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1	Investigation of degradation products of cocaine and benzoylecgonine in
2	the aquatic environment
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14 ABSTRACT

15 In this work, ultra-high-performance liquid chromatography (UHPLC) coupled to a 16 hybrid quadrupole time-of-flight mass spectrometer (QTOF MS) has allowed the discovery 17 and elucidation of degradation products of cocaine and its main metabolite benzoylecgonine 18 (BE) in water. Spiked surface water was subjected to hydrolysis, chlorination and photo-19 degradation (both ultraviolet irradiation and simulated sunlight). After degradation of cocaine, 20 up to sixteen compounds were detected and tentatively identified (1 resulting from hydrolysis; 21 8 from chlorination; 7 from photo-degradation), three of which are well known cocaine 22 metabolites (BE, norbenzoylecgonine and norcocaine). Regarding BE degradation, up to ten compounds were found (3 from chlorination; 7 from photo-degradation), including one 23 24 known metabolite (norbenzoylecgonine). Since reference standards were available for the 25 major metabolites, they could be confirmed using information on retention time and fragment 26 ions. The other degradates resulted from chlorination, dealkylation, hydroxylation and nitration, or from a combination of these processes. Several influent and effluent sewage 27 28 water, and surface water samples were then screened for the identified compounds (known 29 and unknown) using UHPLC-tandem MS with triple quadrupole. BE, norcocaine and 30 norbenzoylecgonine were identified in these samples as major metabolites. Four previously 31 unreported degradates were also found in some of the samples under study, illustrating the 32 usefulness and applicability of the degradation experiments performed in this work.

33

34 Keywords

35 Cocaine, degradation and transformation products, water, time-of-flight mass spectrometry.

37 1. INTRODUCTION

38 Cocaine use has increased during the last decade and is the illicit drug with the 39 second-highest consumption in Europe, behind only cannabis (EMCDDA 2010). After 40 consumption and excretion, cocaine enters the sewage treatment plants (STPs) as the parent 41 drug or as human metabolites (mainly benzoylecgonine (BE)) and may end up in the 42 receiving surface waters as a consequence of incomplete elimination in the STPs. In most 43 studies, if the presence of cocaine in the aquatic environment is reported, only the parent 44 compound and a few relevant metabolites, commonly BE and cocaethylene or ecgonine 45 methyl ester are included (Baker and Kasprzyk-Hordern 2011). Occasionally, in monitoring 46 studies dealing with sewage- and surface water, some minor metabolites have been found, 47 such as norBE and norcocaine (e.g. Chiaia et al. 2008; Zuccato et al. 2008; Bijlsma et al. 48 2009; Bisceglia et al. 2010). Although concentrations reported in surface water are generally 49 low (i.e. 7 - 60 ng/L for cocaine and 15 - 191 ng/L for BE (Huerta-Fontela et al. 2008; Gheorghe et al. 2008)), there is a potential negative impact of their presence in the aquatic 50 51 ecosystem (Binelli et al. 2012). Especially, the effects of combined exposure to multiple 52 compounds are of potential concern.

53 In order to evaluate the hazard in the water cycle, not only removal of the parent 54 compounds and metabolites in the treatment processes must be taken into account, but also 55 the possible formation of degradation/transformation products (TPs). In some countries (e.g. 56 Italy), chlorination is progressively abandoned because of its potential for generating 57 unwanted TPs and replaced by UV irradiation (Antonelli et al. 2008). Furthermore, after 58 incomplete elimination during chlorination (Huerta-Fontela et al. 2008; Boleda et al. 2009), 59 cocaine and BE which ended up in surface water may be exposed to natural sunlight and 60 produce photo-degradation products. The same would occur for cocaine and BE still present 61 in treated wastewater when no tertiary treatment is applied in the STP (e.g. Gheorghe et al. 62 2008; Huerta-Fontela et al. 2008; Bijlsma et al. 2009; Bisceglia et al. 2010). Despite the fact 63 that some TPs are more persistent or might exhibit similar toxicity than their parent 64 compounds (Farré et al. 2008; Kern et al. 2009; Fatta-Kassinos et al. 2011; Metz et al. 2011), 65 the research on TPs of illicit drugs has received little attention. Nevertheless, investigation of 66 TPs is of importance to know the overall contribution of chemicals in the environment. 67 Information on potential TPs that may be present in the environment can be used to set-up 68 monitoring studies in order to get a wider and more realistic view on the impact of cocaine on 69 the aquatic environment.

70 The identification of TPs in the aquatic environment, especially unknown ones, is a 71 challenging task for analytical chemists and commonly various techniques and/or analytical 72 reference standards are necessary for a reliable confirmation (Wick et al. 2011). An important 73 analytical tool in the elucidation of TPs is high resolution mass spectrometry (HRMS), with 74 analyzers like Orbitrap and time-of-flight (TOF). The accurate mass full-spectrum acquisition 75 and the possibility to obtain fragment ions by coupling HRMS to ion trap or quadrupole 76 analyzers is highly suitable and helpful for the proposal of convincing molecular structures 77 (Ibañez et al. 2004; Farré et al. 2008; Quintana et al. 2010; Metz et al. 2011).

