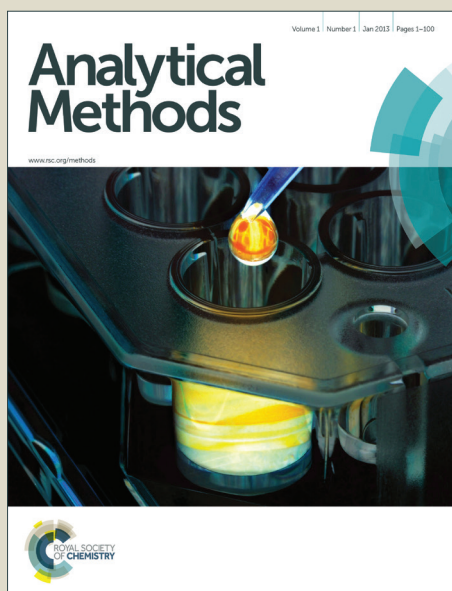


Analytical Methods

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**Simultaneous analysis of kasugamycin and streptomycin in vegetables
by liquid chromatography-tandem mass spectrometry**

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Keywords: mixed-mode chromatography, liquid chromatography, mass spectrometry,
kasugamycin, streptomycin.

25 **Abstract**

26 In this work an LC-MS/MS method has been developed for the simultaneous
27 analysis of kasugamycin (KAS) and streptomycin (STR) in vegetable samples. The use
28 of a mixed-mode HPLC column and a ternary mobile phase acetonitrile:water:0.5%
29 aqueous formic acid in gradient elution mode are proposed providing enough retention
30 and resolution for these highly polar compounds. Heated-electrospray (H-ESI) has been
31 used to ionize KAS and STR when coupling liquid chromatography to mass
32 spectrometry. In contrast to what happens with most aminoglycosides, KAS only
33 showed single charged ions in the full-scan mass spectrum. The $[M+H]^+$ of both KAS
34 and STR are proposed as precursor ions for tandem mass spectrometry since more
35 stable product ions and better ion ratio precisions were obtained. Fragmentation of KAS
36 has been studied, showing that the cleavage of the glycosidic bonds provided the main
37 product ions and the most abundant and selective ones are proposed for quantitation and
38 confirmation purposes in MRM mode. The LC-MS/MS method developed has been
39 applied to the analysis of vegetables. A simple clean-up procedure using hydrophilic-
40 lipophilic-balanced cartridges was applied to several vegetable extracts from tomato,
41 zucchini, chard and lettuce, obtaining recoveries >80% for both compounds. The high
42 matrix suppression observed for KAS in all matrices was reduced by a 1:5 sample
43 dilution with acetonitrile, providing a 100-fold improvement in sensitivity for this
44 compound. Method quality parameters have been established for tomato matrices,
45 obtaining method limits of quantitation in the low $\mu\text{g kg}^{-1}$ level ($5\text{-}10 \mu\text{g kg}^{-1}$) for both
46 compounds, precisions expressed in %RSD better than 7% and accuracies expressed as
47 relative error better than 8% at $100 \mu\text{g kg}^{-1}$ level, making this method suitable for
48 routine analysis.

1. Introduction

Kasugamycin (KAS) and streptomycin (STR) are two aminoglycoside (AGs) antibiotics active against certain types of gram-negative bacteria¹. Their general structure includes several aminosugars linked by glycosidic bonds to a streptidine (STR) or a 2-deoxystreptamine (KAS) unit. STR, produced by *Streptomyces griseus*, was the first antibiotic applied against plant diseases² to control fire blight of apple and pear, wild fire of tobacco and bacterial leaf blight of the rice plant, among others and its use is being reassessed due to the spread of resistant bacterial strains.³ KAS is produced by *Streptomyces kasugaensis*, it has a strong preventive effect against rice blast caused by *Piricularia oryzae*, and it has been also used for the treatment of bacterial diseases in bell peppers, tomatoes, citrus, cucumbers and many other fruiting crops. Moreover, the use of KAS as a substitute of STR for the treatment and prevention of fire blight in pears and apples has been suggested⁴ due to the low toxicity of KAS to crops and mammals and its activity against resistant strains. Thus, residues of these aminoglycosides might be present in fruits and vegetables.

Since public concern over pesticide residues in foods has become an important issue, recommended maximum residue levels (MRL) for these compounds in agricultural products have been established. Nevertheless, different levels have been set depending on the food commodity and the country. For instance, in the United States MRLs for KAS have been set at 0.04 mg kg⁻¹ in fruiting crops and at 0.05 mg kg⁻¹ for apples⁵ whereas for STR the MRLs range from 0.25 to 0.5 mg kg⁻¹, depending on the matrix.⁶ In the European Community their use is not authorised since 2005⁷ and thus a default MRL of 0.01 mg kg⁻¹ is recommended.⁸

Liquid chromatography coupled to mass spectrometry is the analytical technique of choice for the analysis of KAS and STR, which are highly hydrophilic compounds that lack chromophore groups. Due to the amino groups present in their structures, both STR and KAS are poorly retained in reversed-phase columns such as C18⁹⁻¹¹ and for this reason, alternative chromatographic separation modes such as ion pair liquid chromatography (IPLC), hydrophilic interaction liquid chromatography (HILIC) and mixed-mode chromatography have been used. The analysis of KAS presents a high analytical challenge and this compound is not currently included in the analysis of other AGs. In fact, only two studies have been published dealing with the determination of KAS in foods. The first one proposes a method for the analysis of both KAS and STR,

