2

3

4 **Table II. Results of the Premi®Test analyses on spiked juice samples (step 1).**

Antibiotic family	SULPHONAMIDE	TETRACYCLINE	MACROLIDE	BETA-LACTAM	AMINOGLYCOSIDE	BETA-LACTAM	
Antibiotic	Sulfamethazine	Oxytetracycline	Tylosin	Amoxicillin	Gentamicin	Ceftiofur	<b>Global</b>
MRL(muscle) ( $\mu\text{g kg}^{-1}$ )	100	100	100	50	50	1000	/
Tested concentrations ( $\mu\text{g kg}^{-1}$ )	50/ <b>100</b> /200	50/ <b>100</b> /200	50/ <b>100</b> /200	25/ <b>50</b> /100	<b>50</b> /100/200	100/200/400	/
Detection rate at 0.5 MRL	0%	60%	80%	100%	(0%)*	-	<b>48%</b>
Detection rate at MRL	20%	80%	100%	100%	0%	(100%)	<b>67%</b>
Detection rate at 2* MRL	100%	100%	100%	100%	40%	(100%)	<b>90%</b>
<b>Detection capability</b>	<b>2xMRL</b>	<b>2xMRL</b>	<b>MRL</b>	<b>0.5xMRL</b>	<b>&gt; 2xMRL</b>	<b>0.5xMRL</b>	/

7 \*MRL in bold character

9 \* The percentage is between brackets when there were no analyses performed at this level of concentration. i.e. All the tested concentrations for ceftiofur were below MRL and even 0.5 MRL. At 400  $\mu\text{g kg}^{-1}$ , 100 % of the results were positive or doubtful.

12 Table III. Analyses of naturally incurred samples with the 3 screening methods (step 2).

13

Antibiotic	OTC/SDMX	Amoxicillin	Tylosin	Global results of incurred samples	Blank	Fp+	Fp-
MRL (pig muscle) ( $\mu\text{g}/\text{kg}$ )	100 / 100	50	100				
Concentrations by LC/MS-MS ( $\mu\text{g}/\text{kg}$ )	760 / 150	270	750				
Number of positive results with Premi <sup>®</sup> Test	5 / 5	5 / 5	5 / 5	15 / 15	0 / 5	0 %	0 %
Number of positive results with STAR	5 / 5	5 / 5	5 / 5	15 / 15	3 / 5	60 %	0 %
Number of positive results with FPT	5 / 5	4 / 5	1 / 5	10 / 15	2 / 5	40 %	33 %

14 *Fp+*: false positive rate; *fp-*: false negative rate; OTC: Oxytetracycline; SDMX: sulfadimethoxine

15



16 Table IV. Correlation between field laboratories and AFSSA, comparison of the 3 screening  
 17 tests and LC/MS-MS method in relation to the species (step 3).

18

	Specie	Bovine	Porcine	Poultry	Others	Unknown	TOTAL
LVD	Number of samples analysed at LVD <sup>b</sup>	379	671	205	26	146	1427
	Number of positive and doubtful results at LVD Premi®Test	63	26	1	5	7	102
AFSSA	Number of samples analysed at AFSSA	65	31	4	5	7	112 <sup>a</sup>
	Number of positive and doubtful results with Premi®Test	45	19	2	5	7	78
	Correlation LVD/AFSSA (%)	69	61	50	100	100	76
	Number of positive samples with FPT	3	5	2	0	0	10
	Number of positive samples with STAR	13	13	4	0	2	32
AFSSA LC/MS-MS	Number of tested samples	38	18	4	1	0	61
	Number of positive samples (identified molecule)	14	8	3	0	0	25
	Rate of positive confirmation (%)	37	44	75	0	/	41

19

20 <sup>a</sup> Total of the samples sent to AFSSA for confirmation

21 <sup>b</sup> LVD means "Laboratoire Vétérinaire Départemental" in French which is a field veterinary laboratory.

22

23 Table V. Correlation between the Premi®Test and the FPT (step 3).

	1st analysis (112 samples)	2 <sup>nd</sup> analysis (1427 samples)
Relative accuracy AC (%)	33.9	94.8
Relative specificity SP (%)	30.4	95.0
Relative sensitivity SE (%)	70.0	70.0

24 *Relative accuracy:  $AC = (PA + NA)/N*100\%$*

25 *Relative specificity:  $SP = (NA/N-)*100\%$*

26 *Relative sensitivity:  $SE = (PA/N+)*100\%$*

27 *Where: NA is the negative agreement, PA: the positive agreement, ND: the negative discrepancy, PD: the*  
 28 *positive discrepancy;*

29  *$N = NA + Pa + PD + ND$ : total number of samples*

30 *N- is the total number of negative samples obtained with the reference method (NA + PD)*

31 *N+ is the total number of positive samples obtained with the reference method (PA + ND)*

32

33 Table VI. Results of the identification of positive samples with LC/MS-MS at the AFSSA:  
 34 Comparison of Premi®Test, Four Plate Test and STAR protocol in 2005 (step 3).

