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# When other separation techniques fail – Compound-specific carbon isotope ratio analysis of sulfonamide containing pharmaceuticals by HT- LC-IRMS

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Compound-specific isotope analysis (CISA) of non-volatile analytes has been enabled by the introduction of the first commercial interface to hyphenate liquid chromatography with an isotope ratio mass spectrometer (LC-IRMS) in 2004, yet carbon isotope analysis of unpolar and moderately polar compounds is still a challenging task since only water as eluent and no organic modifiers can be used to drive the separation in LC. The only way to increase the elution strength of aqueous eluents in reversed phase LC is the application of high temperatures to mobile and stationary phase (HT-LC-IRMS). In this context we present the first method to determine carbon isotope ratios of pharmaceuticals that cannot be separated by already existing separation techniques for LC-IRMS, such as reversed phase chromatography at normal temperatures, ion-chromatography and mixed mode chromatography. The pharmaceutical group of sulfonamides, which is generally mixed with trimethoprim in

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3 pharmaceutical products, has been chosen as probe compounds. Substance amounts as low as  
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5 0.3  $\mu$ g are sufficient to perform a precise analysis. The successful applicability and  
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7 reproducibility of this method is shown by the analysis of real pharmaceutical samples. The  
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9 method provides the first tool to study the pharmaceutical authenticity as well as degradation  
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11 and mobility of such substances in the environment by using the stable isotopic signature of  
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13 these compounds.  
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## 20 **Introduction**

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23 Compound-specific isotope analysis of non-volatile organic compounds is often performed by  
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25 gas chromatography-isotope ratio mass spectrometry (GC-IRMS) with prior derivatization.<sup>1</sup>  
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27 Although these methods give accurate and reproducible results, they are not applicable for  
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29 many non-volatiles. The hyphenation of liquid chromatography (LC) to isotope ratio mass  
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31 spectrometry enables the determination of carbon isotope ratios of very polar, thermolabile  
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33 and high molecular weight compounds without the need for derivatization.<sup>2</sup> But due to the  
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35 design of the interface organic modifiers cannot be used in LC eluents.<sup>3</sup> In order to precisely  
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37 measure carbon isotope ratios, the analytes need to be detected as CO<sub>2</sub>. In the interface, the  
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39 whole column effluent is non-selectively converted to CO<sub>2</sub> by peroxodisulfate in a heated  
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41 capillary, extracted from the aqueous phase into a carrier gas stream via a membrane and  
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43 transported to the IRMS inlet. Consequently, any additional carbon compound (either as  
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45 buffer or solvent) in the eluent besides analyte carbon would result in a severe decrease of  
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47 sensitivity, wrong carbon isotope ratios or a saturation of the isotope ratio mass spectrometer  
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49 signal.. This restricts the type of applicable separation techniques for LC-IRMS. Several  
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51 methods have been reported for the determination of carbon isotope ratios of carbohydrates,  
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53 amino sugars and amino acids based on ion exchange chromatography.<sup>4-6</sup>  
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3 However, with ion exchange as sole separation mechanism, it is difficult to obtain a full  
4 separation of proteinogenic amino acids.<sup>2</sup> The resolution of many amino acids can be  
5 substantially improved by adding the possibility for hydrophobic interactions by a further RP-  
6 column.<sup>2</sup> Almost all amino acids can be resolved by mixed-mode chromatography, i.e. the  
7 deliberate combination of RP and ion exchange interaction sites in one stationary phase,  
8 which is nowadays the most published method for LC-IRMS analysis of underivatized amino  
9 acids.<sup>7</sup> The stationary phase contains ion exchange groups bound to alkyl chains or phenyl  
10 rings, which also allow an operation with 100 % aqueous eluents without the risk of a loss of  
11 retention due to particle pore dewetting.  
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24 There are no studies, which use ion chromatography or mixed mode chromatography for  
25 compound-specific isotope analysis of pharmaceuticals. Solely, carbon isotope ratios of  
26 Paracetamol and Aspirin<sup>TM</sup> have been measured to show the applicability of the LC-IRMS  
27 interface system. In this study a reversed phase (RP) silica based column at ambient  
28 temperature was used to separate those two compounds.<sup>3</sup> However, it is obvious, that also ion  
29 chromatography as well as mixed mode chromatography can be used for those  
30 pharmaceuticals. In the following, we will discuss, why reversed phase separation at ambient  
31 temperatures, HT-LC separation with porous graphitic carbon materials, ion chromatography  
32 and mixed-mode phases cannot be not feasible for the here investigated pharmaceutical  
33 compound classes of sulfonamides and trimethoprim.  
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47 Sulfonamides are amphoteric target compounds which are negatively charged at pH values  
48 greater than eight and positively at very low pH. In theory, both anion as well as cation  
49 exchange chromatography may work under these circumstances. However, a change of the  
50 pH to 8 would result in negatively charged sulfonamides but there is no anionic group at any  
51 pH for trimethoprim. To this end, a separation of the chosen target compounds would not be  
52 possible. Since trimethoprim is usually combined in pharmaceutical products containing  
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3 sulfonamides, valuable information on its carbon isotope ratio would be lost. Another  
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5 problem at pH 8 is an increased background caused by a much higher solubility of CO<sub>2</sub> from  
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7 air in the employed eluents, which requires a sophisticated eluent preparation and  
8  
9 conservation in order to maintain a low baseline in the chromatogram. Apart from the above  
10  
11 mentioned criteria against anion exchange, cation exchange is not a good option, since the pH  
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13 of the mobile phase should be two pH units below  $pK_a$  for good peak shape and so, high  
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15 concentration of acid would be necessary in the eluent. Despite these rather practical reasons  
16  
17 the retention of the target sulfonamides and trimethoprim on such columns will remain a  
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19 mixture of ion exchange and reversed phase chromatography due to hydrophobic interactions  
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21 with the polymeric support of the ion exchange resin.  
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26 Such an intentional combination of retention mechanisms is utilized in the mixed-mode  
27  
28 columns mentioned above. In case of amino acids it was shown that these stationary phases  
29  
30 work well because the elution strength of water is high enough to elute hydrophobic amino  
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32 acids from the mixed-mode phase without assistance of organic modifiers.<sup>7</sup> In contrast to  
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34 this, using comparable mixed-mode phases for sulfonamides and trimethoprim a much higher  
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36 retention than for a reversed phase separation is expected due electrostatic and strong  
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38 hydrophobic interactions. Thus, this method would require an organic proportion in the  
39  
40 mobile phase to aid elution, which is not compatible with LC-IRMS, or operate the  
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42 chromatography at higher temperatures. But the temperature stability of mixed-mode columns  
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44 remains to be examined.  
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49 An alternative approach that overcomes these described restrictions is high temperature liquid  
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51 chromatography (HT-LC) with reversed phases. HT-LC coupled to IRMS has already been  
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53 used to determine carbon isotope ratios of small organic acids<sup>8</sup> and caffeine derivatives.<sup>9</sup> The  
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55 elution strength of aqueous mobile phases without organic modifiers is often too low to elute  
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57 compounds from unpolar stationary phases at ambient temperatures. Especially, for the here  
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3 investigated compounds elution times increase dramatically, which results in inadequate  
4 retention times, a loss in resolution and signal sensitivity. The application of temperatures  
5 notably higher than ambient to mobile phase and column has been proposed to substitute  
6 organic solvents.<sup>10</sup> With increasing temperatures the static permittivity and viscosity of water  
7 decreases. At 150 °C (at a pressure of 50 bar) the static permittivity of water is nearly the  
8 same as that of about 70 % methanol or acetonitrile/water mixtures.<sup>10</sup> Thus, elution strength  
9 of pure aqueous mobile phases in RP chromatography at high temperatures is similar to  
10 solvent/water mixtures encountered in RP-LC. Temperature gradients instead of solvent  
11 gradients can be used to separate compounds on nonpolar stationary phases such as octadecyl  
12 silica or porous graphitized carbon (PGC). Temperature-programmed elution can even be  
13 predicted from two measurements with different temperature gradients.<sup>11</sup>  
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28 As the LC-IRMS detector is very sensitive to any carbon containing compound, a major  
29 concern using high temperature liquid chromatography (HT-LC) is the carbon feed from the  
30 column into the interface. At high temperature column bleed is much higher due to faster  
31 hydrolysis of the support material and bonded phase. Column bleed invisible for most  
32 common LC detection techniques may overlay the analyte peak or produce elevated  
33 background levels. Thus, Zhang et al.<sup>9</sup> evaluated four stationary phases which have proven  
34 their temperature stability in previous studies.<sup>12</sup> Hybrid particles from ethylene bridged silica  
35 and zirconium dioxide based materials can be used with excellent results for LC-IRMS at HT-  
36 LC conditions. Up to column-specific maximum temperatures column bleed did not affect  
37  $\delta^{13}\text{C}$ -values of various probe compounds.<sup>9</sup> It also has to be considered that carry over or late  
38 elution of hydrophobic compounds retained on very unpolar phases like PGC can alter  $\delta^{13}\text{C}$ -  
39 values of the analytes due to ghost peaks or occasional change in background  $^{13}\text{C}/^{12}\text{C}$ -ratio.  
40 Additionally,  $\delta^{13}\text{C}$ -values of thermolabile analytes may be influenced by isotopic  
41 fractionations resulting from thermal degradation on the column.<sup>13</sup> Porous graphitic carbon  
42 columns such as the Hypercarb™ possess very high retention at ambient temperatures.<sup>13</sup>  
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3 Although the retention times are decreased at higher temperatures, an unacceptable column  
4 bleed can be observed during temperature gradient elution.<sup>8</sup>  
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11 The aim of this work was to show that a HT-LC can be used for the determination of carbon  
12 isotope ratios of sulfamerazine, sulfadiazine, sulfamethoxazole, sulfathiazole and  
13 trimethoprim. These antibiotic substances have been widely used in veterinary and human  
14 medicine for some decades. Although they have been subject of many research campaigns in  
15 environmental<sup>14</sup> and pharmaceutical science<sup>15</sup> so far, but no determination of isotope ratios at  
16 natural abundance has been published. Several HT-LC methods for trimethoprim and  
17 sulfonamide measurements have been published in literature<sup>11, 16, 17</sup> for quantitative analysis,  
18 but to our best knowledge no carbon-free eluents have been used and no stable isotope  
19 signatures of these compounds have been determined. Up to now, isotope ratios of polar  
20 active pharmaceutical ingredients can only be determined either by <sup>13</sup>C-NMR or labor-  
21 intensive offline techniques like preparative LC followed by elemental analyzer (EA-IRMS).  
22 Here, we demonstrate a pure aqueous separation of these compounds for its use in CSIA and  
23 proved its potential for verifying the authenticity of antibiotics containing sulfamethoxazole  
24 and trimethoprim via  $\delta^{13}\text{C}$ -values.  
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## 43 **Experimental**