78 Laboratory degradation experiments in combination with HRMS are one of the most 79 useful tools to identify TPs that can be formed in the aquatic environment. They have been 80 applied mainly to elucidate pesticide and pharmaceutical TPs formed in water (Ibañez et al. 81 2004; Hernández et al. 2008; Quintana et al. 2010; Wick et al. 2011). Treatment conditions applied by STPs, e.g. chlorination and UV irradiation, can be simulated, as well as natural 82 83 sunlight. The most important TPs identified can subsequently be included in multi-residue LC 84 tandem MS methods with triple quadrupole. This has allowed the detection of parent 85 compounds and of their related TPs in sewage-, surface- and/or drinking water (Hernández et al. 2008; Quintana et al. 2010; Wick et al. 2011), and illustrates the importance of 86 87 investigating TPs.

The use of MS^E is an attractive option, which is feasible working with hybrid QTOF 88 89 MS instruments. Using this approach, information on both (de)protonated molecules and their 90 fragment ions is acquired simultaneously in a single injection (Hernández et al. 2011). The 91 accurate mass measurement of the (de)protonated molecule generally allows the assignment 92 of a highly probable molecular formula. Subsequently, fragment ions as well as neutral losses 93 can be investigated in order to elucidate the structure of the TPs detected. Available software 94 for the detection of metabolites and TPs are usually offered by MS manufacturers. They 95 compare and contrast data of a presumptive positive sample with a control or blank sample. 96 This facilitates data processing and might even detect (low abundant) compounds overlooked 97 by visual inspection.

98 The objective in this paper was to perform a study on TPs of cocaine and BE that 99 might be found in the aquatic environment. Several laboratory controlled degradation 100 experiments (i.e. hydrolysis, chlorination, and photo-degradation under ultraviolet (UV) 101 irradiation and simulated sunlight) have been carried out and the TPs formed investigated by 102 LC-QTOF under MS^E mode. To the best of our knowledge, several unknown TPs reported in

- 103 this study have not previously described in the literature. In a subsequent step, influent and
- 104 effluent sewage water, and also surface waters, were searched for the identified TPs.

106 2. MATERIALS AND METHODS

107

108 **2.1. Reagents and chemicals**

109 Cocaine, norcocaine, BE and norbenzoylecgonine (norBE) reference standards were 110 purchased from the National Measurement Institute (Pymble, Australia) and Cerilliant (Round 111 Rock, TX, USA). Standard solutions of cocaine and BE were prepared at 500 mg/L in 112 acetonitrile (ACN) and methanol (MeOH), respectively. Intermediate work solutions (50 113 mg/L) were made by diluting the solution ten times with MeOH.

HPLC-grade MeOH, ACN and formic acid (FA) were acquired from Scharlau
(Barcelona, Spain). Sodium hypochlorite solution (available chlorine 10%) was obtained from
Sigma-Aldrich. A Milli-Q ultra-pure water system from Millipore (Bedford, MA, USA) was
used to obtain the HPLC grade water. Leucine enkephalin and imazalil were purchased from
Sigma-Aldrich and Dr. Ehrenstorfer (Augsburg, Germany), respectively.

119 Solid-phase extraction (SPE) cartridges (Oasis-HLB; 3mL, 60 mg) were purchased 120 from Waters (Milford, MA, USA). Prior to use, the SPE cartridges were conditioned by 121 washing and rinsing with 3 mL of MeOH and 3 mL of Milli-Q water.

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- 123

3 **2.2. Degradation experiments**

Blank surface water from the Mijares River (Castellón, Spain) was collected in November 2010 and used for all laboratory controlled experiments. Surface water (pH 8.1) was selected in order to simulate reality, as it contains matrix components which may affect degradation.

Surface water used for hydrolysis, chlorination and photo-degradation experiments was spiked with cocaine or BE at a concentration of 0.5 mg/L. This relatively high concentration allowed better evaluation of degradation products, and especially facilitated the detection of minor TPs. Non-spiked surface water samples were subjected to the same degradation processes and used as control samples.

The hydrolysis and chlorination experiments were performed at room temperature and in darkness. Regarding chlorination, 40 μ L of ten-fold diluted sodium hypochlorite solution was added to 50 mL of each surface water sample. During the experiment, 2 mL aliquots of the water sample were collected at several time intervals (0, 30 min, 1, 3, 10 h, 1, 3, 7, 11 and

- 137 15 days for hydrolysis; and 0, 30 min, 1 and 3 hours for chlorination), after stirring of the 138 water solutions, and were immediately stored at -20 °C until analysis.
- 139 Photo-degradation experiments were carried out under UV irradiation and simulated 140 sunlight. UV irradiation was performed using a mercury lamp with its main output at 254 nm. 141 250 mL surface water samples were kept in quartz glass vessels at a distance of ~15 cm from 142 the lamp. The experiment was carried out in a fume hood at room temperature over a period 143 of 72 h under constant stirring of the samples. Sunlight was simulated using a solar simulation 144 system (Suntest XLS+, Atlas MTT, Linsengericht, Germany), equipped with a xenon arc 145 lamp as radiation source and a solar light filter allowing a wavelength in the range of 300 -800 nm. The radiation intensity was set to 500 W/m^2 and the light dose per hour of irradiation 146 147 to 1.8 MJ/h. In this way, 90 irradiation hours corresponds to 15 days of natural sun light (dose: 288 MJ/m²). The degradation was performed using 250 mL closed quartz glass vessels 148 149 and sample temperature was set to 25°C in order to minimize sample evaporation and possible 150 thermal transformation. Aliquots were sampled after stirring of the water solution. The first 2 151 mL water aliquots were analysed, prior to the irradiation experiments (t = 0). During 152 irradiation experiments, 2 mL water samples were taken at different time intervals (see 153 hydrolysis experiment), and immediately stored at -20 °C until analysis.
- 154

155 **2.3. Instrumentation**

For identification and elucidation of TPs, a Waters Acquity ultra-high-performance liquid chromatography (UHPLC) system was interfaced to a hybrid quadrupole orthogonal acceleration time-of-flight mass spectrometer (Q-TOF Premier, Waters Micromass) equipped with an orthogonal Z-spray electrospray ionization interface (ESI) operating in both positiveion and negative-ion mode and controlled by MassLynx v 4.1 software. Leucine enkephalin was used as the lock mass (m/z 556.2771 in positive-ion and m/z 554.2615 in negative-ion mode) ensuring typically mass errors below 2 mDa.

