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Bjorn Berendsen, Ingrid Elbers, Linda Stolker

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The stability of antibiotics in matrix and reference solutions determined using a straight-forward procedure applying mass spectrometric detection

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Abstract:	The stability of an antibiotic is a very important characteristic, especially in the field of antibiotic residue analysis. During method development or validation, the stability of the antibiotic has to be demonstrated no matter if the method is used for screening, confirmation, qualitative or quantitative analysis. A procedure for testing the stability of antibiotics in solutions and food samples using LC-MS/MS is described here. The procedure is based upon the assumption that the antibiotics are stable when stored at -70 °C. Representative solutions or spiked samples containing the antibiotic are stored at the temperature to be tested (-18 or 4 °C) and at -70 °C. After a selected storing time samples are moved from the chosen storage temperature to -70 °C. At the end of the study, all samples -per class of antibiotic- are analysed in one batch. By applying statistical models it is finally concluded at which circumstances the antibiotic is stable. The stability of 60 antibiotics belonging to the classes of tetracyclines, sulphonamides, quinolones, penicillins, macrolides and aminoglycosides are tested. The stability of solutions containing tetracyclines and penicillins is only guaranteed for three months while stored at -18 °C. Solutions of all other antibiotics tested are stable for at least 6 or 12 months when

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	stored at 4 °C. In muscle tissue stored at -18 °C no severe degradation of the tested antibiotics was observed with the exception of the penicillins. The presented stability data are useful as a reference for laboratories carrying out validation studies of analytical methods for antibiotic (residue) detection. It saves them time needed for long term stability testing of solutions and samples.

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**The stability of antibiotics in matrix and reference solutions
determined using a straight-forward procedure applying mass
spectrometric detection**

B.J.A. Berendsen*, I.J.W. Elbers, A.A.M. Stolker

*RIKILT – Institute of Food Safety, Wageningen UR (University and Research centre), Akkermaalsbos
2, 6708WB, P.O. Box 230, 6700AE, Wageningen, The Netherlands.*

*Corresponding author: Email: bjorn.berendsen@wur.nl

Abstract

The stability of an antibiotic is a very important characteristic, especially in the field of antibiotic residue analysis. During method development or validation, the stability of the antibiotic has to be demonstrated no matter if the method is used for screening, confirmation, qualitative or quantitative analysis. A procedure for testing the stability of antibiotics in solutions and food samples using LC-MS/MS is described here. The procedure is based upon the assumption that the antibiotics are stable when stored at -70 °C. Representative solutions or spiked samples containing the antibiotic are stored at the temperature to be tested (-18 or 4 °C) and at -70 °C. After a selected storing time samples are moved from the chosen storage temperature to -70 °C. At the end of the study, all samples -per class of antibiotic- are analysed in one batch. By applying statistical models it is finally concluded at which circumstances the antibiotic is stable. The stability of 60 antibiotics belonging to the classes of tetracyclines, sulphonamides, quinolones, penicillins, macrolides and aminoglycosides are tested. The stability of solutions containing tetracyclines and penicillins is only guaranteed for three months while stored at -18 °C. Solutions of all other antibiotics tested are stable for at least 6 or 12 months when stored at 4 °C. In muscle tissue stored at -18 °C no severe degradation of the tested antibiotics was observed with the exception of the penicillins. The stability data reported here are useful as a reference for laboratories carrying out validation studies of analytical methods for antibiotic (residue) detection. The data should save time needed for long term stability testing of solutions and samples.

Keywords: stability, degradation, antibiotics, mass spectrometry

Introduction

Nowadays, many antibiotics are used in animal production, particularly in intensive animal rearing like pigs, poultry and veal calves. For food safety and prevention of antibiotic resistance only the use of registered antibiotics is allowed and maximum residue limits (MRLs) in food products are established in EU/37/2010 (2010) to protect the consumer from being exposed to antibiotics. To monitor at these MRLs sensitive quantitative analytical methods are needed. In most cases, for the quantitative analysis, solutions of antibiotics are used as reference standards. To obtain a correct quantification, it is very important that knowledge is available on the stability of the antibiotic in the solution.

Next to the stability of the antibiotics in reference solutions it is also important to have knowledge regarding the stability of the antibiotic in the sample material. Only then suitable storage conditions can be chosen in case the analysis cannot take place immediately

after sample collection. To obtain information on the stability of antibiotics in matrix a suitable quantitative analytical method has to be available. Additionally a statistical procedure for the data evaluation is mandatory.

It is obvious that characterisation of the method of analysis is very important. For that reason the validation of analytical testing methods is a primary requirement ISO 17025 (2005) accredited laboratories. Guidelines for the validation of the analysis of antibiotics in food matrices are established in EC/2002/657 (2002) and these state that for qualitative and quantitative methods, may it be either screening or confirmatory, the stability of the analyte in solution and in matrix are main characteristics to be determined. Analyte stability information has to be obtained either from experimental data or from literature. Unfortunately stability testing can be time-consuming and only limited literature is available about antibiotic stability.