### 44 **Chemicals and Reagents**

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46 Analytical standards of sulfadiazine, sulfathiazole, sulfamerazine, sulfamethoxazole and  
47 trimethoprim were purchased from Sigma-Aldrich GmbH (Steinheim, Germany) in a purity of  
48 at least 98 %. Sulfamethoxazole and trimethoprim were purchased from Sigma-Aldrich  
49 GmbH (Steinheim, Germany) as well but in >99 % purity (VETRANAL<sup>TM</sup>). Structures and  
50  $pK_a$ -values of these compounds are listed in Table 1. Sodium peroxodisulfate ( $\text{Na}_2\text{S}_2\text{O}_8$ ),  
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3 phosphoric acid ( $\text{H}_3\text{PO}_4$ ) and sodium hydrogenphosphate ( $\text{NaH}_2\text{PO}_4$ ) for eluent and reagent  
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5 solutions were supplied by Fluka (Steinheim, Germany). Reagent solutions and eluents were  
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7 degassed in an ultrasonic bath (Bandein Eletronic, Berlin, Germany) under vacuum  
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9 (Vacuubrand, Wertheim, Germany). In order to avoid regassing these solutions were  
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11 continuously purged with helium of 99.999 % purity (Air Liquide, Oberhausen, Germany) at  
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13 a flow rate of approximately  $30 \text{ mL min}^{-1}$ .  
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### 16 17 **Analytical standard solutions**

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19 Standard solutions of sulfonamides were freshly prepared prior to each set of analyses by  
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21 dissolving a required amount of substance in 15 mL HT-LC buffer solution at approximately  
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23  $60 \text{ }^\circ\text{C}$ .  
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### 26 27 **LC conditions**

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29 HT-LC system consisted of a Rheos Allegro binary pump (Flux Instruments, Buchs,  
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31 Switzerland) and a HT-HPLC 200 column oven (SIM Scientific Instruments Manufacturer  
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33 GmbH, Oberhausen, Germany). The compounds were separated on an XBridge  $\text{C}_{18}$  column  
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35  $100 \times 2.1 \text{ mm}$ ,  $3.5 \text{ }\mu\text{m}$  particle size equipped with a  $10 \times 2.1 \text{ mm}$  pre-column packed with the  
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37 same material (Waters, Eschborn, Germany). A sodium phosphate buffer of  $5 \text{ mM}$  and  $\text{pH } 3$   
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39 was used as the eluent at a flow rate of  $500 \text{ }\mu\text{L min}^{-1}$ . The temperature gradient was  $60 \text{ }^\circ\text{C}$  for  
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41  $3 \text{ min}$ , then to  $80 \text{ }^\circ\text{C}$  at  $3 \text{ }^\circ\text{C min}^{-1}$ , held for  $15 \text{ min}$ , then to  $100 \text{ }^\circ\text{C}$  at  $3 \text{ }^\circ\text{C min}^{-1}$  and held for  $5$   
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43  $\text{min}$ .  
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48 Flushing of the organic solvent from shipping or storage of the column prior to measurements  
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50 was done while the column was installed in the HT-LC-IRMS system. In order to prevent  
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52 precipitation the column was first flushed for  $2 \text{ hours}$  with pure water and afterwards  
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54 equilibrated with the phosphate buffer for  $1 \text{ hour}$  at a flow rate of  $500 \text{ }\mu\text{L min}^{-1}$ .  
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### 58 59 **LC-IRMS interface conditions**