Identified AB family	Quantification / MRL	Number of analysed	Premi®Test results			FPT results		STAR results	
			-	D	+	-	+	-	+
MACRO	<MRL	1		1			1		1
SULFA	<MRL	1		1		1		1	
TTC	<MRL	10	4	3	3	6	4	2	8
TTC	>MRL	2	1		1		2		2
BL	<MRL	10		2	8	10		7	3
BL	>MRL	2		2		1	1		2
TTC + SULFA	<MRL	1		1			1		1
BL + sulfa	>MRL	1		1		1		1	
MACRO + TTC + BL	<MRL	1		1		1		1	
TTC + BL	<MRL	1		1		1		1	
AMINO + BL	<MRL	1		1		1		1	
<b>Global</b>	<LMR	26	4	7	15	20	6	13	13
	>LMR	5	1	0	4	2	3	1	4
	<b>Total</b>	<b>31</b>	<b>5</b>	<b>7</b>	<b>19</b>	<b>22</b>	<b>9</b>	<b>14</b>	<b>17</b>
Absence	/	57	5	8	44	56	1	43	14
Not analysed	/	24	24			24		24	
<b>TOTAL</b>	/	<b>112</b>	<b>34</b>	<b>15</b>	<b>63</b>	<b>102</b>	<b>10</b>	<b>81</b>	<b>31</b>

38 Table VII. Results of the identification of positive samples with LC/MS-MS at the AFSSA:  
 39 Comparison of alternative method (Premi®Test) and reference method (FPT) from 2003 to  
 40 2005 (step 3).

Identified AB family	Quantification / MRL**	Number of analysed samples	Premi®Test results			FPT results	
			-	D*	+	-	+
Beta-lactams	<MRL	9	/	1	8	8	1
	>MRL	8	/	/	8	2	6
	<b>Sum***</b>	<b>17</b>					
Tetracyclines	<MRL	13	5	4	4	6	8
	>MRL	12	5	1	6	/	11
	<b>Sum</b>	<b>25</b>					
Sulphonamides	<MRL	2	/	1	1	2	/
	>MRL	2	/	/	2	2	/
	<b>Sum</b>	<b>4</b>					
Quinolones	>MRL	3	1	/	2	/	3
BL + Sulfa	<MRL	2	/	/	2	1	1
	>MRL	2	/	/	2	2	/
	<b>Sum</b>	<b>4</b>					
Tetra + Sulfa	<MRL	1	/	/	1	/	1
	>MRL	3	/	/	3	/	3
	<b>Sum</b>	<b>4</b>					
Tetra + BL	<MRL	1	/	/	1	1	/
Macrolides	<MRL	3	1	1	1	1	2
	>MRL	3	/	/	3	1	2
	<b>Sum</b>	<b>6</b>					
Global	<MRL	31	6	7	18	19	13
	>MRL	33	6	1	26	7	25
<b>TOTAL</b>	/	<b>64</b>	<b>12</b>	<b>8</b>	<b>44</b>	<b>26</b>	<b>38</b>

42 \* D: Doubtful

43 \*\* MRL: Maximum Residue Limit

44 \*\*\* Sum of samples confirmed lower and higher than respective MRLs.

46 **Table VIII. Results of the expert laboratory (AFSSA) during the interlaboratory**  
47 **study.**

48

Level of contamination	Oxytetracycline Porcine	Sulfamethazine Porcine	Ceftiofur Bovine	Tylosin Chicken
L0	0*	0	0	2
L1	0	0	0	2
L2	4	4	4	4
L3	4	4	4	4

49 \* number of positive results

50

51 **Table IX. Raw data of the participants for each material and the 16 combinations.**

Porcine Oxytetracycline																	
		L0 (0)				L1 (20*)				L2 (200)				L3 (400)			
		Mat 1		Mat 5		Mat 2		Mat 6		Mat 3		Mat 7		Mat 4		Mat 8	
Lab	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	
A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	D
B	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+
C	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+
D	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+
E	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
F	-	-	-	-	-	-	-	-	+	+	+	+	+	+	+	+	+
G	-	-	-	-	-	-	-	-	-	-	-	-	-	D	D	D	D
M	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
N	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+

  

Porcine ceftiofur																	
		L0 (0)				L1 (40)				L2 (400)				L3 (800)			
		Mat 9		Mat 13		Mat 10		Mat 14		Mat 11		Mat 15		Mat 12		Mat 16	
Lab	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	
A	-	-	-	-	-	-	-	-	-	D	-	-	+	D	+	+	
B	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+	
C	-	-	+	+	-	-	-	-	-	-	-	-	+	+	+	+	
D	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+	
E	-	-	-	-	-	-	-	-	-	-	-	-	D	D	+	+	
F	D	D	D	D	-	-	D	D	-	D	+	+	+	+	+	+	
G	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+	
M	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+	
N	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+	