163 The UHPLC separation was performed using an Acquity UPLC BEH C18, 1.7 μ m 164 particle size analytical column, 100 mm × 2.1 mm (Waters). The mobile phases used were A 165 = H₂O and B = MeOH, both with 0.01% FA. The percentage of organic modifier (B) was 166 changed linearly as follows: 0 min, 10%; 9 min, 90%; 11 min, 90%; 11.1 min, 10%; 14 min, 167 10%. The flow rate was 0.3 mL/min.

For MS^E experiments, two acquisition functions were created and simultaneously used within the same run: the low-energy function (LE) with a collision energy of 4 eV, where mainly the (de) protonated intact molecules are observed, and the high energy (HE) function with a collision energy ramp ranging from 15 to 40 eV, where fragmentation is promoted. The
same collision energy ramp was used for additional MS/MS experiments. The optimized cone
voltage (15 V) and collision energy ramp were identical for both cocaine and BE and seemed
therefore most adequate for the screening of their corresponding degradation products.
Further details on instrument operating conditions can be found elsewhere (Hernández et al.
2011).

For screening of TPs in sewage waters, a TQD triple quadrupole (QqQ) mass spectrometer with electrospray ionization source (Waters) was used. Chromatographic separation was performed using the same analytical column and gradient as used in UHPLC-QTOF analysis. The analysis of surface waters was performed under similar conditions using the TQS (QqQ) mass spectrometer (Waters).

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183 **2.4. Elucidation / identification procedure**

Waters MetaboLynx software (an application manager within MassLynx) was used to
compare accurate mass data of spiked and blank (non-spiked) samples from the laboratory
experiments.

The data comparison by MetaboLynx was performed in two ways. First, for expected TPs, (bio)transformation processes reported in the literature were included in the processing settings. These consisted of a mass window \pm 10 mDa for extracted ion chromatograms (XICs) of each specific exact mass; peaks with less than 10 area units were eliminated. Second, searching for unexpected TPs was performed by mass spectral comparison of nonspiked *versus* spiked samples. XICs were automatically generated for each sample (spiked and non-spiked) over a range from m/z 70 to 550 Da, at 1 Da mass window, and compared.

The most likely elemental compositions of (de)protonated molecules were calculated based on accurate mass LE spectra of the peaks of interest. The accurate mass HE spectra were then used to calculate possible elemental compositions of fragment ions. Assuming that most TPs share similar fragmentation pathways with the parent drug (Wang and Bartlett 198; Bijlsma et al. 2011), fragmentation was compared to that of cocaine and BE, and the TP structures were proposed.

200

201 **2.5. Water samples**

Five influent and five effluent sewage water samples (24-hour composite) and five surface water grab samples from different locations of the Comunidad Valenciana (Eastern Spain) were collected and immediately stored at -20 °C. Sewage water was collected from STPs of Castellón and Benicàssim, while surface water was collected from the Albufera
national park of Valencia.

100 mL of five-fold diluted (with MilliQ) influent wastewater, 100 mL of effluent wastewater or 100 mL surface water was taken for analysis. The samples were loaded onto the HLB cartridges by gravity, and then cartridges were vacuum-dried for 10 min. Analytes were eluted with 5 mL of MeOH. The extracts were evaporated to dryness at 35°C under a gentle stream of nitrogen and reconstructed in 1 mL of 10:90 MeOH:H₂O. Analyses of cocaine and BE TPs were performed by injecting 20 μ L of the final extract into the UHPLC-TQD system (sewage water) or 100 μ L in the UHPLC-TQS system (surface water).

215 **3. RESULTS AND DISCUSSION**

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Many known TPs of environmental contaminants share similar fragmentation pathways as their parent molecules. Then, knowledge of structures of fragment ions and basic fragmentation rules are helpful for achieving confident TP structure proposals. Isotope fit, Double Bound Equivalent (DBE), and accurate mass of fragments observed in the HE function were used to discard potential chemical formulas in order to obtain the most plausible structures of TPs.

223 The fragmentation of cocaine and BE has been studied previously by our own group 224 (Bijlsma et al. 2011) and by others (Wang and Bartlett 1998). This has facilitated the 225 elucidation of some of the TPs found in this work. The most abundant fragment ions in the 226 mass spectra of both compounds are m/z 105 (C₆H₅CO⁺) and a fragment due to the neutral 227 loss of benzoic acid (122 Da). Subsequent fragmentation of the resulting ion $[M+H - 122]^+$ 228 can produce fragments with m/z 150, 122, 119, 108, 91 and 82, involving a further loss of 229 methanol or water or elimination of part of the bicyclic ring system, followed by hydrogen 230 rearrangement.

231

Proposed structures for the TPs found in this work are shown in Figure 1.

232

3.1. Hydrolysis

Gheorge et al. (2008) performed a detailed study on the stability of BE and cocaine in surface and wastewater, testing at different temperatures and pH values in order to establish optimal conditions for sample storage. Degradation of cocaine was minimal at -20 °C and pH 2. However, in our study, realistic environmental conditions were chosen for the experiments without any adjustment of pH and temperature. It is therefore likely that besides hydrolysis, potential biodegradation might also occur. To some extent, these processes may yield the same products.