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3 As interface between HPLC and IRMS an LC-IsoLink (Thermo Scientific, Bremen,  
4 Germany) was used. The wet chemical oxidation of the compounds was realized by online  
5 mixing of HPLC column effluent with phosphoric acid (1.5 M) and sodium peroxodisulfate  
6 (200 g L<sup>-1</sup>) at a reactor temperature of 99.9 °C. For all experiments flow rates of both reagents  
7 were set to 50 μL min<sup>-1</sup>. The formed CO<sub>2</sub> was separated from the aqueous phase by a gas  
8 exchange membrane and transferred to the open split in a Helium stream (Air Liquide,  
9 Oberhausen, purity 99.999 %) at a flow rate of 1.2 mL min<sup>-1</sup>.  
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19 Without HPLC column this interface can be used for bulk carbon isotope ratio determination  
20 by flow injection analysis (FIA). The carrier flow rate to transport the bulk sample from the  
21 sample loop to the interface was set to 200 μL min<sup>-1</sup>, 350 μL min<sup>-1</sup> and 500 μL min<sup>-1</sup> whereas  
22 reagent flow rates remained at 50 μL min<sup>-1</sup> each.  
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29 The mass spectrometric detection was performed on a Delta V Advantage (Thermo Electron,  
30 Bremen, Germany) tuned on maximum linearity. Linearity and precision of the instrument  
31 were checked regularly by reference gas pulses of different amplitudes.  
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### 36 EA-IRMS

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39 Elemental Analyzer CE 1110 (CE Instruments, Milano, Italy) coupled with a ConFlo IV  
40 Interface to an MAT 253 isotope ratio mass spectrometer (both Thermo Scientific, Bremen,  
41 Germany) was used to determine δ<sup>13</sup>C-values of the analytical standards and bulk isotope  
42 ratios of the homogenized pharmaceutical pills. EA-IRMS values were obtained as described  
43 by Werner and Brand.<sup>18</sup> Reference materials for calibration of the working standard  
44 acetanilide to IAEA scale were NBS-22 (oil; δ<sup>13</sup>C = -30.031 ‰), IAEA-CH-6 (sucrose; δ<sup>13</sup>C  
45 = -10.449 ‰) and IAEA 600 (caffeine; δ<sup>13</sup>C = -27.771 ‰).  
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### 56 Sample Preparation

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3 Samples of pharmaceutical tablets (trade-names *cotrim* and *cotrimoxazole*) of six different  
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5 manufacturers each containing 800 mg sulfamethoxazole and 160 mg trimethoprim were  
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7 homogenized and dissolved with the HPLC eluent to 100 mg L<sup>-1</sup> of the specific analyte  
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9 according to the concentration declaration in the package inserts. Besides the active  
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11 ingredients the pills contained excipients like starch and stearate salts.  
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### 14 15 **Data acquisition and handling**

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18 Acquisition and processing of data was performed by Isodat 2.5. The background subtraction  
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20 algorithm used for these measurements was “individual background”. All reported  $\delta^{13}\text{C}$ -  
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22 values are normalized to the VPDB scale. All standard deviations refer to triplicate  
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24 measurements if not stated otherwise.  $\delta^{13}\text{C}$ -values of the pharmaceutical samples were  
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26 corrected with the difference of  $\delta^{13}\text{C}$ -values between EA/IRMS measurements of the  
27  
28 particular compound and HT-LC/IRMS analyses. This corresponds to the principle of  
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30 identical treatment as it was proposed by Werner and Brand.<sup>18</sup>  
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34 The reference gas pulses were used as an internal standard to calculate relative peak area per  
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36 carbon. The pressure of the reference gas remained unchanged during the FIA measurements  
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38 so that any variation of ionization between the runs is eliminated and peak areas can be  
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40 compared.  
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44 Chromatographic resolution  $R_s$  was calculated according to equation:<sup>19</sup>  
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$$47 \quad R_s = 2 \left( \frac{t_{r(2)} - t_{r(1)}}{w_{h(2)} + w_{h(1)}} \right)$$

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52 Here,  $t_r$  is the retention time of peak 1 or 2 and  $w_h$  is the peak width at half height for the two  
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54 peaks.  
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## Results and discussion

### Chromatographic conditions

All selected compounds can be baseline separated by the use of a temperature gradient. Figure 1 shows a chromatogram and the temperature program applied. Resolution between sulfadiazine and sulfathiazole was three, which meets the requirement of a full separation ( $R = 2$ )<sup>19</sup> for compound-specific isotope analysis due to chromatographic isotope effects<sup>20</sup>.

A temperature gradient with a low steepness of  $3\text{ }^{\circ}\text{C min}^{-1}$  was chosen to keep the rising of the baseline as low as possible. Godin et al.<sup>8</sup> reported low precision and accuracy when high rising backgrounds were observed due to steep temperature gradients on a PGC column. In contrast, Zhang et al.<sup>9</sup> did not observe significant loss in precision or accuracy using gradients between  $6$  and  $9\text{ }^{\circ}\text{C min}^{-1}$  for hybrid particles and zirconium based stationary phases. These findings demonstrate the importance of background subtraction algorithms. Godin et al.<sup>8</sup> and Zhang et al.<sup>9</sup> used the “dynamic background” function in ISODAT. In this work the slope of the background was between  $0.17$  and  $0.5\text{ mV s}^{-1}$  so that individual background algorithm was used as suggested by Zhang et al.<sup>9</sup>.

All sulfonamides have shown to be thermally stable during the chromatographic run. Peak shape as well as comparable  $\delta^{13}\text{C}$ -values to FIA of single compounds showed no indications of degradation inside the column. Only trimethoprim showed a large tailing which can either be a sign of thermal decomposition in the column or column overload. These triangular peak shapes at low pH have been associated to column overload of protonated basic substances on silica-based columns due to charge repulsion between retained protonated molecules<sup>21, 22</sup> (see **Error! Reference source not found.**).