Some approaches for stability testing of reference solutions and samples are reported (Okerman 2007, Croubels 2003, Jiménez 2004). Okerman *et al.* (2007) apply microbial techniques to determine the decrease in microbial activity over a period of six months. Croubels *et al.* (2003) recommend ultra violet (UV) detection for analytes possessing good UV absorbing properties because the between-day variation of the UV signal is low compared to mass spectrometric (MS) detection. Jiménez *et al.* (2004) applied several detection techniques among which UV and gas chromatography in combination with MS detection of trimethylsilyl derivatives. These techniques, but especially microbial inhibition and derivatization techniques, are non-specific and thus degradation products might not be distinguishable from the native compound. Liquid chromatography coupled to triple quadrupole mass spectrometry (LC-QqQ/MS) is more likely to discriminate degradation products from the native compound and is therefore the preferred detection technique if day to day variation is overcome. Another disadvantage of the reported approaches is that analyses are carried out on different occasions (Croubels 2003, Jiménez 2004). Therefore fresh reference solutions are to be prepared often or, in case of matrix stability testing, a sample preparation has to be carried out several times. This introduces additional errors and is less time and thus cost efficient.

The publically available information on the stability of reference solutions of veterinary drugs is limited (Okerman 2007, Mathijssen 2010, Chédru-Legros 2010). The availability of stability information of antibiotics in animal matrices is even worse (Verdon 2000). According to 2002/657/EC (2002) the stability in matrix is preferably tested in incurred materials, but if these are not available the use of fortified materials is acceptable.

In this paper we present a straight-forward and time efficient procedure for stability testing of antibiotics in reference solutions and muscle matrix using LC-QqQ/MS as a specific detection technique. This method was applied to determine the stability of solutions of 60 antibiotics used in veterinary practice belonging to the tetracycline (n=4), sulfonamide (n=16),

91 quinolone (n=10), macrolide (n=11), penicillin (n=8) and aminoglycoside (n=11) group.
92 Furthermore, the stability of a selection of 12 antibiotics, representative for different
93 compound groups, in muscle matrix was studied. The results of this research can be used as a
94 reference for the validation of testing methods aiming for the analysis of antibiotics.

96 **Material and methods**

98 *Chemicals, reagents and solutions*

99 HPLC grade methanol and acetonitrile (Biosolve, Valkenswaard, The Netherlands), 25% and
100 32% ammonia, ammonium formate, formic acid, citric acid monohydrate, disodium
101 hydrogenphosphate dihydrate, Na₂EDTA, potassium dihydrogen phosphate, trichloro acetic
102 adic, sodium hydroxide (Merck, Darmstadt, Germany), piperidine (>99%) and
103 heptafluorobutyric acid (HFBA) (Sigma-Aldrich, St. Louis, MO, USA) were used. Milli-Q
104 water was prepared using a Milli-Q system at a resistivity of at least 18 MΩ cm⁻¹
105 (Millipore, Billerica, MA, USA). EDTA-McIlvain buffer was prepared by mixing 500 mL
106 citric acid solution (0.1 M, 21.0 g citric acid mono hydrate) with phosphate buffer (0.2 M,
107 35.6 g Na₂HPO₄ dihydrate in 1 L of water) until a pH of 4.0 is obtained. 74.4 g Na₂EDTA is
108 added and the volume adjusted to 2 L with water. Potassiumphosphatebuffer containing
109 EDTA and trichloro acetic acid was prepared by dissolving 1.36 g KH₂PO₄, 0.15 g Na₂EDTA
110 and 20.0 g trichloro acetic acid in 1 L water.

111 The following reference standards (purity is mentioned between brackets) were all
112 obtained from Sigma-Aldrich: *Tetracyclines*: oxytetracycline (99%), tetracycline (98%),
113 chlortetracycline (93%) and doxycycline (99%). *Sulfonamides*: sulfadiazine (99.6%),
114 sulfathiazole (100%), sulfapyridine (>99%), sulfamerazine (99.8%), sulfamoxole (98.5%),
115 sulfadimidine (also called sulfamethazine or sulfadimerazine, 100%), sulfamethizole (99.9%),
116 dapsone (98.4%), sulfamethoxypyridazine (99.6%), sulfamonomethoxine (98.5%),
117 sulfachloropyridazine (99.4%), sulfadoxine (99.9%), sulfamethoxazole (100%), sulfisoxazole
118 (99.6%), sulfadimethoxine (>99%) and sulfaquinoxaline (>91.6%). *Quinolones*:
119 marbofloxacin (98.8%), norfloxacin (99%), ciprofloxacin (99.9%), enrofloxacin (99.6%),
120 danofloxacin (99.9%), sarafloxacin (97.2%), difloxacin (99.8%), oxolinic acid (99%),
121 nalidixic acid (99.4%) and flumequine (99%). *Penicillins*: amoxicillin (85.6%), ampicillin
122 (96.8%), penicillin G (also called benzylpenicillin, 99.9%), penicillin V (also called
123 phenoxymethylpenicillin, 98.3%), cloxacillin (94.7%), dicloxacillin (95.6%), nafcillin (87.6%)
124 and oxacillin (89.2%). *Aminoglycosides*: apramycin (71.7%), dihydrostreptomycin (75.1%),
125 gentamycin (including C1, C1A, C2 and C2A, 59.5%), kanamycin (77.7%), neomycin
126 (82.5%), paromomycin (82.5%), spectinomycin (69.7%) and streptomycin (76%). *Macrolides*:

tilmicosin (83.7%), tiamulin (99.9%), erythromycin (65.7%), tylosin A (101.3%), valnemulin (>95%), josamycin (101.5%), lincomycin (93%) and spiramycin I (92.6%). Three other macrolides were obtained elsewhere: tulathromycin (97.6%) from Pfizer (New York, NY, USA), aivlosin (also called 3-acetyl-4"-isovaleryl tylosin, 67.9%) from Eco Animal health (London, UK) and pirlimycin (90.3%) from Pharmacia and Upjohn Company (Bridgewater, NJ, USA).

Stock solutions were prepared by accurately weighing in 3 to 6 mg (± 0.02 mg) of reference standard. After correction for purity and counter-ions present of the amount of reference standard taken, it was dissolved in solvent (on weight basis) to obtain the required concentration. Separate methanolic stock solutions were prepared at a concentration of 100 mg L⁻¹ of tetracyclines and penicillins and at a concentration of 1000 mg L⁻¹ for sulfonamides. Separate stock solutions of 1000 mg L⁻¹ were prepared in acetonitrile for all macrolides and in water for the aminoglycosides. Separate stock solutions of quinolones were prepared by dissolving the analyte in 2M ammonia using sonification (30 min) to obtain a concentration of 5000 mg/L after which the solution was diluted to 100 mg L⁻¹ with methanol.

Instrumentation

The liquid chromatography (LC) instrumentation used was an Acquity UPLC system (Waters, Milford, MA, USA) and detection was carried out using a triple quadrupole Quattro Premier mass spectrometer (Waters) (QQQ/MS) operating with electrospray ionization (ESI) in positive mode. The operating parameters were: capillary voltage, 2.5 kV; cone voltage, see Table 1; source temperature, 120 °C; desolvation temperature, 300 °C; cone gas flow, 200 L hr⁻¹; desolvation gas, 500 L hr⁻¹. Data were acquired and processed using MassLynx 4.1 software (Waters).

LC-QQQ/MS analysis

Several multi-compound methods were applied to determine the stability of the antibiotic substances. All analytical methods used for stability testing were fully validated as described in 2002/657/EC (2002) as a quantitative confirmatory method and fulfil the criteria established for trueness, repeatability, within-laboratory reproducibility, robustness and selectivity.

Tetracyclines

The analysis was carried out according to a previously described method (Berendsen 2006).

Sulfonamides

Muscle (1 g) was extracted with 10 mL of water. After mechanical shaking and centrifugation the aqueous extract was filtered using a 30 kD ultrafilter (Millipore) and the filtrate was transferred into a glass vial.

Chromatographic separation was established using a Waters Acquity UPLC BEH C₁₈ analytical column, 50 x 2.1 mm, 1.7 µm. The gradient (mobile phase A, 5 mM formic acid in water; mobile phase B, 5 mM formic acid in water/acetonitrile (1:9, v/v), flow rate 0.8 mL min⁻¹) was: 0-0.5 min, 0% B; 0.5-5.5 min, linear increase to 15% B; 5.5-6.6 min, linear increase to 25% B; 6.6-6.7 min, linear increase to 100% B; 6.7-7.2 min, 100% B; 7.2-7.3 min, return to 0% B. The flow was split 1:1 (MS:waste). The injection volume was 40 µL. Fragmentation was carried out using collision induced dissociation (CID) with the settings presented in Table 1.

Quinolones

Muscle (1 g) was extracted with 10 mL of water. After mechanical shaking and centrifugation the aqueous extract was filtered using a 30 kD ultrafilter (Millipore) and the filtrate was transferred into a glass vial.

A chromatographic separation was established using a Waters Symmetry C₁₈ analytical column, 150 x 3 mm, 5 µm. The gradient (mobile phase A, 5 mM formic acid in water; mobile phase B, 5 mM formic acid in water/acetonitrile (1:9, v/v), flow rate 0.4 mL min⁻¹) was: 0-1 min, 0% B; 1-11 min, linear increase to 100% B; 11-13 min, 100% B; 13-14 min, return to 0% B. The injection volume was 50 µL. Fragmentation was carried out using CID with the settings presented in Table 1.

Macrolides

Muscle (2 g) was extracted with 20 mL EDTA-McIlvain bufffer. After mechanical shaking and centrifugation 10 mL of the aqueous extract was transferred onto a conditioned OASIS® MCX 3CC, 60 mg solid phase extraction cartridge (Waters). The cartridge was washed with 3 mL MeOH / water / 32% ammonia (10:85:5 v/v) and dried under vacuum. The macrolides were eluted from the cartridge using 3 ml MeOH / 32% ammonia (95:5 v/v). The solvent was evaporated (45°C, N₂) and the residue reconstituted in 700 µL 2% ACN in 10 mM ammonium formate buffer, adjusted to pH=4.0 with formic acid. The extract was transferred into a glass vial.