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3 89 and other AGs in animal tissues, milk and eggs using IPLC with trifluoroacetic acid¹²
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5 90 and in the second one, KAS is determined in soils and chili peppers¹³ using HILIC. This
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7 91 last study is the only one published regarding the analysis of KAS in vegetables.
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9 92 Moreover, most of the developed methods for STR have been established for its
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11 93 determination in foods from animal origin. Some of them use IPLC with ion pair
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13 94 reagents like heptafluorobutyric acid,¹⁴⁻²⁰ perfluoropentanoic acid,²¹ and trifluoroacetic
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15 95 acid,¹² while others use HILIC, often with high buffer concentrations in the mobile
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17 96 phase, such as ammonium acetate or formate at concentrations between 150 and 200
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19 97 mM, and with percentages of formic acid ranging from 0.1 to 1%.²²⁻²⁶ Additionally,
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21 98 mixed-mode chromatography with low ionic strength mobile phases has also been
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23 99 proposed for the analysis of STR and other AGs in meat.²⁷ However, only two methods
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25 100 dealing with the analysis of STR in fruits and vegetables can be found in the literature.
26
27 101 One of these studies proposes the use of IPLC with heptafluorobutyric acid as ion pair
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29 102 reagent for the determination of STR and tetracycline in pomegranate²⁸ and the other
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31 103 one applies HILIC to determine STR in apples.²⁹

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33 104 All these methods use electrospray (ESI) in positive mode as the ionization
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35 105 technique because protonation is favored by the presence of amine groups in their
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37 106 chemical structure. Low resolution analyzers such as triple quadrupole^{12-17,19-21,24-27,29}
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39 107 and quadrupole-ion trap^{22, 28} combined with multiple reaction monitoring (MRM)
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41 108 acquisition have been used to determine KAS and STR at low concentration levels. The
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43 109 fragmentation pathways of STR has been studied,^{30, 31} but to the best of our knowledge,
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45 110 the collision-induced dissociation (CID) mass fragmentation and the product ions
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47 111 assignment for KAS have not been proposed. Moreover, when coupling HILIC and
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49 112 IPLC methods to mass spectrometry, the presence of ion pair reagents and the high
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51 113 concentration of salts needed for ion exchange chromatography or HILIC could affect
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53 114 the performance of the mass spectrometric systems in both sensitivity and maintenance.
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55 115 So there is a need for LC-MS methods to analyze aminoglycosides that avoid the use of
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57 116 ion pair reagents or high concentration buffers in the mobile phase.

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59 117 The aim of this work is to develop a liquid chromatography-tandem mass
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118 spectrometry method for the simultaneous analysis of KAS and STR, which are the only
119 aminoglycoside antibiotics used as pesticides, in vegetables. For this purpose, the use of
120 a mixed-mode chromatographic column that combines different retention mechanisms
121 is evaluated in order to retain and separate both compounds without using ion pair
122 reagents or high buffer concentrations as required for HILIC. The ionization of these

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3 123 compounds under electrospray conditions is studied and their tandem mass
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5 124 spectrometry fragmentation is discussed, especially for KAS which has not been
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7 125 previously reported. Finally, the LC-MS/MS method proposed is applied to the
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9 126 simultaneous determination of KAS and STR in vegetables at low $\mu\text{g kg}^{-1}$ level.
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12 128 **2. Experimental**

13 129 14 130 **2.1. Reagents and materials**

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16 132 Streptomycin sulfate (STR, 98%, CAS No. 3810-74-0), and kasugamycin
17 133 hydrochloride hydrate (KAS, 79%, CAS No. 19408-46-9) were purchased from Dr.
18 134 Ehrenstorfer GmbH (Augsburg, Germany). Figure 1 shows the structures of the studied
19 135 compounds. The glycosidic rings of the structures have been labeled with capital letters
20 136 to help with the discussion of the fragmentation patterns. Ethylenediaminetetraacetic
21 137 acid (EDTA) disodium salt dihydrate (98.5%) was purchased from Sigma-Aldrich
22 138 (Steinheim, Germany). LC/MS grade methanol, acetonitrile (ACN) and water and
23 139 trichloroacetic acid (TCA) (99.5%) were obtained from Fluka (Steinheim, Germany)
24 140 and formic acid (98-100%) was provided by Merck (Darmstadt, Germany). Stock
25 141 standard solutions of KAS and STR (2.0 g kg^{-1}) were individually prepared by weight
26 142 in water and stored at -20°C . Intermediate solutions were prepared monthly from stock
27 143 standard solution by appropriate dilution. As aminoglycosides present high sorption
28 144 affinity to polar surfaces all contacts with glass were avoided and only polypropylene
29 145 laboratory material was used. All samples and mobile phases were filtered through a
30 146 $0.22 \mu\text{m}$ nylon membrane filters purchased from Whatman (Clifton, NJ, USA) to avoid
31 147 clogging. Nitrogen (99.8% pure) supplied by gas line from Air Liquide (Madrid, Spain)
32 148 was used for the API source. The collision-induced dissociation gas used for tandem
33 149 mass spectrometry experiments was high-purity Argon (ALPHAGAZ 1 Ar) also from
34 150 Air Liquide (Madrid, Spain). Sample clean-up was performed using Oasis HLB
35 151 cartridges (150 mg , 6 cm^3) and extraction manifold from Waters (Milford, MA, USA).
36 152 For the control of the pH of buffered solvents a pHmeter Basic 20 from Crison
37 153 Instruments (Alella, Spain) was used.
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154 155 **2.2. Sample treatment**

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3 157 Samples of tomato, chard, lettuce, zucchini and red pepper bought at local
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5 158 supermarkets in Barcelona (Spain), were cut, ground and homogenized with an
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7 159 Ultraturrax T25 basic (IKA-Werke, Staufen, Germany) and kept frozen at -20°C until
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9 160 analysis. Subsamples of 2 g were weighed in a 15 mL polypropylene centrifuge tube
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11 161 (Serviquimia, Barcelona, Spain) and extracted in a Sonorex RK100 ultrasonic bath
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13 162 (Bandelin Electronic GmbH & Co., Berlin, Germany) for 10 min with 2 mL of
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15 163 acetonitrile:aqueous 5% TCA, 2 mM EDTA (v/v 1:1) and then centrifuged at 4,000 rpm
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17 164 (Selecta-Macrotronic, J.P. SELECTA S.A., Abrera, Spain) for 10 min. The supernatant
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19 165 was then loaded onto an Oasis HLB SPE cartridge previously conditioned with 3 mL of
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21 166 methanol and 3 mL of water for clean-up. After discarding the first 0.75 mL of the
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23 167 eluate the rest was collected and filtered through a 0.22 µm nylon membrane. Finally,
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25 168 200 µL of the filtered extract were diluted with 800 µL of acetonitrile and injected into
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27 169 the LC-MS/MS system. Matrix-matched calibration standards were prepared with blank
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29 170 matrix samples spiked at 6 concentration levels ranging from 10 to 500 µg kg⁻¹ and
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31 171 submitting them to the same sample treatment described above.
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34 173 **2.3. HPLC-MS/MS**