As pointed out by Godin et al.<sup>8</sup> thermal decomposition inside the column can influence  $\delta^{13}\text{C}$ -values. Since results obtained by FIA and HT-LC are comparable (see Table 2), we suggest

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3 that even if thermal degradation of trimethoprim takes place all metabolites elute as one peak  
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5 and leave  $\delta^{13}\text{C}$ -values unaffected.  
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### 8 **Method detection limits and accuracy**

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11 Method detection limits (MDL) were calculated by the moving mean procedure<sup>23</sup> in order to  
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13 account for any method-related offsets of  $\delta^{13}\text{C}$ -values. Here, the detection limit is defined as  
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15 lowest concentration which is necessary to achieve the measurement of a  $\delta^{13}\text{C}$ -value with a  
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17 defined precision ( $\pm 0.5$  ‰) and accuracy with respect to signal height-independent isotope  
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19 ratio<sup>23</sup>.  
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23 MDLs of the selected sulfonamides and trimethoprim are given in Table 2. Sulfathiazole  
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25 showed the lowest MDL of the selected compounds being 0.3  $\mu\text{g}$  on column. If a poorer  
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27 precision of isotope data is acceptable measurements of sulfadiazine, sulfamethoxazole and  
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29 sulfathiazole can be performed even below those MDLs given in Table 2 (see **Error!**  
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35 Comparing the results obtained by EA and both LC-IRMS measurement modes it can be seen  
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37 that there is a discrepancy of  $\delta^{13}\text{C}$ -values for all selected compounds (see Table 2). This is in  
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39 agreement with other studies where substance-specific differences of more than 1‰ have  
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41 been reported for various substances<sup>9, 24, 25</sup>. Although  $\delta^{13}\text{C}$ -values from EA measurements of  
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43 the pure phase analytical standards might be biased by a contamination with other carbon  
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45 compounds, the authors believe that even with a compound with a high  $\delta^{13}\text{C}$ -value, such as a  
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47 carbonate, the error is smaller than the analytical precision.  
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51 FIA/IRMS analyses have been conducted at different liquid carrier flow rates but constant  
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53 reagent flow rates to simulate the influence of HT-LC flow rate without compromising peak  
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55 shape in order to investigate this offset. As can be seen from **Error! Reference source not**  
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57 **found.** the lower the carrier flow rate in the interface, the more do  $\delta^{13}\text{C}$ -values of FIA  
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3 approach  $\delta^{13}\text{C}$ -values of EA measurements. The peak area also shows a dependency on flow  
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5 rate which confirms that oxidation is incomplete. Lower peak areas are observed in FIA when  
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7 higher flow rates are applied because of reduced reaction time inside the oxidation reactor and  
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9 lower reagent concentrations. The reagent is always delivered in far excess relative to the  
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11 analyte so that concentration effects are irrelevant. In **Error! Reference source not found.**  
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13 the same data were used to compare the difference between EA measurements and FIA  
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15 measurements with the relative carbon sensitivity (peak area per nmol C relative to the  
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17 reference gas peak area) from flow rate variation. It shows that the sulfonamides have a  
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19 similar mineralization mechanism assuming that  $^{13}\text{C}$  and  $^{12}\text{C}$  are equally distributed in the  
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21 standard compounds.  
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26 Incomplete extraction of formed  $\text{CO}_2$  in the interface is not the reason for this bias since  
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28 trimethoprim would follow the same trend as the sulfonamides and lead to a constant offset  
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30 independent of substance. For all analytes  $\delta^{13}\text{C}$ -values were not dependent on concentration  
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32 (see Figure 2). If incomplete gas extraction caused an isotopic fractionation,  $\delta^{13}\text{C}$ -values of all  
33  
34 analytes would be dependent on concentration to the same degree. Thus, this reproducible  
35  
36 offset obviously derives from incomplete oxidation.  
37  
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41 Smaller peak areas from incomplete conversion are unfavourable because of the loss in  
42  
43 sensitivity resulting in the need to use lower flow rates in the HT-LC. On the other hand, a  
44  
45 reduction of the HT-LC flow rate could lead to poor peak shape especially at the end of the  
46  
47 chromatographic run where higher temperatures are employed and actually higher flow rates  
48  
49 should be used in order not to affect efficiency<sup>26</sup>. Columns with a smaller inner diameter can  
50  
51 be used due to their lower optimum flow rate but they might be overloaded by the injected  
52  
53 sample amount required for LC/IRMS.<sup>8</sup>  
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3 For sulfamethoxazole, sulfamerazine and trimethoprim  $\delta^{13}\text{C}_{\text{HTLC}}$  values fit  $\delta^{13}\text{C}_{\text{FIA}}$  values  
4  
5 obtained at a flow rate of  $500 \mu\text{L min}^{-1}$ . Sulfadiazine and sulfathiazole show a small  
6  
7 difference of about 0.9 ‰.  
8

9  
10 Therefore,  $\delta^{13}\text{C}$  values from samples have to be corrected by external standards if absolute  
11  
12 values are needed.  
13

### 14 15 16 17 18 19 **Sample analysis**

20  
21 Using common detection methods such as UV/vis or organic mass spectrometry many  
22  
23 carbon-bearing substances are mostly not interfering with detection of the target analytes, in  
24  
25 particular if analyte-specific wavelengths or mass-to-charge ratios are selected. Here,  
26  
27 unexpected matrix compounds can cause errors in analyte peak area, e.g. if they co-elute or  
28  
29 suppress ionization of the target analyte. But a full baseline is not a prerequisite for an  
30  
31 accurate concentration measurement; coelutions are accepted for the sake of analysis time. In  
32  
33 contrast, in LC-IRMS, any carbon-containing compound is detected making baseline  
34  
35 separation of matrix from the target compounds a crucial step in compound-specific  
36  
37 LC/IRMS measurements. Overlapping peaks often cause inaccurate isotopic results as pointed  
38  
39 out by McCullagh<sup>7</sup>.  
40  
41  
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44  
45 Pharmaceutical samples gave no noteworthy contribution of matrix peaks to the  
46  
47 chromatogram (see **Error! Reference source not found.B**). In the small set of samples  
48  
49 analyzed in this work the variation of the corrected  $\delta^{13}\text{C}$  values of sulfamethoxazole and  
50  
51 trimethoprim from different manufacturers was small. However, there was a substantial  
52  
53 difference between bulk sample  $\delta^{13}\text{C}$  values and compound-specific  $\delta^{13}\text{C}$  values (see Table 3).  
54  
55

56  
57 It was observed that an aged column (column 1 in Table 3; >300 injections), showed an  
58  
59 increased background signal due to column bleeding. To investigate the effect of an aged  
60

1  
2  
3 column on the measurement of the isotopic signature of the compounds, we compared the  
4  
5 results with a second new column (see column 2 in Table 3). As shown in Table 3, repeated  
6  
7 analysis of the pharmaceutical samples also proved a good repeatability of this method for  
8  
9 most of the compounds. The same  $\delta^{13}\text{C}$ -values were obtained for five of the six measured  
10  
11 samples. An exception was observed for sulfamethoxazole of brand B between the old  
12  
13 (column 1) and the new column (column 2) and trimethoprim, which gave no reliable results.  
14  
15 The difference of about 1 ‰ can be explained by a steep increase of the background caused  
16  
17 by column bleeding.  
18  
19

20  
21 A two dimensional plot of the  $\delta^{13}\text{C}$ -values of trimethoprim and sulfamethoxazole suggests  
22  
23 that the combined analysis could identify falsified antibiotics containing these compounds  
24  
25 (see **Error! Reference source not found.**). Together with bulk  $\delta^{13}\text{C}$ -values the individual  
26  
27 samples can be distinguished clearly from each other. But the analyzed set of samples is too  
28  
29 small to draw statistically confirmed conclusions.  
30  
31  
32