Complete cocaine and some BE degradation was observed in surface water after keeping the solution in darkness at room temperature for 15 days (data not shown). Cocaine was mainly transformed into BE through chemical hydrolysis of cocaine ester bonds. Ecgonine methyl ester (EME), another hydrolytic product reported for cocaine (Postigo et al. 2011), was not observed. EME is presumed to be solely an *in vivo* metabolite as a result of enzymatic hydrolysis and for that reason it is unlikely to be formed during cocaine degradation in water (Klette et al. 2000). Gheorghe et al. (2008) had similar results to the present work, where cocaine and EME degraded in spiked surface water, while BE initiallyincreased owing to the possible chemical hydrolysis of cocaine.

250

251 **3.2. Chlorination**

Table 1 summarizes the TPs of cocaine and BE formed during chlorination. Retention times and experimental m/z-values, proposed elemental composition of the protonated TPs and their fragment ions, the mass error in mDa, and the double bond equivalent (DBE) are given.

256 Chlorination TPs of cocaine and BE were investigated under the experimental 257 conditions described in section 2.2. High chlorine concentration (8 mg/L) was used, similar to 258 the conditions employed by STPs for wastewater treatment. In previous studies on acidic 259 pharmaceutical TPs, ascorbic acid was found to be an effective quenching agent to prevent 260 further degradation with chlorine (Quintana et al. 2010). However, in the present study, we 261 observed that it affected the stability of some TPs (the monochlorinated TP-C4, -C7 and -C8 were no longer observed after adding ascorbic acid to the sample vials). Therefore, we did not 262 263 use ascorbic acid addition in our experiments. The sample aliquots taken at different times 264 were frozen, stored and thawed just before analysis. In any case, quenching chlorination 265 seemed not much important in this case, as a fast degradation of cocaine and BE occurred 266 (after 30 minutes neither cocaine nor BE was observed in sample aliquots analyzed).

267 The simultaneous acquisition of accurate mass LE and HE spectra and useful isotopic pattern information (distribution of the ³⁷Cl isotope) obtained in the MS^E mode, allowed the 268 detection and tentative identification of several TPs, in a single injection. Among these TPs, 269 270 some well-known cocaine metabolites, BE (TP-C1), norBE (TP-C2), and norcocaine (TP-C3), 271 were identified and subsequently confirmed by using reference standards. All TPs were 272 determined and identified in the positive-ion mode. Besides the protonated molecules 273 $[M+H]^+$, their sodium adducts $[M+Na]^+$ were also observed surely owing to the presence of 274 sodium in NaClO. Some TPs that contain a carboxylic group could also be analyzed in 275 negative-ion mode; however, analysis under negative mode did not reveal additional TPs to 276 those observed in positive mode.

277 TP-C4 (C₉H₁₅ClNO₃⁺, m/z 220.0740), with an abundant peak at 4.81 min, may be 278 generated via benzoylester cleavage and chlorination. Initial fragmentation involves losses of 279 water and methanol (to ions with m/z 202 and 188, respectively), suggesting that this TP is a 280 secondary product from TP-C3 (norcocaine). 281 Chlorination of cocaine yielded an intense peak at 8.27 min with $[M+H]^+$ m/z 282 324.1003, named as TP-C8 ($C_{16}H_{19}CINO_4^+$), corresponding to demethylation of the 283 bridgehead nitrogen and consecutive halogenation (Figure S1A). The fragmentation of this 284 TP was comparable to cocaine and its metabolites, where the most abundant fragments ions 285 are m/z 105 (C₆H₅CO⁺) and m/z 202 (loss of benzoic acid, 122 Da) (Wang and Bartlett 1998). 286 Secondary fragmentation of the ion m/z 202 ([M+H–122]⁺) involves the loss of either HCl or 287 CH₃OH to ions m/z 166 and 170, respectively, the later indicating that initially N-288 demethylation rather than O-demethylation occurred. The complete fragmentation pathway 289 for TP-C8 is proposed in Figure S1B. The characteristic chlorine isotopic pattern confirms 290 the presence of Cl in the fragment ions with m/z 202, 170 and 142, whereas it is absent in the 291 ions with *m/z* 288, 166, 134 and 105.

Another TP of cocaine, TP-C6 ($[M+H]^+$, m/z 318.1336) is has the same nominal mass as cocaethylene ($C_{18}H_{24}NO_4^+$, m/z 318.1705), but they could be differentiated both chromatographically and by HRMS, as a difference of 36.9 mDa was observed. The most likely molecular formula for TP-C6 is $C_{17}H_{20}NO_5^+$ (m/z 318.1341, Δ 0.5 mDa). Thus, TP-C6 would result from oxidation (+O–2H) of cocaine during chlorination experiments, which probably occurs on the bicyclic ring system, since the characteristic fragment ion with m/z105 ($C_6H_5CO^+$) is still present.

299 Chlorination of BE resulted in TP-C5 ($[M+H]^+$, m/z 304.1185) and TP-C7 300 ($C_{15}H_{17}CINO_4^+$, m/z 310.0846) at retention times of 5.89 min and 7.53 min, respectively 301 (**Table 1**). These compounds show similar fragmentation pathways to TP-C6 and TP-C8, 302 respectively, although with an expected mass shift of -14 in several of the m/z values. These 303 TPs were also observed after cocaine chlorination, where BE probably acted as an 304 intermediate.