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36 175 HPLC separation was performed on an Open Accela liquid chromatography
37 176 system (Thermo Fisher Scientific, San José, CA, USA), equipped with a quaternary
38 177 UHPLC pump, and a CTC PAL autosampler (CTC Analytics, Zwingen, Switzerland).
39 178 An Obelisc R (150 x 2.1 mm, 5 µm) column (SIELC Technologies, Prospect Heights,
40 179 IL, USA) was used for the proposed method. Mobile phases consisted of a mixture of
41 180 acetonitrile (A), aqueous 0.5% formic acid (B) and water (C). The chromatographic
42 181 initial conditions were 85% A, 3.8% B and 11.2% C, and a linear ramp was performed
43 182 in 1.5 min to 85% A, 7.5% B and 7.5% C. A second linear ramp was carried out in 4.5
44 183 min to 5% A, 50% B and 45% C and these conditions were maintained for 4 min,
45 184 moving back to initial conditions in 2 min and equilibrating the column for 8 min. The
46 185 flow rate was 300 µL min⁻¹ and the column was kept at ambient temperature. The
47 186 injection mode was full loop with a 5 µL loop.

48 187 Mass spectrometry was performed in a TSQ Quantum Ultra AM (Thermo Fisher
49 188 Scientific, San José, CA) triple quadrupole mass spectrometer equipped with heated-
50 189 electrospray (H-ESI) as ionization source. Electrospray voltage was set at 3.0 kV and
51 190 the temperatures of the ion transfer tube and the vaporizer were both held at 300°C.

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3 191 Sheath gas, auxiliary gas and ion sweep gas flow rate were 35, 10 and 0 a.u. (arbitrary
4 units), respectively. In tandem mass spectrometry experiments, both quadrupoles
5 192 worked in low resolution mode (0.7 full width half maximum, FWHM), the collision
6 193 gas pressure was 1.5 mTorr and the collision energies (CE) ranged from 11 eV to 18
7 194 eV. Two transitions were monitored for each compound using a dwell time of 250 ms (1
8 195 scan/cycle time). Transitions for quantitation and confirmation purposes and the
9 196 corresponding collision energies and ion ratios are listed in Table 1. Other instrument
10 197 settings were optimized by infusion of standard solutions (10 mg L⁻¹) at a flow rate of 5
11 198 $\mu\text{L min}^{-1}$, using the built-in syringe pump and a zero dead volume T-piece to mix them
12 199 with the mobile phase.
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22 202 **3. Results and discussion**

26 204 **3.1. HPLC-MS/MS**

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28 206 Due to the high polarity of KAS and STR their analysis by LC is usually
29 207 performed by IPLC or HILIC. However, the use of these chromatographic methods
30 208 requires the addition of ion pair reagents or buffers at high concentration to the mobile
31 209 phase, thus making difficult the compatibility with mass spectrometry. In this work a
32 210 mixed-mode Obelisc R column has been used to explore the simultaneous
33 211 chromatographic separation of KAS and STR, as this column has been proposed for the
34 212 separation of STR and other AGs using gradient elution with a mobile phase composed
35 213 of a mixture of acetonitrile:1% aqueous formic acid:water.²⁷ These conditions worked
36 214 well for STR but KAS eluted too early in the chromatogram due to the high elution
37 215 strength of the mobile phase, so alternative chromatographic conditions were optimized.
38 216 To this end, different aqueous formic acid gradients were tested. When the initial formic
39 217 acid concentration was lowered, the retention of KAS increased because at low pHs
40 218 hydrophilic interaction is the prevailing retention mechanism, enhancing the retention
41 219 with the decrease of the ionic strength. However, the retention and peak width of STR
42 220 also increased for the same reason, resulting in poor chromatographic efficiency (peak
43 221 width > 2 min) and long analysis times. To achieve good retention and peak efficiency,
44 222 a ternary gradient simultaneously decreasing acetonitrile and increasing the formic acid
45 223 concentration was used, providing the optimal conditions. The optimized gradient has
46 224 been indicated in the experimental section and Figure 2 shows the chromatogram

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3 225 obtained under those chromatographic conditions. As can be seen, the compounds were
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5 226 baseline separated in less than 6 minutes with good peak shape and far enough from the
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7 227 elution front.