### 33 **Conclusion**

34  
35  
36 In this work we present the first method for compound-specific carbon isotope analysis of  
37  
38 pharmaceuticals, which cannot be separated by conventionally used techniques for LC-IRMS.  
39  
40 Substance amounts as low as 0.3  $\mu\text{g}$  are sufficient to perform a precise analysis. The  
41  
42 applicability of this method for pharmaceutical research and product authentication was  
43  
44 demonstrated. It is also the first step towards a study of the degradation and mobility of these  
45  
46 substances in the environment on an isotopic level. Following appropriate sample enrichment  
47  
48 compound-specific  $\delta^{13}\text{C}$ -values of highly contaminated samples can be easily measured, but  
49  
50 need to be corrected by external standards.  
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3 There are also some indications that the frequent offset of  $\delta^{13}\text{C}$ -values from LC/IRMS and  
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5 EA/IRMS measurements derives from incomplete oxidation suggesting the need for  
6  
7 improvements of the interface for higher sensitivity and accuracy.  
8  
9

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15  
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17  
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23  
24 No. 16120 N) as well as financial support from the German Research Foundation (DFG).  
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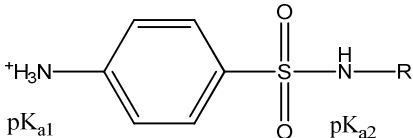
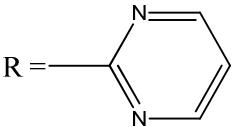
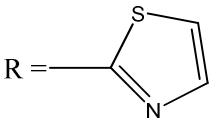
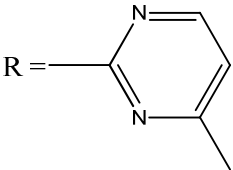
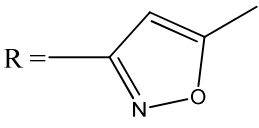
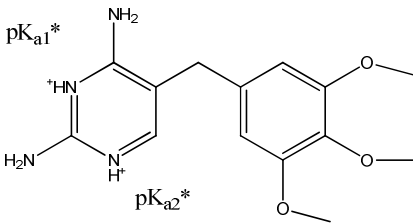
### 28 29 30 **References**

- 31 1. L. T. Corr, R. Berstan and R. P. Evershed, *Rapid Communications in Mass Spectrometry*, 2007,  
32 **21**, 3759-3771.
- 33 2. J. P. Godin, J. Hau, L. B. Fay and G. Hopfgartner, *Rapid Communications in Mass*  
34 *Spectrometry*, 2005, **19**, 2689-2698.
- 35 3. M. Krummen, A. W. Hilker, D. Juchelka, A. Duhr, H. J. Schluter and R. Pesch, *Rapid Commun.*  
36 *Mass Spectrom.*, 2004, **18**, 2260-2266.
- 37 4. S. Bode, K. Deneff and P. Boeckx, *Rapid Communications in Mass Spectrometry*, 2009, **23**,  
38 2519-2526.
- 39 5. D. J. Morrison, K. Taylor and T. Preston, *Rapid Communications in Mass Spectrometry*, 2010,  
40 **24**, 1755-1762.
- 41 6. J. P. Godin, D. Breuille, C. Obled, I. Papet, H. Schierbeek, G. Hopfgartner and L. B. Fay, *Journal*  
42 *of Mass Spectrometry*, 2008, **43**, 1334-1343.
- 43 7. J. S. O. McCullagh, *Rapid Communications in Mass Spectrometry*, 2010, **24**, 483-494.
- 44 8. J. P. Godin, G. Hopfgartner and L. Fay, *Anal. Chem.*, 2008, **80**, 7144-7152.
- 45 9. L. Zhang, D. M. Kujawinski, M. A. Jochmann and T. C. Schmidt, *Rapid Communications in*  
46 *Mass Spectrometry*, 2011, **25**, 2971-2981.
- 47 10. Y. Yang, M. Belghazi, A. Lagadec, D. J. Miller and S. B. Hawthorne, *Journal of Chromatography*  
48 *A*, 1998, **810**, 149-159.
- 49 11. S. Wiese, T. Teutenberg and T. C. Schmidt, *Journal of Chromatography A*, 2011, **1218**, 6898-  
50 6906.
- 51 12. T. Teutenberg, K. Hollebekkers, S. Wiese and A. Boergers, *Journal of Separation Science*,  
52 **2009**, **32**, 1262-1274.
- 53 13. J. D. Thompson and P. W. Carr, *Analytical Chemistry*, 2002, **74**, 1017-1023.
- 54 14. K. Schauss, A. Focks, H. Heuer, A. Kotzerke, H. Schmitt, S. Thiele-Bruhn, K. Smalla, B.-M.  
55 Wilke, M. Matthies, W. Amelung, J. Klasmeier and M. Schloter, *Trac-Trends Anal. Chem.*,  
56 **2009**, **28**, 612-618.  
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15. P. Huovinen, *Clinical Infectious Diseases*, 2001, **32**, 1608-1614.
16. L. Pereira, S. Aspey and H. Ritchie, *Journal of Separation Science*, 2007, **30**, 1115-1124.
17. S. Giegold, T. Teutenberg, J. Tuerk, T. Kiffmeyer and B. Wenclawiak, *Journal of Separation Science*, 2008, **31**, 3497-3502.
18. R. A. Werner and W. A. Brand, *Rapid Communications in Mass Spectrometry*, 2001, **15**, 501-519.
19. J. V. Hinshaw, *Lc Gc Europe*, **23**, 362-+.
20. M. Blessing, M. A. Jochmann and T. C. Schmidt, *Anal. Bioanal. Chem.*, 2008, **390**, 591-603.
21. L. R. Snyder, J. J. Kirkland and J. W. Dolan, *Introduction to Modern Liquid Chromatography*, Wiley, Hoboken, New Jersey; USA, 2010.
22. S. M. C. Buckenmaier, D. V. McCalley and M. R. Euerby, *Analytical Chemistry*, 2002, **74**, 4672-4681.
23. M. A. Jochmann, M. Blessing, S. B. Haderlein and T. C. Schmidt, *Rapid Communications in Mass Spectrometry*, 2006, **20**, 3639-3648.
24. S. Reinnicke, A. Bernstein and M. Elsner, *Analytical Chemistry*, 2010, **82**, 2013-2019.
25. C. I. Smith, B. T. Fuller, K. Choy and M. P. Richards, *Analytical Biochemistry*, 2009, **390**, 165-172.
26. T. Teutenberg, *High-Temperature Liquid Chromatography - A User's Guide for Method Development*, RSCPublishing, 2010.
27. Z. M. Qiang and C. Adams, *Water Research*, 2004, **38**, 2874-2890.

**Table 1** Physico-chemical properties and structures of the investigated substances

Compound	$pK_{a1}^{27}$	$pK_{a2}^{27}$	Molecular formula ( $MW/g\ mol^{-1}$ )	CAS-No.
				