The most abundant TPs formed after cocaine chlorination corresponded to TP-C8 and TP-C1 (BE), whereas TP-C5 could be considered as minor TP. The abundance of TP-C2 (norBE), solely formed after chlorination of BE, was in the same order of magnitude as TP-C3 (norcocaine),-C4,-C6 and -C7.

The data obtained was not sufficient to predict the exact position of the chlorine or keto group in the unknown TPs (from C4 to C8). The combination of several spectroscopic techniques, such as further analysis by nuclear magnetic resonance (NMR), would be required to definitely elucidate the molecular structure of these compounds. Nevertheless, the information obtained in this study regarding the elemental composition of protonated TPs and their fragment ions will allow screening of these compounds in future monitoring studies. This is of interest to have more realistic and complete information, as these TPs are not included in environmental studies related with the presence of cocaine.

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318 **3.3. Photo-degradation**

Photo-degradation of cocaine and BE in aqueous solution under simulated sunlight and/or UV irradiation resulted in eight TPs (**Figure 1**) including two known metabolites: BE (TP-P4) and norBE (TP-P5). The TPs and the data obtained from the QTOF experiments are summarized in **Table 2**. Initially, TP-P2 isomers and TP-P4 were also generated after UV irradiation, but these TPs were effectively removed after 3 and 8 hours, respectively.

TP-P1 (C₉H₁₄NO₂⁺, m/z 168.1025), with an abundant peak at 1.93 min, was generated 324 under simulated sunlight of BE. TP-P1 may be produced via cleavage of the benzoyl ester 325 326 bond, reduction of the carboxylic acid group and dehydrogenation. Its fragmentation under 327 HE acquisition mode involved the loss of water $(m/z \ 150)$ and subsequent loss of 328 formaldehyde (m/z, 120) and of the bridgehead nitrogen (m/z, 93). The fact that there is no loss 329 of both H₂O and CO indicates the absence of a carboxylic acid function in this molecule. No 330 confirmative position of the double-bond could be given. However most likely it would be 331 located in the part away from the hydroxyl and aldehyde group.

332 After photo-degradation of both cocaine and BE, three isomeric products named as 333 TP-P2 ($C_{16}H_{20}NO_5^+$, m/z 306.1341) were detected at retention times of 2.78, 3.07 and 3.91 334 min (Figure 2A, left). They would correspond to a hydroxylation product of BE. During 335 photo-degradation of cocaine, BE was formed and readily transformed afterwards acting as a photo-intermediate (Figure 3). In MS^E, the three TP-P2 isomers showed similar 336 337 fragmentation with fragment ions m/z 168 (loss of 138 Da, i.e., benzoic acid+O) and m/z 121 338 $(C_7H_5O_2^+, \text{ corresponding to } C_6H_5CO^+ + O, m/z \text{ 105+16}), \text{ indicating that hydroxylation occurs}$ 339 at the phenyl ring. Accordingly, the TP-P2 isomers were presumably ortho-, meta- and para-340 hydroxy-BE. In vivo, cocaine metabolizes to BE and then to norBE and/or to meta-hydroxyl-341 BE and para-hydroxyl-BE (Klette et al. 2000). The suggested hydroxylation products here are 342 similar to the monohydroxylated cocaine products generated by hydrogen peroxide treatment 343 (Tanaka et al. 2002), as a result of solar photo-degradation using a catalyst (titanium dioxide, 344 TiO₂), or by a photo-Fenton reaction (Postigo et al. 2011). In our work, the three isomers seemed to be formed in the photo-degradation experiments as the XIC at the $[M+H]^+$ exact 345 346 mass (m/z 306.1341) revealed. All the three isomers gave fragment ions with m/z 168, 150 347 and 121, whereas only two compounds generated the ion with m/z 186 (Figure 2A). This is 348 probably due to the fact that the loss of 120 Da (C₇H₄O₂) results in resonance-stabilized

neutrals only for the *ortho-* and *para-*analogues (**Figure 2B**). Thus, the peak at 3.07 min should be the *meta-*hydroxy-BE. Combining this information with literature data on the elution order of *para-* and *meta-*hydroxy-BE (Pichini et al. 2005; Bisceglia et al. 2010), one can conclude that the isomer at 2.78 min is the *para-* isomer and the one with 3.90 min the *ortho-* isomer.

NorBE (named as TP-P5 in this section) co-eluted with one of the hydoxylated derivatives (Rt = 3.96 and 3.91, respectively) and as a consequence overlapping spectra were obtained in MS^E . In this case, additional product ion MS/MS experiments were performed to obtain "clean" accurate mass spectra of both compounds to confirm their identities.