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9 228 Regarding mass spectrometry, thermally assisted electrospray (H-ESI) was used
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11 229 as ionization source in LC/MS, as it improved desolvation, especially with mobile
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13 230 phases with a high percentage of water. Full-scan mass spectra were studied to select
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15 231 the most adequate precursor ions for tandem mass spectrometry. The mass spectrum of
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17 232 KAS (Figure 2) is very simple showing only the protonated molecule $[M+H]^+$ (m/z 380)
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19 233 and the loss of CO_2 from the protonated molecule due to in-source collision-induced
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21 234 dissociation (m/z 336). In contrast, the mass spectrum of STR (Figure 2) is more
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23 235 complex, because STR has two guanidine groups with high pKa values (13.40 ± 0.70)³²
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25 236 that favored the generation of multiple charged ions. So, in the mass spectrum of STR
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27 237 the single and double charged ions, $[M+H]^+$ and $[M+2H]^{2+}$ (m/z 582 and 292,
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29 238 respectively) due to protonation and also the single and double charged ions due to the
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31 239 hydrated form of the aldehyde group, $[M+H_2O+H]^+$ and $[M+H_2O+2H]^{2+}$ (m/z 600 and
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33 240 301 respectively), can be observed. Moreover, some in-source fragmentation from the
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35 241 cleavage of the glycosidic bonds is also present (Figure 2). The $[M+H]^+$ was selected as
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37 242 precursor ion for tandem mass spectrometry since it showed the best fragmentation
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39 243 performance, leading to more stable product ions and better precisions than the double
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41 244 charged ions. As regards the hydrated form of the single charged ion, it showed the
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43 245 unspecific loss of water as the most intense product ion, which is more susceptible to
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45 246 interferences than the fragmentation from the protonated molecule. The chosen SRM
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47 247 transitions for STR are listed and assigned in Table 1. For KAS, the $[M+H]^+$ was also
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49 248 selected as precursor ion and the triple quadrupole product ion scan spectrum is shown
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51 249 in Figure 3. The fragmentation pattern of KAS has been discussed using the
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53 250 nomenclature defined by Goolsby and Brodbelt,³⁰ which proposes the use of the
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55 251 subscript “1” for the cleavage of the bond between the ring designated with the
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57 252 uppercase letter and the glycosidic oxygen, and the subscript “2” for that between the
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59 253 glycosidic oxygen and the next ring. As happens with most AGs, the main product ions
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254 of the $[M+H]^+$ of KAS involved the cleavage of the glycosidic bond (Figure 3), whether
255 arising from the loss of the glucose ring ($[M+H - B_2]^+$, m/z 200) or from the combined
256 loss of the glucose ring and the amino(imino)acetic side chain ($[M+H - B_2 -$
257 $C_2H_4O_2N_2]^+$, m/z 112). The presence of the carboxylic group in the side chain favored
258 also the loss of CO_2 from the protonated molecule (m/z 336) or from the $[M+H - B_2]^+$

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3 259 (m/z 156). Additionally, further cross-ring cleavage and rearrangement of the m/z 112
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5 260 would produce the product ion at m/z 70. Fragmentation of KAS is shown in Figure 3
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7 261 and the elemental compositions of all product ions have been confirmed by MS/HRMS
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9 262 experiments in a Q-Exactive (Thermo Fisher Scientific) at a mass resolution of 35,000
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11 263 FWHM (full width half maximum) at m/z 200, obtaining mass errors below 2 ppm. The
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13 264 most abundant product ions, at m/z 112 and m/z 200, were selected for quantitation and
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15 265 confirmation, respectively, of KAS when working in MRM mode (Table 1).

16 266 Instrumental quality parameters including limits of detection (LOD), linearity
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18 267 and run-to-run and day-to-day precisions were calculated (Table 2). LODs, based on a
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20 268 signal-to-noise ratio of 3:1 for the confirmation transition, were estimated by injecting
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22 269 standard solutions at concentrations down to $0.1 \mu\text{g kg}^{-1}$, obtaining results lower than 10
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24 270 pg injected for both compounds. KAS provided slightly better instrumental sensitivity,
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26 271 probably because for this compound only one precursor ion was obtained in ESI in
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28 272 contrast with what happened to STR, where the total ion signal was split into different
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30 273 species. Standard calibration curves with 6 concentration levels between 5 and $300 \mu\text{g}$
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32 274 kg^{-1} showed good linearity ($R^2 > 0.996$). Run-to run ($n=5$) and day-to-day ($n=5$, 3 days)
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34 275 precisions were calculated at two concentration levels, $15 \mu\text{g kg}^{-1}$ and $150 \mu\text{g kg}^{-1}$, and
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36 276 were always lower than 7% for both compounds. Retention time precision was also
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38 277 evaluated in the same way as the concentration precision, and low values, even for the
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40 278 day-to-day precision (below 1%), were obtained. Regarding the ion ratio (relative areas
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42 279 for quantifier vs qualifier transitions) precisions, the %RSD values were below 10% for
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44 280 both compounds.

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282 3.2. Sample treatment

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284 To analyse KAS and STR in vegetables, a sample treatment previously proposed
285 for the analysis of STR and other AGs in meat²⁷ was evaluated, consisting of an
286 extraction with an aqueous solvent containing a strong acid (TCA) and a chelating agent
287 (EDTA) followed by a clean-up with hydrophilic-lipophilic-balanced cartridges (HLB).
288 The clean-up with HLB cartridges served to retain the matrix coextractives, that gave a
289 cloudy and slightly coloured aspect to the extract. The highly polar AGs were not
290 retained in this sorbent and eluted directly. Recoveries were evaluated by means of the
291 analysis of blank tomato samples (2 g) spiked at $250 \mu\text{g kg}^{-1}$, and good recoveries
292 ($>80\%$) were obtained for STR, which were similar to those obtained in the previous

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3 293 work. However, KAS, which had not been previously studied, was not recovered at all.
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5 294 Recoveries with solvent based standards were close to 100%, and when blank tomato
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7 295 extracts spiked after sample treatment were analysed poor results for KAS were also
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9 296 obtained. Thus, the lack of signal for KAS was due to high matrix suppression for this
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11 297 compound and not to poor recoveries. Variations of the sample treatment method such
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13 298 as changes in the extraction solvent, or the use of other SPE strategies like ion-exchange
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15 299 cartridges not only did not provide successful results for KAS but also worsened the
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17 300 recoveries for STR as well, so alternative method were sought. As a first step, the use of
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19 301 an ionization source less susceptible to matrix effects such as Atmospheric Pressure
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21 302 Chemical Ionization (APCI) was tested. Although the high polarity and low volatility of
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23 303 AGs would produce lower ionization efficiency than ESI, the decrease in the ion
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25 304 suppression might produce an overall improvement in sensitivity. In fact, the use of
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27 305 APCI has been previously proposed to overcome ion suppression due to ion pair
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29 306 reagents for the LC-MS/MS analysis of gentamicin impurities.³³ As expected, STR
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31 307 provided lower signal intensities in APCI than in ESI, but no signal was observed for
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33 308 KAS in APCI in both, positive and negative modes. The lack of response of KAS might
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35 309 be explained by the presence of a carboxylic group in a side chain in KAS (Figure 1)
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37 310 which might favor the formation of an internal hydrogen bond with the primary amino
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39 311 substituent in ring A that would make KAS difficult to ionize in APCI in both positive
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41 312 and negative modes. Anyhow, as the ionization of KAS was not possible, the use of the
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43 313 APCI source was discarded.