Sulfadiazine	1.6	6.8	C <sub>10</sub> H <sub>10</sub> N <sub>4</sub> O <sub>2</sub> S (250.28)	68-35-9
				
Sulfathiazole	2.01	7.11	C <sub>9</sub> H <sub>9</sub> N <sub>3</sub> O <sub>2</sub> S <sub>2</sub> (255.32)	144-74-1
				
Sulfamerazine	2.07	6.9	C <sub>11</sub> H <sub>12</sub> N <sub>4</sub> O <sub>2</sub> S (264.31)	127-58-2
				
Sulfamethoxazole	1.85	5.60	C <sub>10</sub> H <sub>11</sub> N <sub>3</sub> O <sub>3</sub> S (253.28)	723-46-6
				
Trimethoprim	3.23	6.76	C <sub>14</sub> H <sub>18</sub> N <sub>4</sub> O <sub>3</sub> (290.32)	738-70-5
				

\*single mesomeric structure showing the initial

protonation sites

**Table 2 HTLC-IRMS performance evaluation**

	MDL*	$\delta^{13}\text{C}_{HT-LC}$	$\delta^{13}\text{C}_{FIA}^{\ddagger}$	$\delta^{13}\text{C}_{EA}$	$\delta^{13}\text{C}_{EA}-\delta^{13}\text{C}_{HT-LC}$
	[ $\mu\text{g}$ ]	[ $\text{‰}$ ] <sup>#</sup>	[ $\text{‰}$ ] <sup>#</sup>	[ $\text{‰}$ ] <sup>#</sup>	[ $\text{‰}$ ]
Sulfadiazine	0.4	$-30.8 \pm 0.1$	$-31.6 \pm 0.4$	$-29.2 \pm 0.2$	1.6
Sulfathiazole	0.3	$-28.5 \pm 0.1$	$-29.6 \pm 0.4$	$-26.64 \pm 0.04$	1.9
Sulfamerazine	0.5	$-31.3 \pm 0.3$	$-31.2 \pm 0.4$	$-28.90 \pm 0.03$	2.4
Sulfamethoxazole	0.5	$-30.7 \pm 0.3$	$-30.3 \pm 0.2$	$-27.91 \pm 0.02$	2.8
Trimethoprim	0.4	$-37.6 \pm 0.3$	$-37.6 \pm 0.2$	$-34.15 \pm 0.04$	3.4

\*as absolute amount on column

<sup>#</sup>Standard deviations refer to at least triplicate measurement<sup>‡</sup>at a flow rate of 500  $\mu\text{L min}^{-1}$ **Table 3** Bulk and corrected compound-specific  $\delta^{13}\text{C}$ -values of the measured pharmaceutical samples.

Brand	$\delta^{13}\text{C}_{bulk}$	Sulfamethoxazole	Sulfamethoxazole	Trimethoprim
	[ $\text{‰}$ ]	$\delta^{13}\text{C}$ [ $\text{‰}$ ] column 1 <sup>a)</sup>	$\delta^{13}\text{C}$ [ $\text{‰}$ ] column 2 <sup>b)</sup>	$\delta^{13}\text{C}$ [ $\text{‰}$ ] column 2
A	$-29.6 \pm 0.1$	$-27.6 \pm 0.2$	$-27.7 \pm 0.4$	$-31.1 \pm 0.3$
B	$-29.7 \pm 0.4$	$-27.6 \pm 0.3$	$-28.6 \pm 0.4$	$-30.9 \pm 0.3$
C	$-28.46 \pm 0.09$	$-28.4 \pm 0.3$	$-28.4 \pm 0.2$	$-31.9 \pm 0.7$
D	$-28.5 \pm 0.1$	$-28.0 \pm 0.2$	$-28.1 \pm 0.2$	$-32.1 \pm 0.3$
E	$-30.0 \pm 0.4$	$-27.4 \pm 0.2$	$-27.2 \pm 0.4$	$-30.8 \pm 0.4$
F	$-29.67 \pm 0.04$	$-28.2 \pm 0.2$	$-28.4 \pm 0.1$	$-31.7 \pm 0.4$

<sup>a)</sup>Old column >300 measurements.<sup>b)</sup>New column

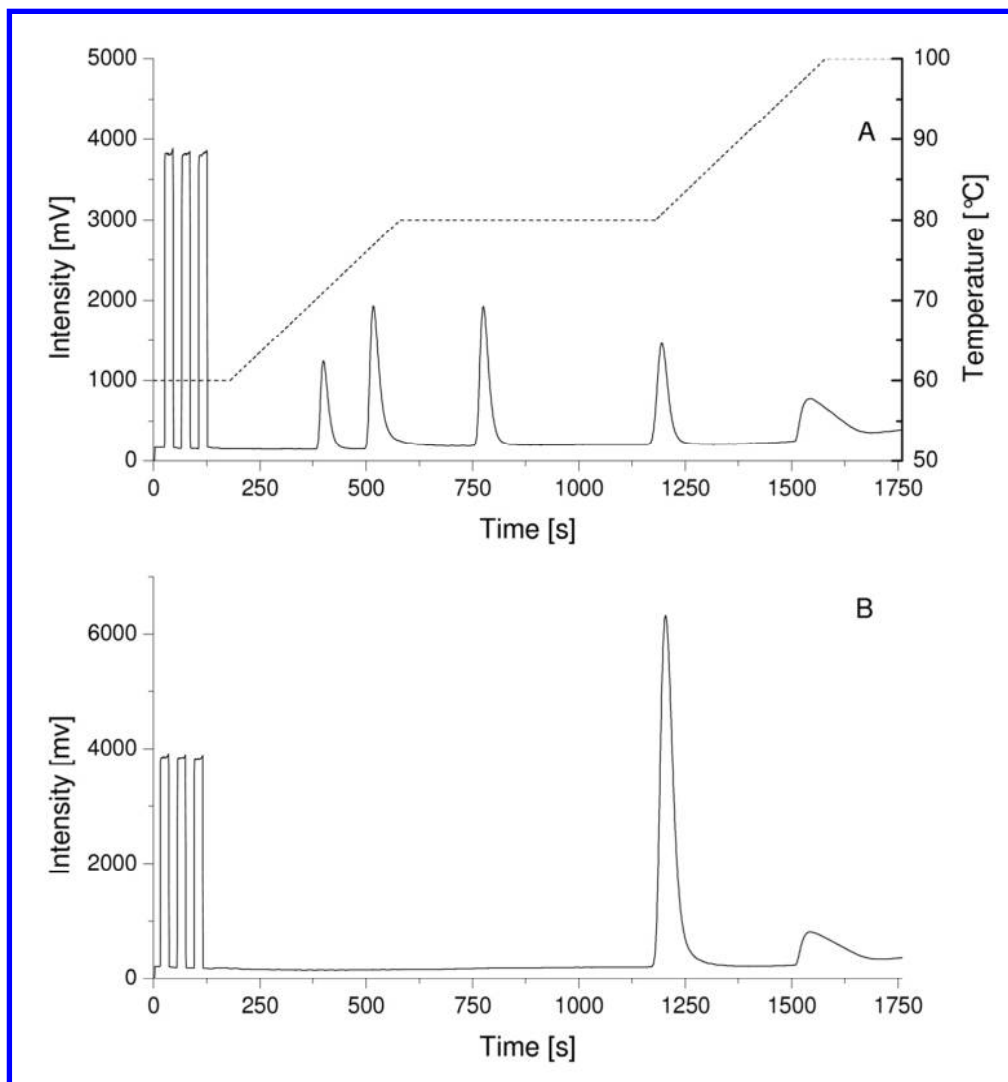


Figure 1 A) Chromatogram ( $m/z$  44) of a standard mixture separated by a temperature gradient indicated by the dashed line. The elution order is sulfadiazine, sulfathiazole, sulfamerazine, sulfamethoxazole and trimethoprim. The concentration of each substance was  $100 \text{ mg L}^{-1}$ . B) Chromatogram ( $m/z$  44) of a pharmaceutical sample containing sulfamethoxazole and trimethoprim obtained by the same temperature gradient as above.  
122x131mm (300 x 300 DPI)