  

Bovine sulfamethazine																	
		L0 (0)				L1 (20)				L2 (200)				L3 (400)			
		Mat 17		Mat 21		Mat 18		Mat 22		Mat 19		Mat 23		Mat 20		Mat 24	
Lab	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	
A	-	-	-	-	-	-	-	-	-	-	-	-	-	D	-	-	
B	-	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	
C	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	
D	-	-	-	-	-	-	-	-	-	D	-	D	+	+	+	+	
E	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	
F	-	-	-	-	-	-	-	-	-	-	+	+	-	-	D	D	
G	-	-	-	-	-	-	-	-	-	-	-	-	-	-	D	D	
M	-	D	-	-	-	-	-	D	-	-	-	-	-	-	-	-	
N	+	-	-	-	-	-	-	-	-	-	-	-	-	D	+	D	

  

Chicken Tylosin																	
Conc	L0 (0)				L1 (10)				L2 (100)				L3 (200)				
		Mat 25		Mat 29		Mat 26		Mat 30		Mat 27		Mat 31		Mat 28		Mat 32	
Lab	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	A1	A2	
A	-	-	-	-	-	-	-	-	+	D	-	D	D	+	D	+	
B	-	-	-	-	-	-	-	-	+	+	+	+	+	+	+	+	
C	-	-	-	-	-	-	-	-	+	+	+	+	+	+	+	+	
D	-	-	-	-	-	-	-	-	+	+	+	+	+	+	+	+	
E	-	-	-	-	-	-	-	-	-	-	-	D	+	+	+	+	
F	-	-	-	-	-	-	-	-	+	+	+	+	+	+	+	+	
G	-	-	-	-	-	-	-	-	+	+	-	D	+	+	+	+	
M	-	-	-	-	-	-	-	-	D	-	D	D	+	+	+	+	
N	-	-	-	-	-	-	-	-	D	+	+	+	+	+	+	+	

52 *A1: first analysis; A2: second analysis; Mat: Material; \*Conc: concentrations (ng ml<sup>-1</sup>).*

53 **Table X. Analysis of the results of all the participating laboratories in term of**  
 54 **percentages of positive results for each material and in term of reproducibility, by**  
 55 **material (combination animal/antibiotic species) and global reproducibility.**

Material	Number of sample (number of the pair)	Level of contamination	False positive and true positive rates	Percentages of positive results for each material	Reproducibility (%)
Porcine OTC	1	L0	FP0 a)*	0%	100.0
	2	L1	FP1 b)	0%	100.0
	3	L2	TP2 c)	11%**	88.9***
	4	L3	TP3 d)	69%	69.4
Porcine ceftiofur	5	L0	FP0 a)	11%	83.3
	6	L1	FP1 b)	8%	94.4
	7	L2	TP2 c)	17%	88.9
	8	L3	TP3 d)	53%	100.0
Bovine sulfamethazine	9	L0	FP0 a)	17%	91.7
	10	L1	FP1 b)	6%	91.7
	11	L2	TP2 c)	11%	83.3
	12	L3	TP3 d)	100%	52.8
Chicken Tylosin	13	L0	FP0 a)	0%	100.0
	14	L1	FP1 b)	0%	100.0
	15	L2	TP2 c)	83%	83.3
	16	L3	TP3 d)	100%	100.0
<b>Total</b>					<b>89.1</b>

57  
58  
59 *FP : False positive rate*

60 *TP : True positive rate*

61 *a) False positive at level L<sub>0</sub>*

62 *b) False positive at level L<sub>1</sub>*

63 *c) True positive at level L<sub>2</sub>*

64 *d) True positive at level L<sub>3</sub>*

65 *\* At each level of contamination, 9 laboratories reported 4 results for 2 materials in blind duplicate. The total number of samples per level is 9\*4 = 36 samples.*

66 *\*\* TP2 c) = 4 TP / 36 samples at level L<sub>2</sub> \*100 = 11 %*

67 *\*\*\* Reproducibility at L<sub>2</sub> = 32 negative results divided by 36 samples \*100 = 88.9 %*

69 **Table XI. Global analysis of the results of all the participants in term of**  
 70 **repeatability of the analyses with the Premi®Test.**

Lab	(Number of identical results for the same sample / N)*100	(Number of identical results for 2 identical samples (pair) / N)*100
A	87.5	87.5
B	100.0	100.0
C	100.0	93.8
D	93.8	100.0
E	71.9	96.9
F	96.9	78.1
G	96.9	90.6
M	90.6	90.6
N	96.9	96.9
<b>Total</b>	<b>94.8</b>	<b>92.7</b>

73 *N: total number of samples (32)*