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45 314 As a second step, the sample extract was diluted with ACN. Blank vegetable
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47 315 samples (lettuce, zucchini, red pepper, chard and tomato) spiked at 500 $\mu\text{g kg}^{-1}$ were
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49 316 submitted to the sample treatment and the obtained extracts were analysed without
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51 317 dilution and diluting 1:1, 1:5, and 1:10 with acetonitrile. For all the samples, an
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53 318 important increase on the response of KAS was obtained when diluting with
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55 319 acetonitrile, which allowed its detection. As an example, Figure 4 shows the LC-
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57 320 MS/MS chromatograms of two aliquots of a blank tomato extract spiked with KAS and
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59 321 STR at 500 $\mu\text{g kg}^{-1}$ without dilution (A) and diluted 1:5 with ACN (B). As it can be
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322 seen, KAS response increased considerably after a 1:5 dilution, showing up to a 100-
323 fold enhancement in signal intensity, which provided a significant decrease of the LODs
324 allowing the detection of KAS at the European MRL value (0.01 mg kg^{-1}) which is the
325 most restrictive one. Further dilution of the extracts did not provide an improvement in
326 signal intensity, probably because the decrease in signal due to dilution was higher than

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3 327 the improvement due to the reduction of matrix suppression. As a result, a 1:5 dilution
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5 328 of the extract is proposed for the analysis of KAS and STR in vegetable samples. As can
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7 329 be seen in Figure 4, signal intensity of STR significantly decreased when diluting the
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9 330 sample, as apparently a less matrix effect was affecting this compound. Nevertheless,
10 331 the response of STR after the dilution is similar to that of KAS.
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14 333 **3.3. Method performance**

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17 335 Method quality parameters such as method limit of quantitation (MLOQ), run-
18 336 to-run and day-to-day precisions, accuracy and ion ratio precision were evaluated using
19 337 tomato samples free of the analytes as blank samples, and the obtained values are listed
20 338 in Table 2. As KAS was affected from ion suppression, matrix-matched calibration was
21 339 used for the quantitation of the samples. Matrix-matched calibration curves were
22 340 prepared in the 10-500 $\mu\text{g kg}^{-1}$ range by spiking blank tomato samples and submitting
23 341 them to the sample treatment detailed in the experimental section. The MLOQs (signal-
24 342 to-noise ratio of 10:1) were estimated by injecting samples spiked at low concentration
25 343 values (below 10 $\mu\text{g kg}^{-1}$). As can be seen in Table 2, in contrast to what happened with
26 344 the instrumental LOD, the MLOQ obtained for STR was better than that of KAS,
27 345 probably as a result of the higher matrix suppression that affected the latter. MLOQs
28 346 were at or below the default MRL values established for these compounds in the
29 347 European Union (10 $\mu\text{g kg}^{-1}$), and far below the MRL allowed in other countries such as
30 348 USA (250 $\mu\text{g kg}^{-1}$ for STR),⁶ Japan (300 $\mu\text{g kg}^{-1}$ for KAS)³⁴ or Canada (100 $\mu\text{g kg}^{-1}$ for
31 349 both compounds).^{35,36} MLOQs in the other vegetable matrices also were studied, and in
32 350 all cases the results were below 15 $\mu\text{g kg}^{-1}$ for both compounds, only being above the
33 351 MRL in the European Union in red pepper matrices. To evaluate run-to-run precision,
34 352 five replicate analyses of blank samples spiked at two concentration levels, 10 $\mu\text{g kg}^{-1}$
35 353 and 100 $\mu\text{g kg}^{-1}$, were analysed. Day-to-day precision was calculated by analysing 5
36 354 replicates each day during three days at the same two concentration levels. %RSD
37 355 values based on concentrations (Table 2) ranged from 13 to 24% for the low
38 356 concentration level, and were below 8% for the high concentration level. Accuracy
39 357 expressed as relative error was also evaluated at the two concentration levels and as can
40 358 be seen in Table 2, KAS provided worse accuracies than STR for the low level (23%
41 359 versus 15%), as it corresponded to the MLOQ for this compound (10 $\mu\text{g kg}^{-1}$). Ion ratio
42 360 relative errors with respect to those of standards were lower than 10% and ion ratio

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3 361 precisions were consistent with those obtained with standard solutions (below 13%). In
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5 362 comparison with previous literature works that analyze either KAS or STR in
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7 363 vegetables,^{13,28,29} the proposed method allows the simultaneous analysis of both
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9 364 compounds without losing sensitivity. Also, ion pair reagents are not needed for the
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11 365 chromatographic separation, that favours MS performance and maintenance, and
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13 366 provides slightly better LODs than those reported in the analysis of KAS and STR in
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15 367 animal tissues.¹²
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17 369 **3.4. Method application**