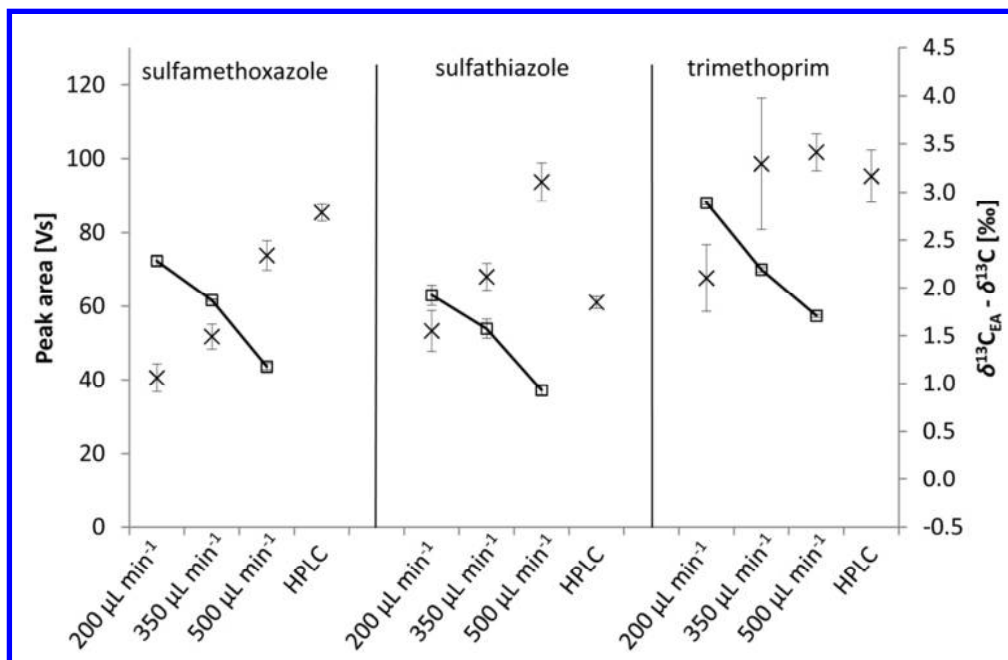


Figure 2 Peak areas (boxes) and  $\delta^{13}\text{C}$  values (crosses) determined by FIA at various carrier flow rates and HTLC at 500  $\mu\text{L min}^{-1}$ . No peak areas are given for HTLC measurements due to incomparable ionization conditions with respect to FIA. Similar trends can be observed for sulfadiazine and sulfamerazine.  
102x66mm (300 x 300 DPI)

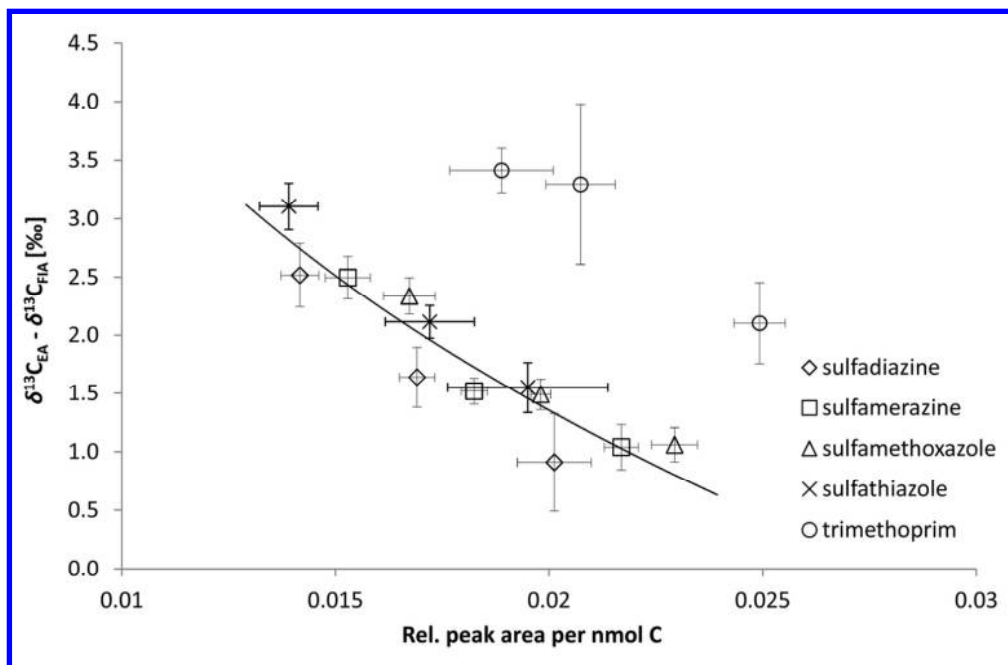


Figure 3 Dependency of the difference between EA/IRMS and FIA/IRMS derived  $\delta^{13}\text{C}$ -values on the relative peak area per nmol carbon obtained by different carrier flow rates. Peak areas per carbon amount were referenced to the reference gas peak in order to account for differences in ionization between the single measurements. The solid line represents a logarithmic fit of the sulfonamide results.  
108x70mm (300 x 300 DPI)

