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

3

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13

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
23 compounds were found (3 from chlorination; 7 from photo-degradation), including one
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
27 nitration, or from a combination of these processes. Several influent and effluent sewage
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.

36

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;
50 Gheorghe et al. 2008)), there is a potential negative impact of their presence in the aquatic
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
82 applied by STPs, *e.g.* chlorination and UV irradiation, can be simulated, as well as natural
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
86 al. 2008; Quintana et al. 2010; Wick et al. 2011), and illustrates the importance of
87 investigating TPs.

88 The use of MS^E is an attractive option, which is feasible working with hybrid QTOF
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.
105

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.

114 HPLC-grade MeOH, ACN and formic acid (FA) were acquired from Scharlau
115 (Barcelona, Spain). Sodium hypochlorite solution (available chlorine 10%) was obtained from
116 Sigma-Aldrich. A Milli-Q ultra-pure water system from Millipore (Bedford, MA, USA) was
117 used to obtain the HPLC grade water. Leucine enkephalin and imazalil were purchased from
118 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.

122

123 2.2. Degradation experiments

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

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

133 The hydrolysis and chlorination experiments were performed at room temperature and
134 in darkness. Regarding chlorination, 40 μ L of ten-fold diluted sodium hypochlorite solution
135 was added to 50 mL of each surface water sample. During the experiment, 2 mL aliquots of
136 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 $-20\text{ }^{\circ}\text{C}$ 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 -
146 800 nm. The radiation intensity was set to 500 W/m^2 and the light dose per hour of irradiation
147 to 1.8 MJ/h . In this way, 90 irradiation hours corresponds to 15 days of natural sun light
148 (dose: 288 MJ/m^2). The degradation was performed using 250 mL closed quartz glass vessels
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 $-20\text{ }^{\circ}\text{C}$ until analysis.

154

155 **2.3. Instrumentation**

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

163 The UHPLC separation was performed using an Acquity UPLC BEH C18, $1.7\text{ }\mu\text{m}$
164 particle size analytical column, $100\text{ mm} \times 2.1\text{ mm}$ (Waters). The mobile phases used were A
165 = H_2O 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 .

168 For MS^E experiments, two acquisition functions were created and simultaneously used
169 within the same run: the low-energy function (LE) with a collision energy of 4 eV, where
170 mainly the (de) protonated intact molecules are observed, and the high energy (HE) function

171 with a collision energy ramp ranging from 15 to 40 eV, where fragmentation is promoted. The
172 same collision energy ramp was used for additional MS/MS experiments. The optimized cone
173 voltage (15 V) and collision energy ramp were identical for both cocaine and BE and seemed
174 therefore most adequate for the screening of their corresponding degradation products.
175 Further details on instrument operating conditions can be found elsewhere (Hernández et al.
176 2011).

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

182

183 **2.4. Elucidation / identification procedure**

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

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

194 The most likely elemental compositions of (de)protonated molecules were calculated
195 based on accurate mass LE spectra of the peaks of interest. The accurate mass HE spectra
196 were then used to calculate possible elemental compositions of fragment ions. Assuming that
197 most TPs share similar fragmentation pathways with the parent drug (Wang and Bartlett
198 1998; Bijlsma et al. 2011), fragmentation was compared to that of cocaine and BE, and the TP
199 structures were proposed.

200

201 **2.5. Water samples**

202 Five influent and five effluent sewage water samples (24-hour composite) and five
203 surface water grab samples from different locations of the Comunidad Valenciana (Eastern
204 Spain) were collected and immediately stored at -20 °C. Sewage water was collected from

205 STPs of Castellón and Benicàssim, while surface water was collected from the Albufera
206 national park of Valencia.

207 100 mL of five-fold diluted (with MilliQ) influent wastewater, 100 mL of effluent
208 wastewater or 100 mL surface water was taken for analysis. The samples were loaded onto
209 the HLB cartridges by gravity, and then cartridges were vacuum-dried for 10 min. Analytes
210 were eluted with 5 mL of MeOH. The extracts were evaporated to dryness at 35°C under a
211 gentle stream of nitrogen and reconstructed in 1 mL of 10:90 MeOH:H₂O. Analyses of
212 cocaine and BE TPs were performed by injecting 20 µL of the final extract into the UHPLC-
213 TQD system (sewage water) or 100 µL in the UHPLC-TQS system (surface water).