A chromatographic separation was established using a Waters Atlantis dC₁₈ analytical column, 150 x 3 mm, 5 µm. The gradient (mobile phase A, 10 mM ammonium formate buffer, pH adjusted to 4.0 with formic acid; mobile phase B, 10 mM ammonium formate buffer in water/acetonitrile (1:9 v/v), adjusted to pH=4.0 using formic acid; flow rate 0.4 mL min⁻¹) was: 0-1 min, 30% B; 1-17 min, linear increase to 100% B; 17-18 min, 100% B; 18-19 min,

200 return to 30% B. The flow was split 2:1 (MS:waste). The injection volume was 50 μ L.

201 Fragmentation was carried out using CID with the settings presented in Table 1.

202

203 *Penicillins*

204 Muscle samples were treated according to a previously reported method (van Holthoon 2010).

205 Before analysis 100 μ L of each mixed standard solution was transferred into a 4 mL glass

206 container. The solvent was evaporated under nitrogen at 45°C after which 1.0 mL 2%

207 piperidine (Sigma-Aldrich) solution in water was added to dissolve the residue. The

208 penicillins were derivatized to penicillin piperidines at room temperature during 30 minutes

209 after which 100 μ L 10% acetic acid (Merck, Darmstadt, Germany) in water and 2.0 mL 100

210 mM ammonium acetate buffer, pH=6.0, were added. These solutions were injected into the

211 previously reported LC-MS/MS system (van Holthoon 2010).

212

213 *Aminoglycosides*

214 Muscle (2 g) was extracted with 20 mL 10 mM potassiumphosphatebuffer pH=4.0 containing

215 EDTA and trichloroacetic acid. After mechanical shaking and centrifugation the pH of the

216 supernatant was adjusted to pH=7.7 using sodium hydroxide (30% in water). The aqueous

217 extract was transferred onto a conditioned CBX 6CC, 500 mg solid phase extraction cartridge

218 (Avantor, Phillipsburg, NJ, USA). The cartridge was washed with 4 mL water and dried under

219 vacuum. The aminoglycosides were eluted from the cartridge using 10% acetic acid in MeOH.

220 The solvent was evaporated (60°C, N₂) and the residue reconstituted in 400 μ L 0.065%

221 HFBA in water. The extract was transferred into a glass vial.

222 A chromatographic separation was established using a Waters Symmetry C₁₈

223 analytical column, 150 x 3 mm, 5 μ m. The gradient (mobile phase A, 0.065% HFBA in water;

224 mobile phase B, 0.065% HFBA in methanol; flow rate 0.4 mL min⁻¹) was: 0-0.5 min, 0% B;

225 0.5-5.5 min, linear increase to 45% B; 5.5-17.0 min, linear increase to 60% B; 17-22 min,

226 60% B; 22-23 min, return to 0% B. The flow was split 1:1 (MS:waste). The injection volume

227 was 20 μ L. Fragmentation was carried out using CID with the settings presented in Table 1.

228 Using this procedure gentamycin C2 and C2A cannot be distinguished.

229

230 Note: please insert Table 1 here

231

232

233 *Methodology*

234 *Stability of antibiotics in solutions*

235 Of each stock solution of quinolones, macrolides and aminoglycosides 1 mL was transferred

236 into ten different glass containers. Of the sulfonamide stock solutions a mixed standard

237 solution of 10 mg L⁻¹ in methanol was prepared of which 1 mL was transferred into ten
238 different glass containers. For each solution at t=0, two containers were stored at -70 °C and 8
239 at 4 °C. After 1, 3, 6 and 12 months each, two containers were moved from 4 °C to -70 °C.
240 After 12 months, all containers were placed at room temperature to warm up and the solutions
241 were diluted to suitable concentrations with water (aminoglycosides in 0.065% HFBA in
242 water) in duplicate (I and II) to prevent overloading of the analytical system. This experiment
243 is schematically presented in Figure 1. These solutions were analysed using LC-QqQ/MS in
244 the following order: I t=0, I t=12, I t=6, I t=3, II t=0, II t=12, II t=6 and II t=3. This series was
245 repeated four times resulting in five injections per dilution. For quinolones this experiment
246 was carried out for a six month period only. For tetracyclines stock solutions and penicillins
247 mixed standard solutions the experiment was carried out with transfer of containers at t=1
248 week, t=2 weeks, t=1 month, t=2 months and t=3 months at both 4 °C and -18 °C because of
249 expected instability.