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21 371 Over 24 vegetable samples, including tomato (2 samples), lettuce (2 samples),
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23 372 chard (2 samples), red pepper (2 samples), zucchini (2 samples) and pear (4 samples),
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25 373 purchased at local supermarkets were analyzed. No residues of KAS or STR were
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27 374 detected at levels above the MLOQ in any of the analyzed samples. This fact shows the
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29 375 compliance of the European Community regulations concerning the use of pesticides
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31 376 and the control of the presence of residues of pesticides in foods. Negative results were
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33 377 also found in the analysis of 10 tomato samples acquired from a vegetable garden that
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35 378 had been treated with both compounds. Although it was expected that some residues
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37 379 should have been found in this samples, the short extinction time of these compounds,
38
39 380 which is lower than 6 days for KAS in chili peppers,¹³ might be a possible reason for
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41 381 the negative results. However, the method provided satisfactory results with spiked
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43 382 blank samples at levels below the MRLs established in most countries (Table 2),
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45 383 making it suitable for routine analysis in laboratories dealing with the control in
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47 384 countries where the use of this compounds as pesticides is allowed (e.g. USA and
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49 385 Canada) and also in imported and exported goods.
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51 387 **4. Conclusions**

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55 389 In this work a method is proposed for the simultaneous analysis of KAS and
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57 390 STR in vegetable samples. A base line chromatographic separation, good peak shapes
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59 391 and short analysis time (less than 6 minutes) are achieved using a mixed-mode column
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392 and an ternary mobile phase composed of acetonitrile:water:0.5% aqueous formic acid
393 in reversed gradient elution. Single charged ions are proposed as precursors for tandem
394 mass spectrometry since they showed better fragmentation performance than the

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3 395 multiple charged ions. The CID MS/MS fragmentation of KAS was studied for first
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5 396 time concluding that the main product ions arose from the cleavage of the glycosidic
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7 397 bonds. The simultaneous loss of glucose and the amino(imino)acetic acid side chain
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9 398 (m/z 112) was used for quantitation, whereas the loss of glucose (m/z 200) was chosen
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11 399 for confirmation purposes. The high matrix suppression that affected the electrospray
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13 400 KAS response when analyzing vegetables can be greatly decreased by diluting the
14
15 401 obtained extract 1:5 with ACN, achieving a 100-fold improvement in signal intensity.
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17 402 Under the optimal working conditions the LC-MS/MS developed method provided
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19 403 MLODs lower enough to determine KAS and STR at the lower MRL established for
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21 404 them in all countries ($10 \mu\text{g kg}^{-1}$). The applicability of the method has been assessed by
22
23 405 analyzing spiked samples, and good method quality parameters have been obtained.
24

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35

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Figure captions

Figure 1. Structures and acronyms of the studied compounds

Figure 2. Chromatogram of a standard mixture (100 ng g⁻¹) using the conditions given in the experimental section and full-scan mass spectra of KAS and STR

Figure 3. Triple quadrupole product ion scan and fragmentation of KAS

Figure 4. LC-MS/MS chromatogram of a blank tomato extract spiked at 500 µg kg⁻¹ level A) without dilution and B) diluting 1:5 with acetonitrile

496 **Tables**

497 Table 1. Tandem mass spectrometry working conditions. Selected SRM transitions, collision energies (CE), ion assignments and ion ratios

Compound	Retention time (min)	Precursor ion		Product ion				Ion ratio
		<i>m/z</i>	Assignment	<i>m/z</i>	Assignment	CE (eV)		
KAS	4.44	380.2	[M+H] ⁺	112.1	[M+H-B ₂ -C ₂ H ₄ O ₂ N ₂] ⁺	18	Quantitation	1.3
				200.1	[M+H-B ₂] ⁺	11	Confirmation	
STR	5.07	582.3	[M+H] ⁺	263.1	[C ₂] ⁺	31	Quantitation	1.7
				246.1	[C ₂ -NH ₃] ⁺	35	Confirmation	

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509 Table 2. Instrumental and method quality parameters.

Compound	Instrumental quality parameters								
	LOD (pg)	Run-to-run precision (%RSD, n=5)				Day-to-day precision (%RSD, n=5, 3 days)			
		Concentration		Retention time precision	Ion ratio precision	Concentration		Retention time precision	Ion ratio precision
		5 µg kg ⁻¹	150 µg kg ⁻¹			5 µg kg ⁻¹	150 µg kg ⁻¹		
KAS	3	4.1	2.5	0.2	4.8	6.0	4.6	0.7	7.1
STR	8	3.6	1.5	0.3	6.7	6.7	3.2	0.4	8.9
	Method quality parameters (tomato matrix)								
	MLOQ (µg kg ⁻¹)	Precision (%RSD)				Accuracy		Ion ratio	
		Run-to-run (n=5)		Day-to-day (n=5, 3 days)		(%Rel. Error)		Rel. Error	Precision
		10 µg kg ⁻¹	100 µg kg ⁻¹	10 µg kg ⁻¹	100 µg kg ⁻¹	10 µg kg ⁻¹	100 µg kg ⁻¹	(%)	(%RSD)
KAS	10	13	5.2	24	6.7	23	8.0	10	13
STR	4	15	5.0	21	5.2	19	5.2	9.2	11