358 The accurate mass of two other, less abundant but interesting, unknown isomeric TPs 359 with retention times 2.94 and 3.54 min was determined to be m/z 351.1180 (TP-P3) (Figure S2). The most likely molecular formula is $C_{16}H_{18}N_2O_7$ (mass error -1.2 mDa). Therefore, 360 361 TP-P3 is suggested to be generated via hydroxylation (+ OH) and nitration (+ NO₂) of BE. 362 The incorporation of a NO₂ group is feasible, since the photo-degradation experiments were 363 carried out using surface water of the Mijares River (Castellón province), where relatively 364 high nitrate concentrations (around 10 mg/L) are normally present owing to the wide use of 365 fertilizers in this agricultural area (Hernández et al. 2008). The presence of the common 366 fragment ions with m/z 168, 150, 119 and 82, indicated that hydroxylation and nitration did 367 not take place on the bicyclic ring system, but on the phenyl ring. This could be confirmed by the presence of a major fragment ion with m/z 166.0140 (C₇H₄NO₄⁺) corresponding to 368 369 $C_6H_5CO^+ + OH + NO_2 - H_2 (m/z \ 105 + 17 + 46 - 2)$. The positions of the NO₂ and OH group 370 could not be definitively determined. From a structural point of view, one might expect more 371 than two chromatographic peaks, since there are various possible combinations regarding the 372 positions of NO₂ and OH. Supposedly, hydroxylation takes place first, because a possible TP 373 corresponding to the nitration of BE ($C_{16}H_{20}NO_4^+ + NO_2$) with m/z 335 was not observed, 374 whereas hydroxyl-BE was in fact found, as previously discussed. Based on the effect of a 375 hydroxyl-group on electrophilic substitution, the entrance of NO₂ is probably ortho- and 376 para- orientated (Morrison and Boyd 1992). Together with the three possible hydroxy-BE 377 structures, this would result in six conceivable combinations (four ortho- and two para-378 orientated). Nevertheless, only two chromatographic peaks were observed at 2.94 and 3.54 379 min (Figure S2). Possibly, the small differences in polarity allowed co-elution of the *ortho*-380 and of the para- orientated isomers. Owing to interaction via intra-molecular H-bonding of 381 the neighbouring -NO₂ and -OH groups in the ortho- position the overall polarity of the 382 molecule decreases. As an example, o-nitrophenol is retained stronger than p-nitrophenol

using a reversed-phase analytical column (Masqué et al. 2000). The presence of nitrated
derivates indicates influence of the matrix on the degradation of the parent compound.

The use of UHPLC allowed decreasing analysis time with excellent chromatographic resolution. These characteristics are important in terms of sample throughput, separation efficiency and sensitivity (Wilson et al. 2005). In **Figure 2** and **Figure S2**, XICs of common fragments show several chromatographic peaks resulting from different TPs. The chromatographic separation of these TPs was important in order to avoid overlapping of spectra acquired using the MS^E approach and to facilitate a reliable identification. Furthermore, the inherent increased sensitivity favoured the detection of less abundant TPs.

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393 **3.4. Screening of water samples**

394 Screening of cocaine TPs has been performed in several sewage and surface water 395 samples, including the highest number of TPs reported until now. To this aim an UHPLC-396 MS/MS (QqQ) system was used for the screening of the above suggested TPs in the water 397 samples. This technique is especially suited for target screening in complex matrices as high 398 sensitivity can be achieved in selected reaction monitoring (SRM) mode. SPE was applied to 399 five influent, five effluent sewage waters and five surface waters in order to pre-concentrate 400 and clean-up the samples (see Section 2.5). Hydrophilic and lipophilic balanced (HLB) 401 cartridges were selected, which demonstrated good efficiency for cocaine, BE and other 402 drugs, pharmaceuticals and metabolites with different physical and chemical characteristics 403 (Gheorghe et al. 2008; Baker and Kasprzyk-Hordern 2011; Gracia-Lor et al. 2011). The 404 precursor and product ions, i.e. the MS/MS transitions acquired, were selected (Table 3) on 405 the basis of the main ions observed in previous QTOF MS analysis performed along the 406 degradation experiments. A more sensitive QqQ analyzer (i.e. TQS) was used for surface 407 water analysis, due to the lower concentrations expected in comparison with sewage water.

408 Besides the known metabolites (norcocaine, BE and norBE), TPs of cocaine and BE 409 have been detected, for the first time, in water samples. TP-P1 was found in one influent, four 410 effluents and four surface waters, and TP-P2 isomers were present in four influent and two 411 effluent sewage waters and four surface waters. The TPs had been elucidated after photo-412 degradation of cocaine and BE by simulated sunlight and/or UV irradiation. Therefore, their 413 presence in influents might be noticed as remarkable, as influent sewage water is normally not 414 exposed to sunlight or UV irradiation. As previously discussed, TP-P2 isomers are suggested 415 to be ortho-, meta- and para-hydroxy-BE. Meta-hydroxy-BE and para-hydroxy-BE have 416 been reported as in vivo metabolites and might therefore be present in influent sewage water

417 as a consequence of excretion. Nevertheless, ortho-hydroxy-BE and TP-P1 were also present 418 in influents. Thus, other processes (e.g. bacterial decomposition) in the sewage system might 419 occur and also play a role in their formation. The presence of TP-P1 and TP-P2 in effluent 420 could be caused by incomplete elimination in the STP or degradation of cocaine and BE 421 during treatment. Furthermore, these TPs were not only found for the first time in influent and 422 effluent sewage waters, but also in surface waters. TPs might enter surface waters by 423 releasing sewage effluents or be formed by photo-degradation via natural sunlight. Figure 4 424 shows a positive finding of TP-P1 and TP-P2 isomers in effluent sewage water and in surface 425 water. Although their reference standards were not available, the fact that all the three SRM 426 transitions acquired and that relative retention times to BE (RT-TP/RT-BE) were in good 427 agreement (< 0.01 min) with the TPs identified in degradation experiments give reliability to 428 these findings.

In future studies, additional degradation experiments should be performed in wastewaters in order to address the presence of some of the identified TPs in influents analyzed. Moreover, reference standards of the discovered TPs are required in order to report concentration levels. Subsequently, extended monitoring studies should be set up analyzing paired wastewater and receiving surface waters. This will give more insight in the environmental fate of cocaine, its metabolites and TPs.