250 After integration of the peaks, for each storage time, the average peak area of dilution
251 I was compared with dilution II using a Students' t-test (n=5) to check if the dilutions were
252 carried out correctly. If no statistical difference was observed between both dilutions the
253 overall average peak area of dilution I and II was calculated (n=10) for each storage time.
254 Next, the average of each storage time was compared to the average peak area of the solution
255 stored at -70 °C from the beginning of the experiment (t=0), again using a Students' t-test. If
256 the average peak area at a certain storage time is above 90 % of the average peak area at t=0
257 the compound is considered stable for that specific storage time. If it drops below 90% the
258 solution was considered to be unstable. If a difference between dilution I and II was found, a
259 worst case scenario was adopted by using the most deviating average of dilution I or II in the
260 calculations.

261 With this approach the instability of a compound is observed by comparing a signal
262 obtained for a solution stored at -70 °C with a solution stored at 4 °C or -18 °C. It is assumed
263 that the antibiotics tested are stable at -70 °C (BCR/01/97 1997), however even if an analyte
264 is unstable at -70 °C a difference in signal is expected because it will by definition be more
265 unstable at higher temperatures.

266

267 Note: please insert Figure 1 here

268

269 *Stability of antibiotics in matrix*

270 The stability of antibiotics in muscle matrix was studied using in-house reference material
271 that was prepared for each compound group individually. For each compound group one or
272 more representative substances were selected. A bulk of blank muscle material was fortified
273 with the selected antibiotics at and around the MRL (EU/37/2010, 2010). Each of the

materials was homogenized under cryogenic conditions and the homogeneity of the materials was tested according to The International Harmonized Protocol for Proficiency Testing of Analytical Laboratories (Thompson 2006) and ISO 13528 (2005), taking into account the insights discussed by Thompson (2000).

Just after preparation of the materials, 12 samples of each material were randomly selected of which 6 were stored at -70 °C and 6 at -18 °C. It is assumed that the antibiotics tested are stable at these storage conditions. After the selected test period all 12 samples were analysed in one batch in random order. After integration of the peaks, the overall average peak area and relative standard deviation (VC%, n=6) were calculated for each storage time and a Students' t-test was applied to test for statistical significant differences between the samples stored at -70 °C and the samples stored at -18 °C.

Results and discussion

With the presented approach, for the analysis of the stability of antibiotics in reference solutions and muscle matrix all analyses for a specific class of antibiotics are carried out within one day using LC-MS/MS as the detection technique. Therefore, additional variation due to day to day variation of the instrumentation is prevented. Because the applied detection technique is highly selective, interference of the detector's response caused by degradation products or other interferences is unlikely and only the response of the native compound is recorded. Therefore, this method allows straight-forward, effective and cost efficient stability testing.

Stability in solution

The results of the stability study of all tested antibiotics in reference solutions, stored at 4 °C, are presented in Table 2. Stability results of the tetracycline and penicillin reference solutions, stored at -18 °C are presented in Table 3. The relative response is calculated by dividing the average response at t=1 by the average response at t=0. A compound is considered instable if the relative response drops below 90 %.

Note: please insert Table 2 here

Note: please insert Table 3 here

It is observed that the results for the tetracyclines vary severely for the different storage times and as a result, for tetracycline at t=1 week and chlortetracycline at t=2 months a result is found below the established criterion whereas the next result is again above this criterion. As a result a higher uncertainty in the establishment of the maximum storage time is obtained and a conservative estimation should be made to prevent the use of degraded reference solutions.

Nevertheless, it is clearly observed that especially oxytetracycline and tetracycline are instable in methanolic solutions which was also demonstrated before (Okerman 2007). This research clearly illustrates that tetracycline and oxytetracycline stock solutions should be prepared fresh on a weekly basis if they are stored at 4 °C and every two months (tetracycline) or every three months (other tetracyclines) if stored at -18 °C. The difference in stability of oxytetracycline and tetracycline compared to chlortetracycline and doxycycline is confirmed by Okerman *et al.* (2007). They reported no loss of activity for chlortetracycline and doxycycline during 6 months storage at -20 °C whereas oxytetracycline and tetracycline loose approximately 25% of their activity in the same period. It is noted that in our experiments the degradation of oxytetracycline and tetracycline is more severe. This difference might be explained due to the fact that Okerman *et al.* (2007) monitored the total antibiotic activity (this might include metabolites) instead of the level of the native compounds or because the concentration of the reference solutions were a factor 10 lower in our experiments.

Next to the tetracyclines, the penicillins show severe instability. During storage at 4 °C at least a 45% degradation is observed within one week for all compounds. If these penicillin solutions are stored at -18 °C, the stability is much more prolonged: under these conditions ampicillin and penicillin V solutions are stable for at least two months, the other penicillin solutions for at least three months. This is not in agreement with the results of Okerman *et al.* (2007) who reported no degradation of aqueous ampicillin solutions for at least 6 months when stored at -20 °C and, in the contrary, approximately a 20% degradation of penicillin G and amoxicillin within three months. These differences might be explained due to the fact that they monitored the total antibiotic activity instead of the level of the native compounds, because the concentration of the reference solutions are a factor 10 lower in our experiments or because different solvents were used: Okerman *et al.* (2007) prepared stock solutions in water, whereas we tested the stability in methanolic solutions. The latter is most likely, because instability of penicillins in the presence of methanol was also reported by Mastovska *et al.* (2008). For oxacillin the results are in agreement with Mathijssen *et al.* (2010) who reported no degradation for aqueous oxacillin solutions during 6 months when stored at -20 °C.