214

215 3. RESULTS AND DISCUSSION

216

217 Many known TPs of environmental contaminants share similar fragmentation
218 pathways as their parent molecules. Then, knowledge of structures of fragment ions and basic
219 fragmentation rules are helpful for achieving confident TP structure proposals. Isotope fit,
220 Double Bound Equivalent (DBE), and accurate mass of fragments observed in the HE
221 function were used to discard potential chemical formulas in order to obtain the most
222 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 ($C_6H_5CO^+$) 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

233 3.1. Hydrolysis

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

241 Complete cocaine and some BE degradation was observed in surface water after
242 keeping the solution in darkness at room temperature for 15 days (data not shown). Cocaine
243 was mainly transformed into BE through chemical hydrolysis of cocaine ester bonds.
244 Ecgonine methyl ester (EME), another hydrolytic product reported for cocaine (Postigo et al.
245 2011), was not observed. EME is presumed to be solely an *in vivo* metabolite as a result of
246 enzymatic hydrolysis and for that reason it is unlikely to be formed during cocaine
247 degradation in water (Klette et al. 2000). Gheorghe et al. (2008) had similar results to the

248 present work, where cocaine and EME degraded in spiked surface water, while BE initially
249 increased owing to the possible chemical hydrolysis of cocaine.

250

251 3.2. Chlorination

252 **Table 1** summarizes the TPs of cocaine and BE formed during chlorination. Retention
253 times and experimental m/z -values, proposed elemental composition of the protonated TPs
254 and their fragment ions, the mass error in mDa, and the double bond equivalent (DBE) are
255 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
262 were no longer observed after adding ascorbic acid to the sample vials). Therefore, we did not
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
268 pattern information (distribution of the ^{37}Cl isotope) obtained in the MS^E mode, allowed the
269 detection and tentative identification of several TPs, in a single injection. Among these TPs,
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 $[\text{M}+\text{H}]^+$, their sodium adducts $[\text{M}+\text{Na}]^+$ 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 ($\text{C}_9\text{H}_{15}\text{ClNO}_3^+$, 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 ($C_{16}H_{19}ClNO_4^+$), 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 ($C_6H_5CO^+$) 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 CH_3OH 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.

292 Another TP of cocaine, TP-C6 ($[M+H]^+$, m/z 318.1336) is has the same nominal mass
293 as cocaethylene ($C_{18}H_{24}NO_4^+$, m/z 318.1705), but they could be differentiated both
294 chromatographically and by HRMS, as a difference of 36.9 mDa was observed. The most
295 likely molecular formula for TP-C6 is $C_{17}H_{20}NO_5^+$ (m/z 318.1341, Δ 0.5 mDa). Thus, TP-C6
296 would result from oxidation (+O-2H) of cocaine during chlorination experiments, which
297 probably occurs on the bicyclic ring system, since the characteristic fragment ion with m/z
298 105 ($C_6H_5CO^+$) is still present.

299 Chlorination of BE resulted in TP-C5 ($[M+H]^+$, m/z 304.1185) and TP-C7
300 ($C_{15}H_{17}ClNO_4^+$, 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.

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

309 The data obtained was not sufficient to predict the exact position of the chlorine or
310 keto group in the unknown TPs (from C4 to C8). The combination of several spectroscopic
311 techniques, such as further analysis by nuclear magnetic resonance (NMR), would be required
312 to definitely elucidate the molecular structure of these compounds. Nevertheless, the
313 information obtained in this study regarding the elemental composition of protonated TPs and
314 their fragment ions will allow screening of these compounds in future monitoring studies.

315 This is of interest to have more realistic and complete information, as these TPs are not
316 included in environmental studies related with the presence of cocaine.