436 4. CONCLUSIONS

437 Data on the presence of TPs of organic contaminants in the aquatic environment are 438 required nowadays to have a realistic overview of water quality. UHPLC-QTOF MS has been 439 demonstrated in this work as a valuable tool for the identification of TPs of cocaine and its 440 main metabolite BE in water. After laboratory-controlled hydrolysis, chlorination and photo-441 degradation experiments, the structures of several TPs have been tentatively established. The 442 applicability of these studies has been demonstrated by analyses of sewage water (influent 443 and effluent) and surface water where, in addition to well-known cocaine metabolites, other 444 TPs identified in laboratory experiments have been found and reported for the first time. The 445 relevance of TPs should not be neglected, as it might be necessary to take them into account 446 in future monitoring studies. Other knowledge gaps, such as ecotoxicity and the effects of 447 multiple compound exposure, need to be adressed in order to perform well-founded 448 environmental risk assessment. This implies that lot of research is still required by 449 environmental scientists and analytical chemists, especially when dealing with emerging 450 contaminants.

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460

461 SUPPLEMENTARY DATA

462 In this section, two figures, one reporting the identification of cocaine chlorination TP463 C8 (Figure S1) and the other including the identification of cocaine photo-degradation TP-P3

464 (Figure S2), are included to have supportive visual information on the written text.

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- 562

 (1		
Compound	Retention time	Accurate mass	Chemical	Mass Error	DBE
 (area)	(min)	(<i>m/z</i>)	formulae	(mDa)	
TP-C1 ^{s, s}	3.74	290.1400	$C_{16}H_{20}NO_4$	+0.8	7.5
BE		168.0979	$C_9H_{14}NO_2$	-4.6	3.5
(2059)		150.0896	C ₉ H ₁₂ NO	-2.3	4.5
		119.0478	C ₈ H ₇ O	-1.9	5.5
		105.0325	C7H5O	-1.5	5.5
		82.0658	C₅H₀N	+0.1	2.5
		02.0000	03.10.1		
TP-C2 ^b	3 95	276 1229		-0.7	75
norBE	0.00	154 0952		1.5	2.5
(COO)		104.0000		-1.5	3.5
(080)		136.0744		-1.8	4.5
		105.0332	C ₇ H ₅ O	-0.8	5.5
TP-C3 ^c	/ 31	200 1301		-0.1	75
	4.51	290.1391		-0.1	7.5
		100.0990		-2.1	3.5
(491)		136.0750	C ₈ H ₁₀ NO	-1.2	4.5
		105.0340	C7H₅O	0.0	5.5
	/ 81	220 0729		-11	25
(532)	4.01	202 0620		-0.6	2.5
(002)		188 0475		-0.0 ⊥0.3	3.5
		100.0475		+0.3	3.5
		120.0210		-0.0	1.5
		114.0103	C5H5CIN	-0.0	3.0
TP-C5 [⊳]	5.89	304.1193	C ₁₆ H ₁₈ NO ₅	+0.8	8.5
(18)		286.1080	C ₁₆ H ₁₆ NO ₄	+0.1	9.5
		182.0823	$C_9H_{12}NO_3$	+0.6	4.5
		154.0855		-1.3	3.5
		136.0740		-2.2	4.5
		105.0337	C ₇ H₅O	-0.3	5.5
TP-C6°	6.45	318.1336	$C_{17}H_{20}NO_5$	-0.5	8.5
(270)		286.1061	$C_{16}H_{16}NO_4$	-1.8	9.5
		196.0951	$C_{10}H_{14}NO_{3}$	-2.3	4.5
		168.1002	$C_9H_{14}NO_2$	-2.3	3.5
		136.0743	C ₈ H ₁₀ NO	-1.9	4.5
		105.0325	C7H₂O	-1.5	5.5
	7 50	040 0007			
IP-C7	7.53	310.0837	C ₁₅ H ₁₇ CINO ₄	-0.9	7.5
(607)		274.1046	$C_{15}H_{16}NO_4$	-3.3	8.5
		188.0493	C ₈ H ₁₁ CINO ₂	+1.5	3.5
		170.0352	C ₈ H ₉ CINO	-2.1	4.5
		152.0700	$C_8H_{10}NO_2$	-1.2	4.5
		142.0417	C7H9CIN	-0.7	3.5
		134.0592	C ₈ H ₈ NO	-1.4	5.5
		105.0329	C7H₅O	-1.1	5.5
TP-C8 ^c	8 27	324 0088		-15	75
(3765)	0.21	288 1216		-20	8.5
(3703)		200.1210		-2.0	0.0
		202.0598		-3.1	3.5
		170.0351	C ₈ H ₉ CINO	-2.2	4.5
		166.0838	$C_9H_{12}NO_2$	-3.0	4.5
		142.0406	C7H9CIN	-1.8	3.5
		134.0587	C ₈ H ₈ NO	-1.9	5.5
		105.0328	C7H₂O	-1.2	5.5

563 Table 1: Proposed elemental composition of protonated TPs and their fragments ions 564 obtained during chlorination of cocaine and BE, retention time (min), accurate mass (m/z), mass error (mDa) and double bond equivalent (DBE). 565

^a Maximum absolute area observed (initial area for parent compounds around 4700) 566

^b Also detected in negative ionization mode (-V) ^c TP only observed from cocaine 567

568

570**Table 2:** Proposed elemental composition of protonated TPs and their fragments ions571obtained during photo-degradation of cocaine and BE, retention time (min),572accurate mass (m/z), mass error (mDa) and the double bond equivalent (DBE).