The other tested antibiotics showed to be stable for the tested period of time at 4 °C indicating that quinolones are at least stable for 6 months and the other antibiotics for at least 12 months.

Stability in matrix

The results of the stability study of the antibiotics in matrix, stored at -18 °C are presented in Table 4.

348 Note: please insert Table 4 here

349 Note: please insert Table 5 here

350

351 From the results it is concluded that all tested antibiotics, with the exception of ampicillin and
352 to less extend cloxacillin, are stable in muscle matrix when stored at -18 °C for the tested
353 period of time. Verdon *et al.* (2000) indicated no degradation of amoxicillin in porcine muscle
354 during storage at -20 °C for 3 months and severe degradation after 8 months of storage. From
355 the results of the stability of solutions, for ampicillin and amoxicillin a similar degradation
356 rate is expected and thus the quick degradation of ampicillin is unexpected. This observation
357 might be explained by a between muscle variation and therefore we recommend to store
358 samples containing penicillins at -70 °C even if the storage period is limited.

359 For the penicillin antibiotics the same experiment was carried out after stabilizing the
360 muscle matrix by adding 2 mL phosphate buffer (0.2 M, pH=6) to 2 g minced muscle (van
361 Holthoon 2010). The results of the stability study for the buffered muscle matrix containing
362 ampicillin and cloxacillin are presented in Table 5. It is concluded that after addition of
363 phosphate buffer pH=6 cloxacillin as well as ampicillin are stable in muscle matrix for at least
364 3 months. This is in agreement with previously reported results by Kondrat'eva *et al.* (1967)
365 and Lu *et al.* (2008) who showed prolonged stability of penicillin G, oxacillin and penicillin
366 V solutions at pH=6 to 7. The stabilisation of muscle samples containing penicillins is
367 recommended if no -70 °C storage capacity is available.

368

369 Conclusions

370 A straight-forward procedure is reported for the stability testing of antibiotics in solution and
371 in muscle samples using LC-MS/MS detection. The method proved to be straight-forward,
372 effective and cost efficient. From this research stability information is obtained that can be
373 used for the construction of a validation dossier. Severe instability at 4 °C was observed for
374 oxytetracycline, tetracycline and all tested penicillin solutions only. When stored at -18 °C the
375 stability of these solutions is prolonged; at least 2 months for tetracycline, ampicillin and
376 penicillin V, and at least 3 months for oxytetracycline and the other tested penicillins.
377 Methanolic solutions of chlortetracycline and doxycycline are stable for at least 3 months
378 when stored at 4 °C. Methanolic solutions of the tested sulfonamides, solutions of macrolides
379 in acetonitrile and aqueous solutions of the aminoglycosides are stable for at least 12 months
380 when stored at 4 °C. Alkaline methanolic solutions of the tested quinolones are stable for at
381 least 6 months when stored at 4 °C.

382 The tested antibiotics remain stable in muscle matrix for at least 3 months with the
383 exception of the penicillins. A significant degradation was observed within 3 months for
384 ampicillin and cloxacillin when stored at -18 °C. After stabilizing these muscle samples at

pH=6, the stability of the penicillins in matrix is prolonged to at least 3 months. Therefore it is recommended to store penicillin containing samples at -70 °C or to stabilize them at pH=6. From the work presented here, a huge amount of stability data is obtained. However, stability data is never complete, because numerous compound matrix combinations (including all species) are to be studied and new antibiotic substances become available regularly. Therefore, we suggest that an open access, easy searchable database should be created in which stability data, gathered by various laboratories, can be stored.