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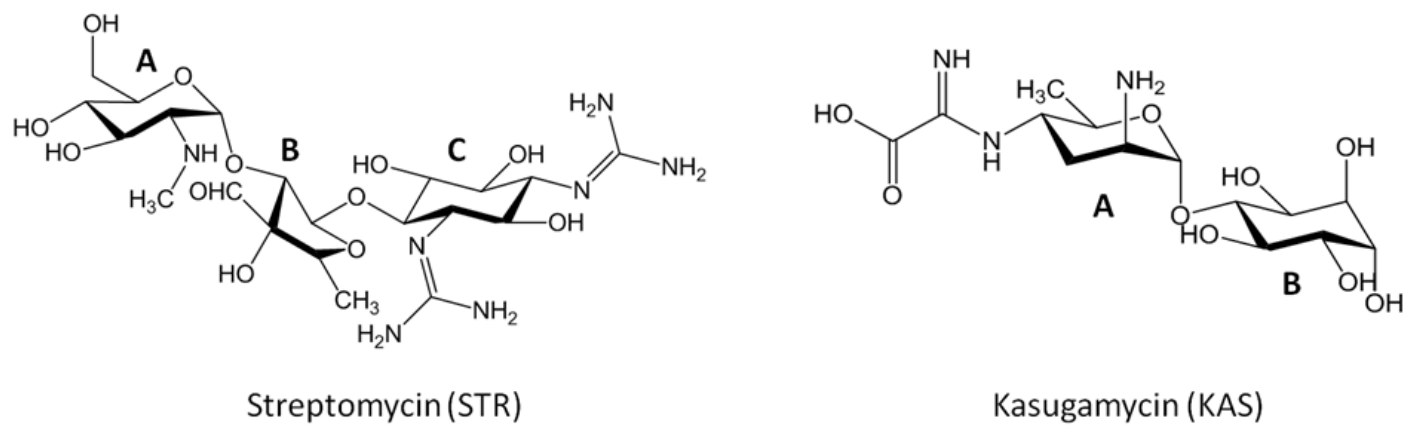
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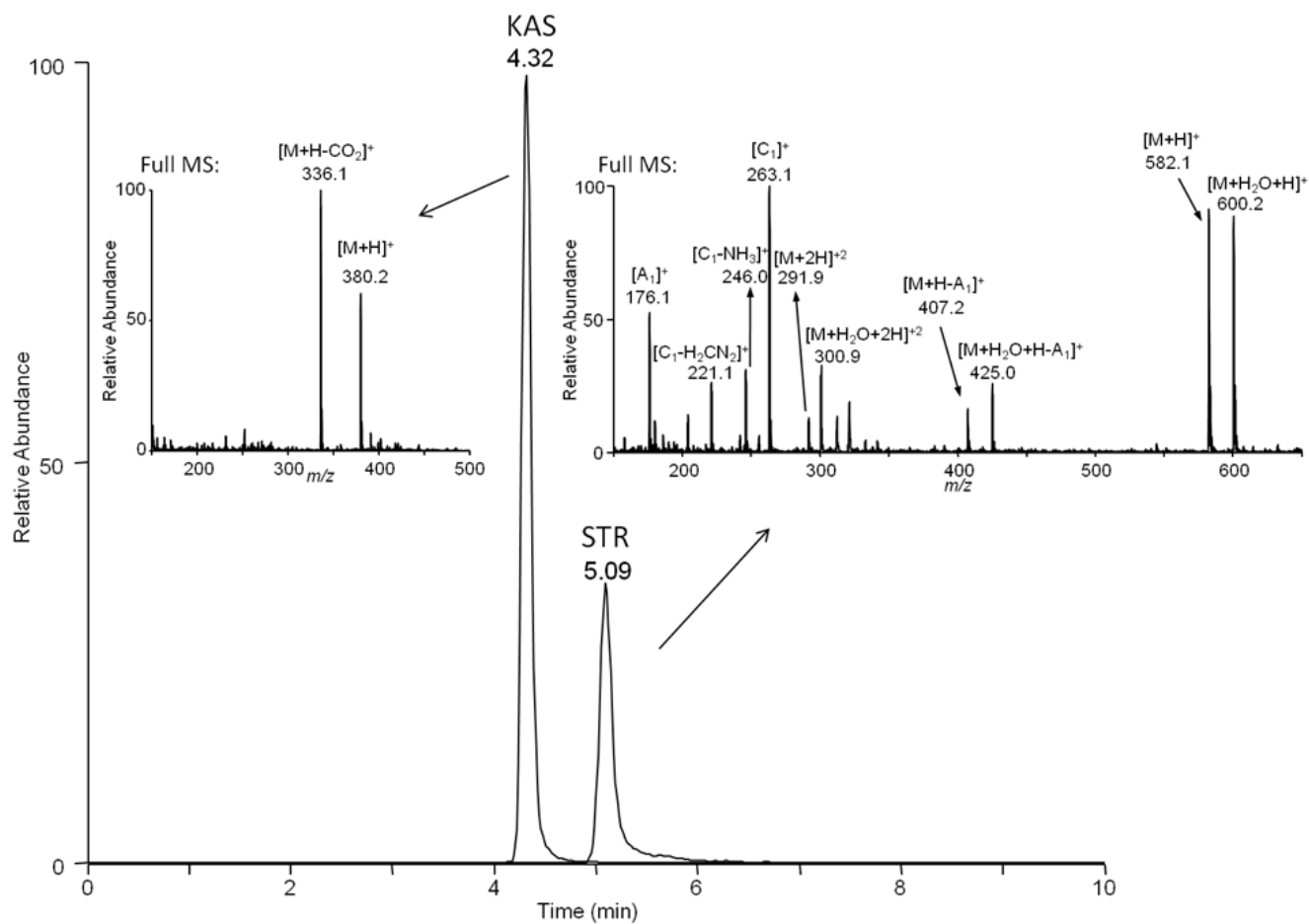
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518 Figure 1. Structures and acronyms of the studied compounds