317

318 3.3. Photo-degradation

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

324 TP-P1 ($C_9H_{14}NO_2^+$, m/z 168.1025), with an abundant peak at 1.93 min, was generated
325 under simulated sunlight of BE. TP-P1 may be produced via cleavage of the benzoyl ester
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 H_2O 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
336 photo-intermediate (**Figure 3**). In MS^E , the three TP-P2 isomers showed similar
337 fragmentation with fragment ions m/z 168 (loss of 138 Da, i.e., benzoic acid+O) and m/z 121
338 ($C_7H_5O_2^+$, corresponding to $C_6H_5CO^+ + O$, m/z 105+16), 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 TiO_2), or by a photo-Fenton reaction (Postigo et al. 2011). In our work, the three isomers
345 seemed to be formed in the photo-degradation experiments as the XIC at the $[M+H]^+$ exact
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 ($C_7H_4O_2$) results in resonance-stabilized

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

354 NorBE (named as TP-P5 in this section) co-eluted with one of the hydroxylated
355 derivatives ($R_t = 3.96$ and 3.91 , respectively) and as a consequence overlapping spectra were
356 obtained in MS^E. In this case, additional product ion MS/MS experiments were performed to
357 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**
360 **S2**). The most likely molecular formula is $C_{16}H_{18}N_2O_7$ (mass error -1.2 mDa). Therefore,
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
368 the presence of a major fragment ion with m/z 166.0140 ($C_7H_4NO_4^+$) corresponding to
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

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

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

392

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.

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

435

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 addressed 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.

451

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460

461 **SUPPLEMENTARY DATA**

462 In this section, two figures, one reporting the identification of cocaine chlorination TP-
463 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.

465

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562

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
 565 mass (m/z), mass error (mDa) and double bond equivalent (DBE).

Compound (area) ^a	Retention time (min)	Accurate mass (m/z)	Chemical formulae	Mass Error (mDa)	DBE
TP-C1 ^{b, c} BE (2059)	3.74	290.1400	C ₁₆ H ₂₀ NO ₄	+0.8	7.5
		168.0979	C ₉ H ₁₄ NO ₂	-4.6	3.5
		150.0896	C ₉ H ₁₂ NO	-2.3	4.5
		119.0478	C ₈ H ₇ O	-1.9	5.5
		105.0325	C ₇ H ₅ O	-1.5	5.5
		82.0658	C ₅ H ₈ N	+0.1	2.5
TP-C2 ^b norBE (680)	3.95	276.1229	C ₁₅ H ₁₈ NO ₄	-0.7	7.5
		154.0853	C ₈ H ₁₂ NO ₂	-1.5	3.5
		136.0744	C ₈ H ₁₀ NO	-1.8	4.5
		105.0332	C ₇ H ₅ O	-0.8	5.5
TP-C3 ^c norcocaine (491)	4.31	290.1391	C ₁₆ H ₂₀ NO ₄	-0.1	7.5
		168.0998	C ₉ H ₁₄ NO ₂	-2.7	3.5
		136.0750	C ₈ H ₁₀ NO	-1.2	4.5
		105.0340	C ₇ H ₅ O	0.0	5.5
TP-C4 ^c (532)	4.81	220.0729	C ₉ H ₁₅ CINO ₃	-1.1	2.5
		202.0629	C ₉ H ₁₃ CINO ₂	-0.6	3.5
		188.0475	C ₈ H ₁₁ CINO ₂	+0.3	3.5
		120.0210	C ₄ H ₇ CINO	-0.6	1.5
		114.0103	C ₅ H ₅ CIN	-0.8	3.5
TP-C5 ^b (18)	5.89	304.1193	C ₁₆ H ₁₈ NO ₅	+0.8	8.5
		286.1080	C ₁₆ H ₁₆ NO ₄	+0.1	9.5
		182.0823	C ₉ H ₁₂ NO ₃	+0.6	4.5
		154.0855	C ₈ H ₁₂ NO ₂	-1.3	3.5
		136.0740	C ₈ H ₁₀ NO	-2.2	4.5
		105.0337	C ₇ H ₅ O	-0.3	5.5
TP-C6 ^c (270)	6.45	318.1336	C ₁₇ H ₂₀ NO ₅	-0.5	8.5
		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 ₉ H ₁₄ NO ₂	-2.3	3.5
		136.0743	C ₈ H ₁₀ NO	-1.9	4.5
		105.0325	C ₇ H ₅ O	-1.5	5.5
TP-C7 ^b (607)	7.53	310.0837	C ₁₅ H ₁₇ CINO ₄	-0.9	7.5
		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 ₈ H ₁₀ NO ₂	-1.2	4.5
		142.0417	C ₇ H ₉ CIN	-0.7	3.5
		134.0592	C ₈ H ₈ NO	-1.4	5.5
		105.0329	C ₇ H ₅ O	-1.1	5.5
TP-C8 ^c (3765)	8.27	324.0988	C ₁₆ H ₁₉ CINO ₄	-1.5	7.5
		288.1216	C ₁₆ H ₁₈ NO ₄	-2.0	8.5
		202.0598	C ₉ H ₁₃ CINO ₂	-3.7	3.5
		170.0351	C ₈ H ₉ CINO	-2.2	4.5
		166.0838	C ₉ H ₁₂ NO ₂	-3.0	4.5
		142.0406	C ₇ H ₉ CIN	-1.8	3.5
		134.0587	C ₈ H ₈ NO	-1.9	5.5
		105.0328	C ₇ H ₅ O	-1.2	5.5