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- 440 Figure 1. Schematic representation of the stability study.
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- 442 Table 1. Precursor ions, collision energy and product ions of the antibiotics.
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- 444 Table 2. Stability of antibiotics in solution stored at 4°C.
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- 446 Table 3. Stability of antibiotics in solution stored at -18°C.
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- 448 Table 4. Stability of antibiotics in muscle stored at -18 °C.
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- 450 Table 5. Stability of penicillins in muscle stored at -18 °C after buffering at pH=6.
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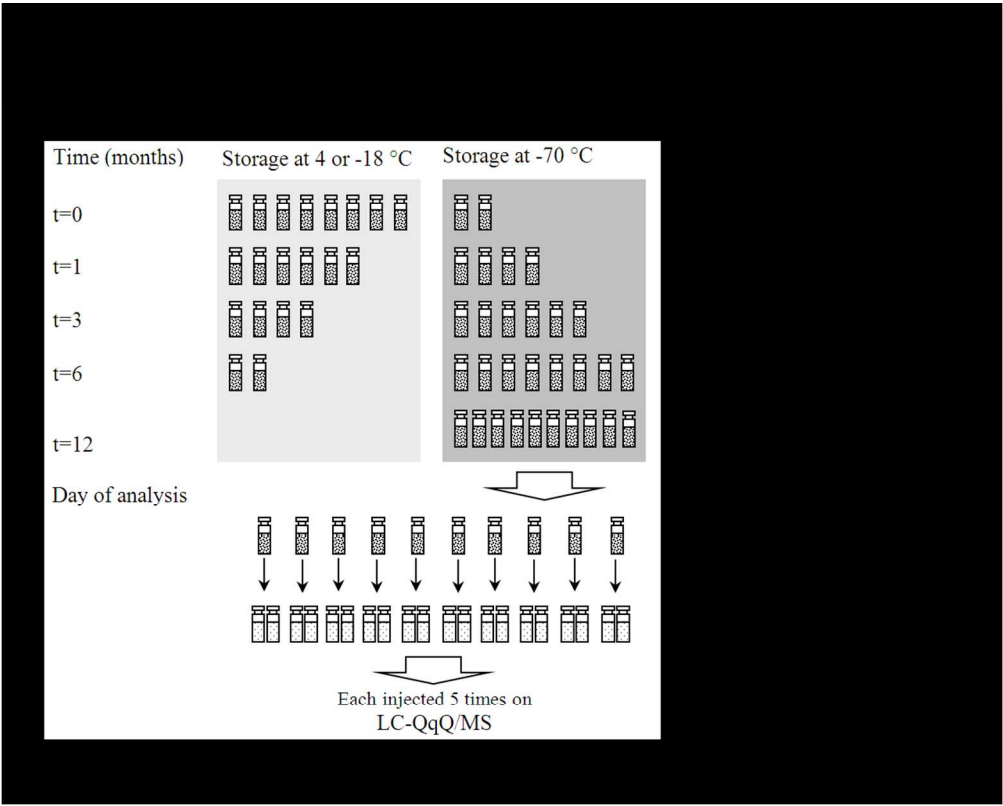


Figure 1. Schematic representation of the stability study.

Table 1. Precursor ions, collision energy and product ions of the antibiotics.

Compound	Cone (V)	Precursor ion (m/z)	Collision energy (eV)	Product ion (m/z)
<i>Sulfonamides</i>				
Sulfadiazine	24	251.2	15	92.2
Sulfathiazole	26	256.2	15	156.1
Sulfapyridine	26	250.2	30	92.2
Sulfamerazole	28	265.2	30	92.2
Sulfamoxole	23	268.2	15	156.1
Sulfadimidine	28	279.2	30	92.2
Sulfamethizole	24	271.1	15	156.1
Dapsone	30	249.2	15	156.1
Sulfamethoxypyridazine	25	281.2	15	156.1
Sulfamonomethoxine	30	281.2	20	156.1
Sulfachloropyridazine	23	285.2	15	156.1
Sulfadoxine	24	311.2	20	156.1
Sulfamethoxazole	22	254.2	25	92.2
Sulfisoxazole	21	268.2	15	156.1
Sulfadimethoxine	28	311.2	20	156.1
Sulfaquinoxaline	30	301.2	15	156.1
<i>Quinolones</i>				
Marbofloxacin	20	363.1	17	72.4
Norfloxacin	20	320.1	18	276.1
Ciprofloxacin	20	332.1	21	288.3
Enrofloxacin	20	360.2	20	316.1
Danofloxacin	20	358.2	35	340.1
Sarafloxacin	20	386.1	23	342.1
Difloxacin	20	400.1	21	356.2
Oxolinic acid	20	262.1	25	244.1
Nalidixic acid	20	233.1	20	215.2
Fumequine	20	262.1	24	244.1
<i>Macrolides</i>				
Tilmicosin	25	435.4	18	695.6
Tiamulin	25	494.3	28	192.2
Erythromycin	25	734.5	25	158.1
Tylosin A	40	916.6	32	174.2
Valnemulin	25	565.4	28	263.2
Josamycin	30	828.5	32	174.2
Lincomycin	28	407.2	21	126.0
Spiramycin I	20	422.4	20	100.9
Tulathromycin	20	403.9	18	158.1
Aivlosin	40	1042.7	42	108.9
Pirlimycin	25	411.2	19	112.0
<i>Aminoglycosides</i>				
Apramycin	25	540.3	20	378.2
Dihydrostreptomycin	35	584.3	30	263.2
Gentamycin C1	20	478.3	15	322.2
Gentamycin C1A	20	450.3	17	322.2
Gentamycin C2 or C2A*	20	464.3	17	322.2

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Gentamycin C2 or C2A*	25	485.2	25	163.1
Kanamycin	30	615.3	35	161.1
Paromomycin	30	616.3	30	163.1
Spectinomycin*	30	351.2	25	100.0
Streptomycin	50	582.3	35	263.2

* analysed as a hydrate salt.

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Table 2. Stability of antibiotics in solution stored at 4°C.