Compound (area) ^a	Retention time (min)	Sun / UV ^d	Accurate mass (<i>m/z</i>)	Chemical formulae	Mass error (mDa)	DBE
TP-P1 [♭]	2.05	Sun	168.1009	$C_9H_{14}NO_2$	-1.6	3.5
(81)			150.0902	C ₉ H ₁₂ NO	-1.7	4.5
			120.0792	$C_8H_{10}N$	-2.1	4.5
			100.0747	C₅H11NO	-1.5	1.5
			93.0686	C_7H_9	-1.8	3.5
TP-P2 ^c	2.78	Sun /	306.1332	$C_{16}H_{20}NO_5$	-0.9	7.5
hydroxy-BE	3.07	UV	186.1103 ^e	$C_9H_{16}NO_3$	-2.7	2.5
(111)	3.91		168.1014	$C_9H_{14}NO_2$	-1.1	3.5
(46)			150.0903	C ₉ H ₁₂ NO	-1.6	4.5
(88)			121.0265	$C_7H_5O_2$	-2.5	5.5
			82.0665	C_5H_8N	+0.8	2.5
TP-P3	2.94	Sun	351.1180	$C_{16}H_{19}N_2O_7$	-1.2	8.5
(22)	3.54		168.1004	$C_9H_{14}NO_2$	-2.1	3.5
(39)			166.0118	C ₇ H ₄ NO ₄	-2.2	6.5
			150.0902	C ₉ H ₁₂ NO	-1.7	4.5
			119.0487	C ₈ H ₇ O	-1.0	5.5
			82.0660	C_5H_8N	+0.3	2.5
TP-P4 ^c	3.74	Sun /	290.1397	$C_{16}H_{20}NO_4$	+0.5	7.5
BE		UV	272.1304	$C_{16}H_{18}NO_{3}$	+1.7	8.5
(6189)			168.1047	$C_9H_{14}NO_2$	+2.2	3.5
			150.0946	C ₉ H ₁₂ NO	+2.7	4.5
			119.0514	C ₈ H ₇ O	+1.7	5.5
			105.0357	C7H₅O	+1.7	5.5
			82.0672	C_5H_8N	+1.5	2.5
TP-P5 ^c	3.96	Sun	276.1227	$C_{15}H_{18}NO_4$	-0.9	7.5
norBE			154.0850	$C_8H_{12}NO_2$	-1.8	3.5
(20)			136.0738	C ₈ H ₁₀ NO	-2.4	4.5
			108.0799	$C_7H_{10}N$	-1.4	3.5
			105.0335	C7H₅O	-0.5	5.5

^a Maximum absolute area observed (initial area for parent compounds around 7800)

574 ^b TP only observed from BE

^c Also detected in negative ionization mode (-V)

^d TP as a result of irradiation under simulated sunlight (Sun) and/or ultraviolet (UV)

^e This fragment ion is not present in the *meta*-hydroxyBE (Tr= 3.07min)

579 Table 3: UHPLC-MS/MS parameters established for the SRM acquisition mode	579	Table 3: UHPLC-MS/MS parameters established for the SRM acquisition mode.
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Compounds	Retention time	Precursor ion	CV ^a	CE ^b	Product ion
	(min)	$(m/z) [M + H]^+$	(V)	(eV)	(m/z)
TP-C1/TP-P4	3.74	290.1	40	20	168.2
(BE)				30	105.0
				30	82.0
TP-C2/TP-P5	3.95	276.2	45	15	154.1
(norBE)				20	136.1
				30	105.0
TP-C3	4.31	290.1	40	20	168.2
(norcocaine)				25	136.1
				30	105.0
TP-C4	4.81	220.1	35	30	188.0
				25	120.0
				30	114.0
TP-C5	5.89	304.1	30	25	154.1
				25	136.0
				30	105.3
TP-C6	6.45	318.1	30	25	286.1
				20	196.2
				25	136.1
TP-C7	7.53	310.1	30	25	188.0
				25	152.0
				35	105.0
TP-C8	8.27	324.1	35	25	288.1
				25	166.1
				25	105.0
TP-P1	2.05	168.1	35	30	150.1
				25	120.1
				30	93.0
TP-P2	2.78	306.1	35	20	168.1
	3.07			15	186.1
	3.91			30	121.0
TP-P3	2.94	351.1	35	35	119.0
	3.54			15	168.1
				35	82.1

581 ^aCV, cone voltage; ^bCE, collision energy

587

586 **FIGURE CAPTIONS**

- 588 Figure. 1 Structure proposals for the identified TPs (TP-Cx were observed after chlorination
 589 and TP-Px were observed after photo-degradation).
- ^a TP only observed from photo-degradation of BE.
- 591 ^b TP detected in sewage and environmental waters

592

- Figure. 2 Detection and identification of cocaine and BE photo-degradation TP-P2 by
 UHPLC-QTOF MS operating under MS^E. (A) narrow-window XICs (± 10 mDa)
 of TP-P2 and structures suggested for [M+H]⁺ of TP-P2 and its fragment ions. (B)
 Structure proposals for the neutral loss of 120 Da (C₇H₄O₂).
 Notice the presence of other chromatographic peaks (marked with *), supporting
 that other TPs share the same fragmentation.
- Figure. 3 Photo-degradation of cocaine where BE acts as photo-intermediate in the formationof TP-P2.
- 602
- Figure. 4 UHPLC-MS/MS chromatograms corresponding to the positive finding of TP-P1
 and TP-P2 in (top) effluent sewage water (analyzed using TQD, December 2011)
 and (bottom) surface water (analyzed using TQS, April 2012). Retention times of
 BE were 3.75 min (TQD) and 3.84 (TQS).





Figure 2