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

567 ^b Also detected in negative ionization mode (-V)

568 ^c TP only observed from cocaine

569

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

Compound (area) ^a	Retention time (min)	Sun / UV ^d	Accurate mass (m/z)	Chemical formulae	Mass error (mDa)	DBE
TP-P1 ^b (81)	2.05	Sun	168.1009	C ₉ H ₁₄ NO ₂	-1.6	3.5
			150.0902	C ₉ H ₁₂ NO	-1.7	4.5
			120.0792	C ₈ H ₁₀ N	-2.1	4.5
			100.0747	C ₅ H ₁₁ NO	-1.5	1.5
			93.0686	C ₇ H ₉	-1.8	3.5
TP-P2 ^c hydroxy-BE (111) (46) (88)	2.78 3.07 3.91	Sun / UV	306.1332	C ₁₆ H ₂₀ NO ₅	-0.9	7.5
			186.1103 ^e	C ₉ H ₁₆ NO ₃	-2.7	2.5
			168.1014	C ₉ H ₁₄ NO ₂	-1.1	3.5
			150.0903	C ₉ H ₁₂ NO	-1.6	4.5
			121.0265	C ₇ H ₅ O ₂	-2.5	5.5
	82.0665	C ₅ H ₈ N	+0.8	2.5		
TP-P3 (22) (39)	2.94 3.54	Sun	351.1180	C ₁₆ H ₁₉ N ₂ O ₇	-1.2	8.5
			168.1004	C ₉ H ₁₄ NO ₂	-2.1	3.5
			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 ₅ H ₈ N	+0.3	2.5		
TP-P4 ^c BE (6189)	3.74	Sun / UV	290.1397	C ₁₆ H ₂₀ NO ₄	+0.5	7.5
			272.1304	C ₁₆ H ₁₈ NO ₃	+1.7	8.5
			168.1047	C ₉ H 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	C ₇ H ₅ O	+1.7	5.5		
	82.0672	C ₅ H ₈ N	+1.5	2.5		
TP-P5 ^c norBE (20)	3.96	Sun	276.1227	C ₁₅ H ₁₈ NO ₄	-0.9	7.5
			154.0850	C ₈ H ₁₂ NO ₂	-1.8	3.5
			136.0738	C ₈ H ₁₀ NO	-2.4	4.5
			108.0799	C ₇ H ₁₀ N	-1.4	3.5
			105.0335	C ₇ H ₅ O	-0.5	5.5

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

574 ^b TP only observed from BE

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

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

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

578

579 **Table 3:** UHPLC-MS/MS parameters established for the SRM acquisition mode.

580

Compounds	Retention time (min)	Precursor ion (<i>m/z</i>) [M + H] ⁺	CV ^a (V)	CE ^b (eV)	Product ion (<i>m/z</i>)
TP-C1/TP-P4 (BE)	3.74	290.1	40	20	168.2
				30	105.0
				30	82.0
TP-C2/TP-P5 (norBE)	3.95	276.2	45	15	154.1
				20	136.1
				30	105.0
TP-C3 (norcocaine)	4.31	290.1	40	20	168.2
				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

582

583

584

585

586 **FIGURE CAPTIONS**

587

588 **Figure. 1** Structure proposals for the identified TPs (TP-Cx were observed after chlorination
589 and TP-Px were observed after photo-degradation).

590 ^a TP only observed from photo-degradation of BE.