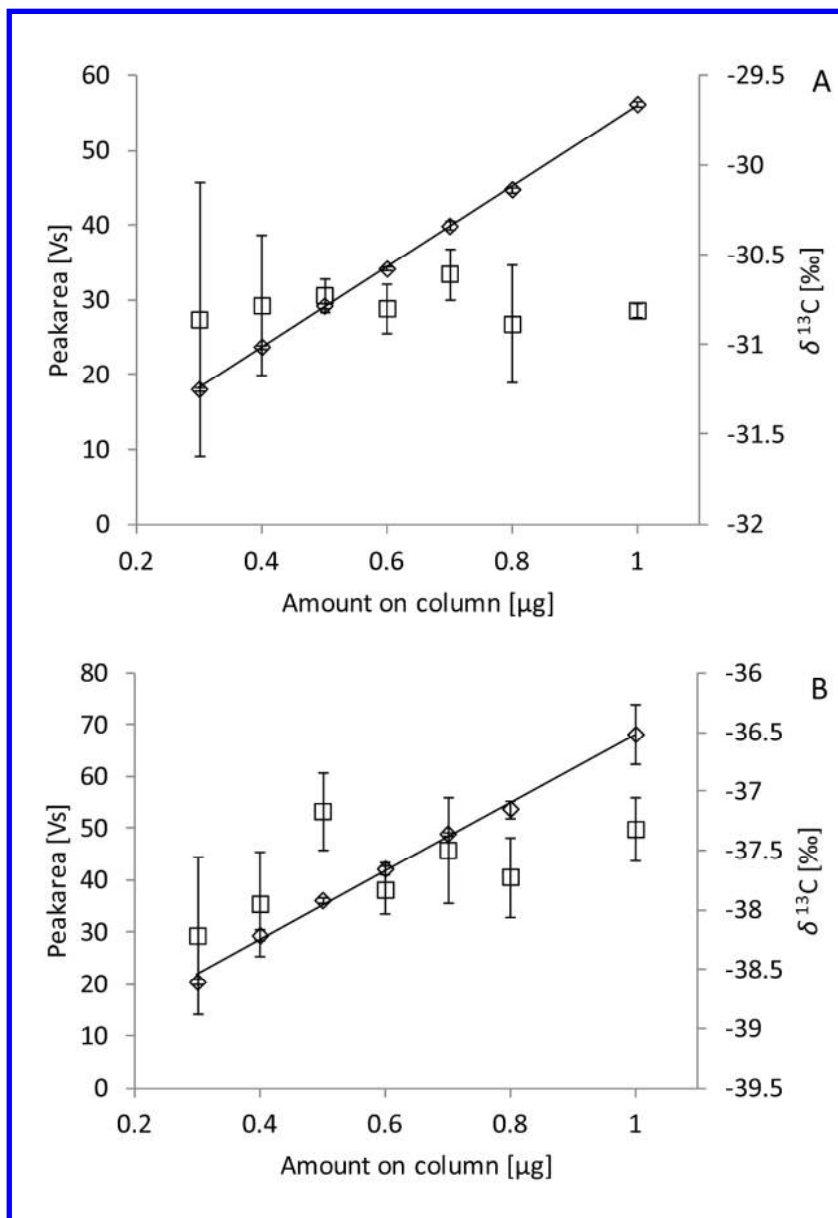


Figure 4 Representative calibration line of sulfadiazine (graph A) and trimethoprim (graph B). Diamonds correspond to the peak area of the  $m/z$  44 trace whereas boxes relate to  $\delta^{13}\text{C}$  values.  
226x327mm (300 x 300 DPI)

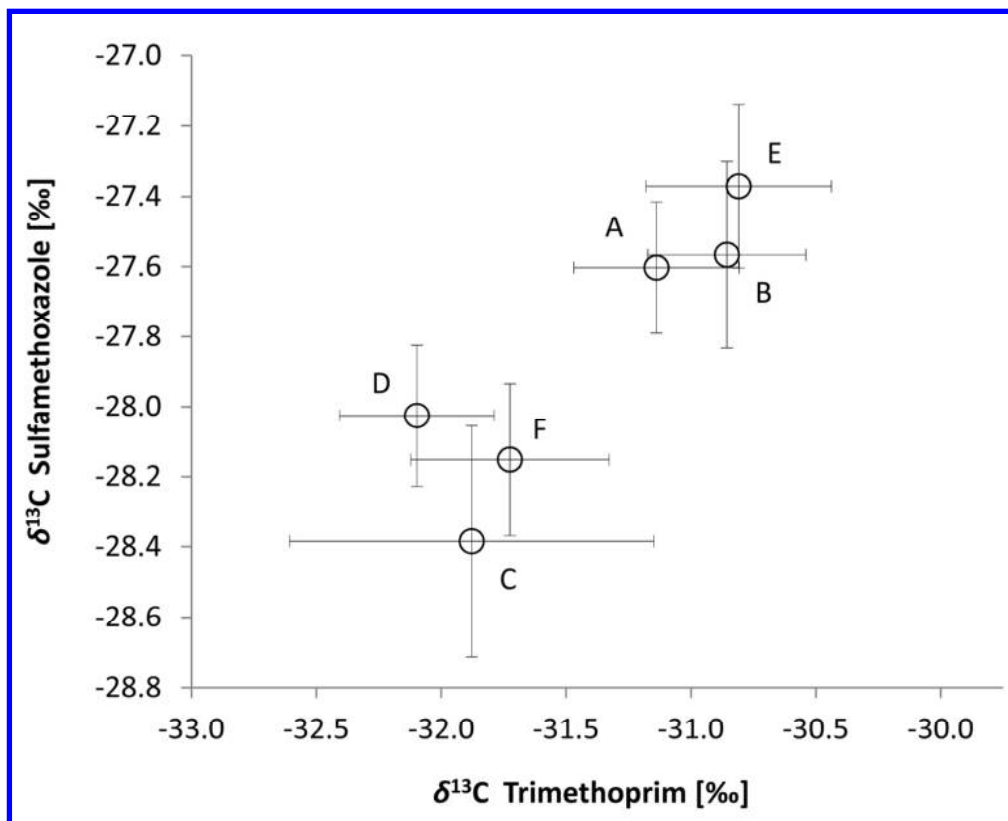
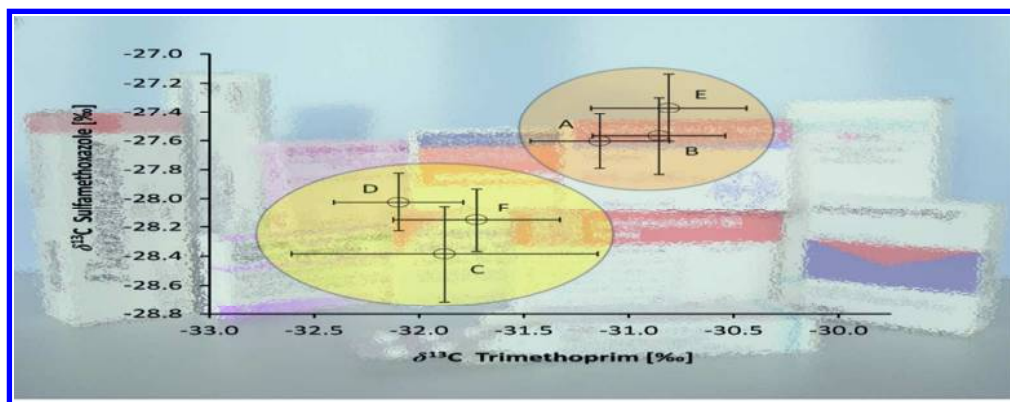


Figure 5 Corrected  $\delta^{13}\text{C}$ -values of trimethoprim and sulfamethoxazole from pharmaceutical products.  
132x107mm (300 x 300 DPI)





For TOC  
89x34mm (300 x 300 DPI)

## **When other separation techniques fail**

Kujawinski, Dorothea M.; Zhang, Lijun; Schmidt, Torsten Claus; Jochmann, Maik A.

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