Compound	Relative response compared to t=0 (%)					Final concluded stability
	t=1w	t=2w	t=1m	t=2m	t=3m	
Tetracyclines						
Oxytetracycline	91	101	45	85	74	2 weeks
Tetracycline	86	93	69	56	45	< 1 week
Chlortetracycline	104	109	92	85	104	≥ 2 months
Doxycycline	107	119	94	102	114	≥ 3 months
Penicillins						
Amoxicillin	31	11				< 1 week
Ampicillin	36	15				< 1 week
Penicillin G	49	27				< 1 week
Penicillin V	30	9				< 1 week
Cloxacillin	38	14				< 1 week
Dicloxacillin	29	9				< 1 week
Nafcillin	54	30				< 1 week
Oxacillin	42	19				< 1 week
	Relative response compared to t=0 (%)					
	t=1m	t=3m	t=6m	t=12m		
Sulfonamides						
Sulfadiazine	99	101	97	98		≥ 12 months
Sulfathiazole	100	101	102	99		≥ 12 months
Sulfapyridine	100	101	98	101		≥ 12 months
Sulfamerazole	100	100	98	100		≥ 12 months
Sulfamoxole	100	96	100	95		≥ 12 months
Sulfadimidine	100	101	100	102		≥ 12 months
Sulfamethizole	101	100	98	101		≥ 12 months
Dapsone	102	102	99	102		≥ 12 months
Sulfamethoxypyridazine	101	102	100	101		≥ 12 months
Sulfamonomethoxine	101	103	99	101		≥ 12 months
Sulfachloropyridazine	101	104	101	102		≥ 12 months
Sulfadoxine	99	101	99	101		≥ 12 months
Sulfamethoxazole	102	102	98	101		≥ 12 months
Sulfisoxizole	100	100	97	99		≥ 12 months
Sulfadimethoxine	98	99	94	99		≥ 12 months
Sulfaquinoxaline	100	101	100	100		≥ 12 months
Quinolones						
Marbofloxacin			97			≥ 6 months
Norfloxacin			100			≥ 6 months
Ciprofloxacin			98			≥ 6 months
Enrofloxacin			102			≥ 6 months
Danofloxacin			106			≥ 6 months
Sarafloxacin			100			≥ 6 months
Difloxacin			101			≥ 6 months
Oxolinic acid			102			≥ 6 months
Nalidixic acid			101			≥ 6 months
Fumequine			103			≥ 6 months
Aminoglycosides						
Apramycin	97	95	100	98		≥ 12 months

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Dihydrostreptomycin	96	98	100	99	≥ 12 months
Gentamycin C1	99	98	99	100	≥ 12 months
Gentamycin C1A	97	97	99	99	≥ 12 months
Gentamycin C2 or C2A*	99	99	100	98	≥ 12 months
Gentamycin C2 or C2A*	98	98	100	98	≥ 12 months
Kanamycin	100	99	99	99	≥ 12 months
Neomycin	101	100	100	99	≥ 12 months
Paromomycin	99	93	99	98	≥ 12 months
Spectinomycin	102	98	97	99	≥ 12 months
Streptomycin	96	99	92	94	≥ 12 months
<i>Macrolides</i>					
Tilmicosin			96	97	≥ 12 months
Tiamulin			114	105	≥ 12 months
Erythromycin			103	112	≥ 12 months
Tylosin A			98	98	≥ 12 months
Valnemulin			108	105	≥ 12 months
Josamycin			109	104	≥ 12 months
Lincomycin			99	98	≥ 12 months
Spiramycin I			102	109	≥ 12 months
Tulathromycin			110	100	≥ 12 months
Aivlosin			116	110	≥ 12 months
Pirlimycin			92	96	≥ 12 months

w = weeks, m=months

The results below 90 % are indicated in bold.

* Gentamycin C2 and C2A cannot be distinguished with the applied methodology.

Table 3. Stability of antibiotics in solution stored at -18°C.

Compound	Relative response compared to t=0 (%)					Final concluded stability
	t=1w	t=2w	t=1m	t=2m	t=3m	
Tetracyclines						
Oxytetracycline	101	111	96	107	116	≥ 3 months
Tetracycline	109	103	104	95	87	2 months
Chlortetracycline	99	117	106	97	100	≥ 3 months
Doxycycline	115	111	111	104	109	≥ 3 months
Penicillins						
Amoxicillin	95	110	92	91	91	≥ 3 months
Ampicillin	95	109	93	93	89	2 months
Penicillin G	95	107	96	98	94	≥ 3 months
Penicillin V	90	110	90	95	88	2 months
Cloxacillin	95	108	96	98	96	≥ 3 months
Dicloxacillin	95	109	97	98	94	≥ 3 months
Nafcillin	95	106	98	100	96	≥ 3 months
Oxacillin	92	107	94	98	94	≥ 3 months

w = weeks, m=months

The results below 90 % is indicated in bold.

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Table 5. Stability of penicillins in muscle stored at -18 °C after buffering at pH=6.

Compound group	Compound	Species	t (days)	Level [VC%] (µg/kg)		Stable?
				Day 0	Day t	
Penicillins	Ampicillin	Porcine	76	4.5 [15]	5.0 [6.0]	Yes
	Cloxacillin	Porcine	76	149 [9.1]	148 [3.0]	Yes

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