591 ^b TP detected in sewage and environmental waters

592

593 **Figure. 2** Detection and identification of cocaine and BE photo-degradation TP-P2 by
594 UHPLC-QTOF MS operating under MS^E. (A) narrow-window XICs (± 10 mDa)
595 of TP-P2 and structures suggested for $[M+H]^+$ of TP-P2 and its fragment ions. (B)
596 Structure proposals for the neutral loss of 120 Da (C₇H₄O₂).

597 Notice the presence of other chromatographic peaks (marked with *), supporting
598 that other TPs share the same fragmentation.

599

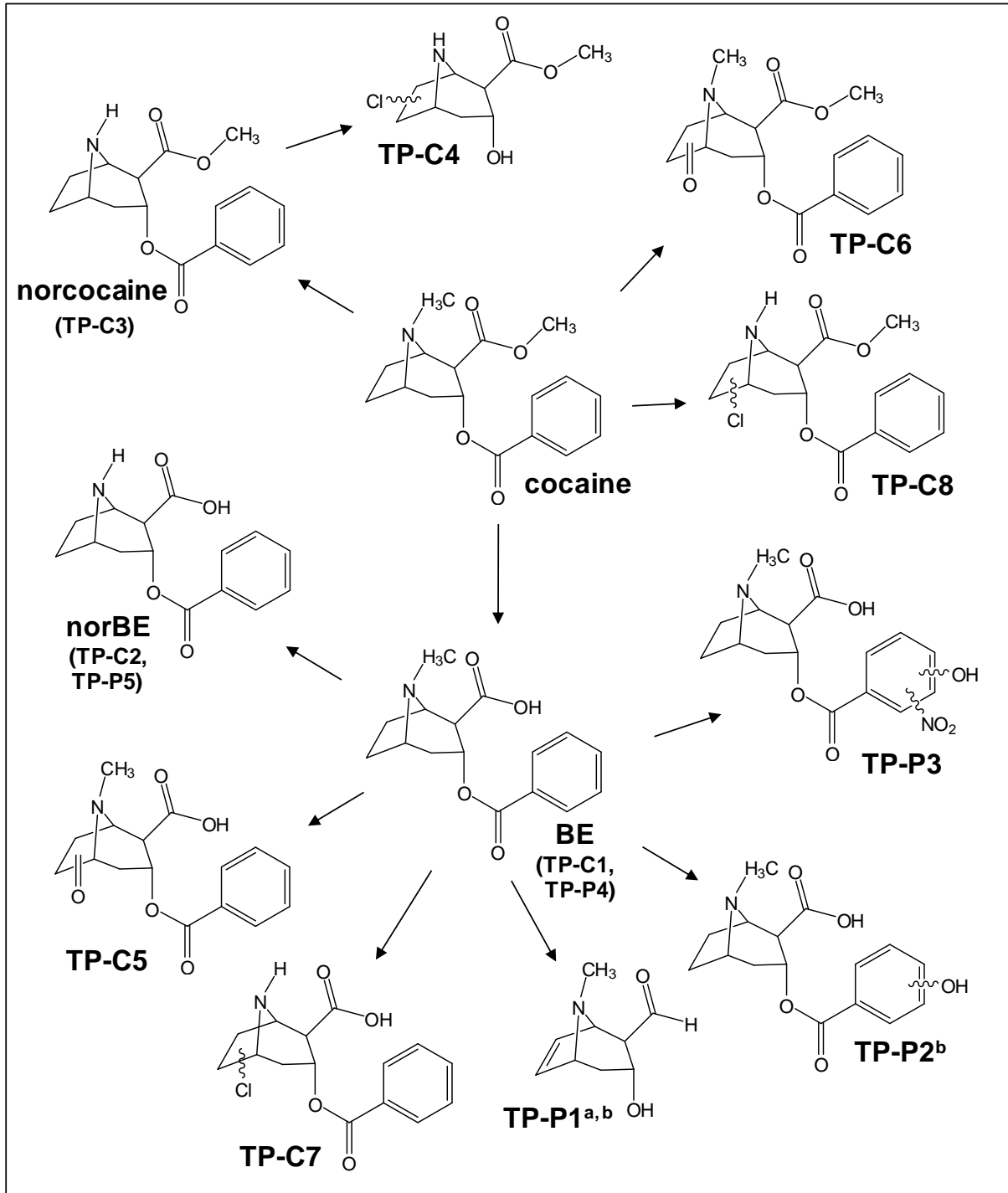
600 **Figure. 3** Photo-degradation of cocaine where BE acts as photo-intermediate in the formation
601 of TP-P2.