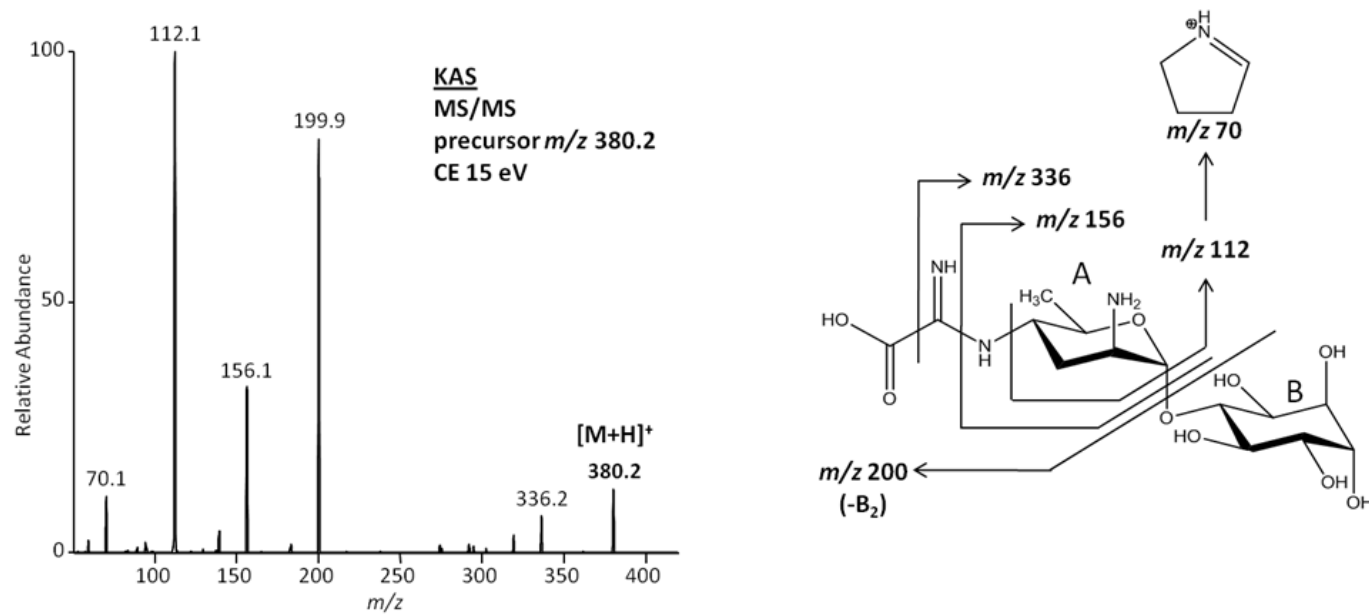
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520 Figure 2. Chromatogram of a standard mixture (100 ng g⁻¹) using the conditions given in the experimental section and full-scan mass spectra of
521 KAS and STR



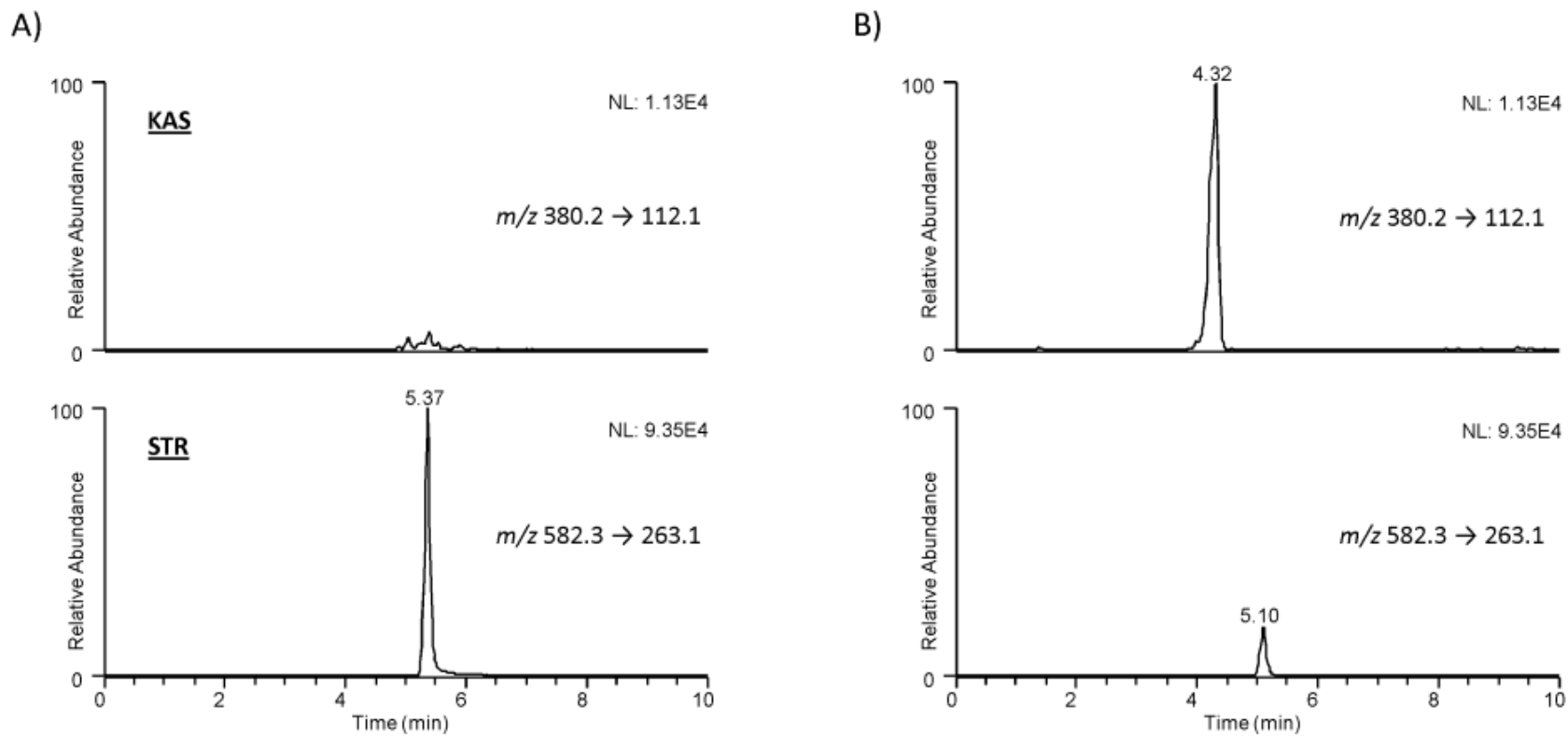
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523 Figure 3. Triple quadrupole product ion scan and fragmentation of KAS



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525 Figure 4. LC-MS/MS chromatogram of a blank tomato extract spiked at 500 $\mu\text{g kg}^{-1}$ level A) without dilution and B) diluting 1:5 with
526 acetonitrile



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