602

603 **Figure. 4** UHPLC-MS/MS chromatograms corresponding to the positive finding of TP-P1
604 and TP-P2 in (top) effluent sewage water (analyzed using TQD, December 2011)
605 and (bottom) surface water (analyzed using TQS, April 2012). Retention times of
606 BE were 3.75 min (TQD) and 3.84 (TQS).

607

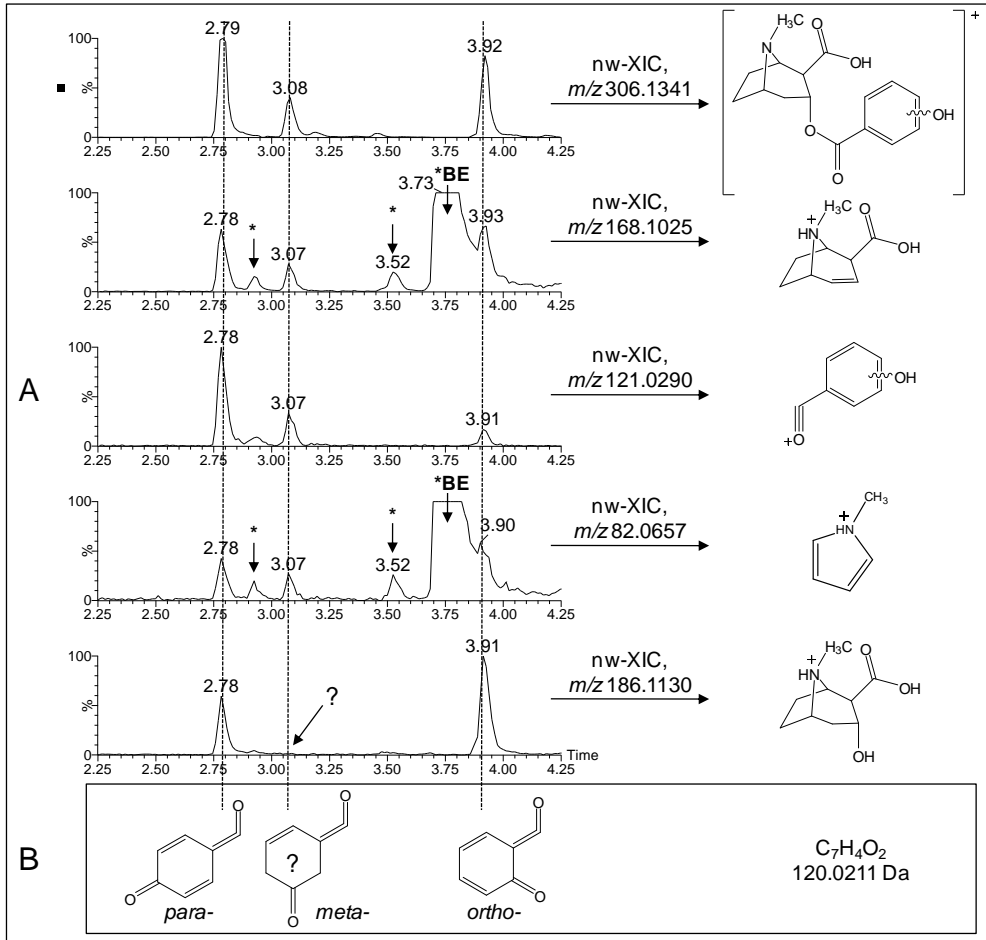
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Figure 1

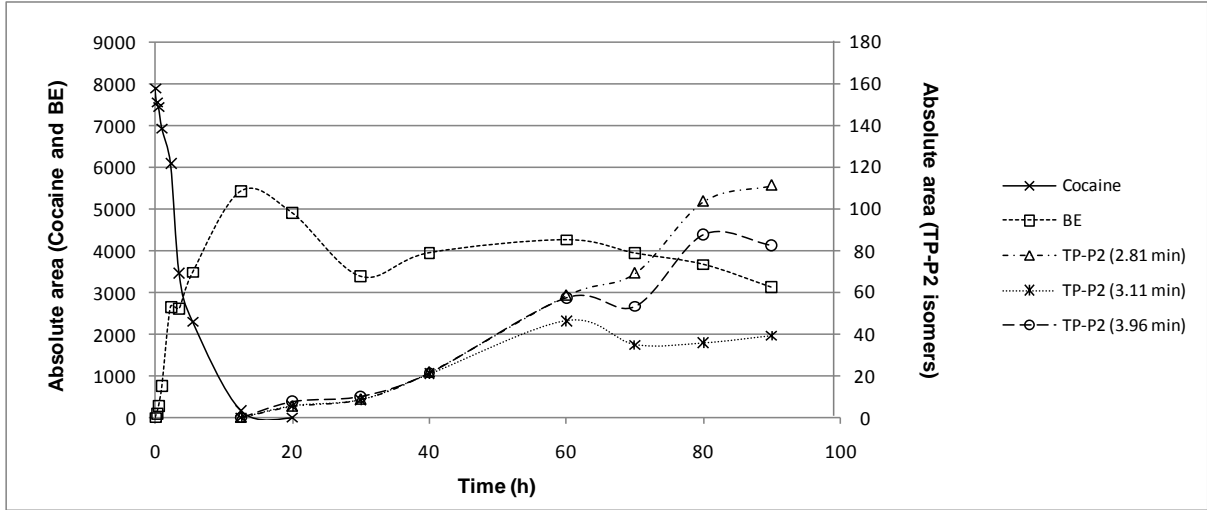
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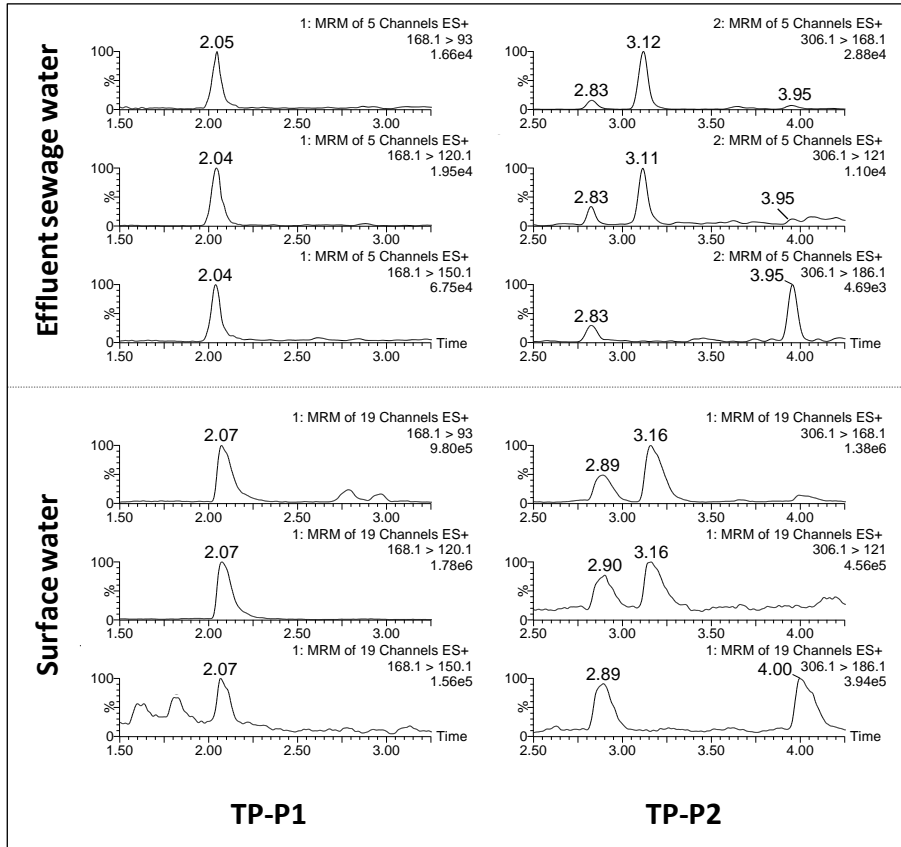
Figure 2

620



621
622
623
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Figure 3



626
627
628

Figure 4