Degradation kinetics and pathways of three calcium channel blockers under UV irradiation

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1. Introduction

In the last few decades, the occurrence of pharmaceuticals and personal care products (PPCPs) in the environment has received increasing attention and induces great challenge to the safety of aquatic ecosystems (Schwarzenbach et al., 2006). These PPCPs are mainly discharged from private households, hospitals and pharmaceutical industries, and eventually reach wastewater treatment plants (WWTPs). However, most of these compounds are only partly removed during wastewater treatment and thus appear in receiving waters and bio-solids (Batt et al., 2008; Chari and Halden, 2011). The occurrence of these CCBs in the WWTP effluents and downstream river waters have been frequently reported in recent years, with a concentration range of 3.0–510.0 ng L−1 (Hummel et al., 2006; Batt et al., 2008; Spongberg and Witter, 2008). Although no direct harmful effects to human have been reported, it is proved that an exposure to a certain concentration of CCBs in the WWTP effluent can induce ecological hazards to fish (Du et al., 2014). Wang and Gardinali (2013) also reported that DIL in reclaimed water can be ingested by fish and accumulate in fish body with a bio-concentration factor larger than 16 and an in-vivo half-life larger than 117 h, indicating a potential ecological risk.

Calcium channel blockers (CCBs), such as amlodipine (AML), diltiazem (DIL), and verapamil (VER), are a group of pharmaceuticals that can selectively block the influx of calcium ions through a calcium channel. They are widely prescribed to lower the blood pressure and to treat arrythmias, angina and other heart diseases (Elliott and Ram, 2011). The occurrence of these CCBs in the WWTP effluents and downstream river waters has been frequently reported in recent years, with a concentration range of 3.0–510.0 ng L−1 (Hummel et al., 2006; Batt et al., 2008; Spongberg and Witter, 2008). Although no direct harmful effects to human have been reported, it is proved that an exposure to a certain concentration of CCBs in the WWTP effluent can induce ecological hazards to fish (Du et al., 2014). Wang and Gardinali (2013) also reported that DIL in reclaimed water can be ingested by fish and accumulate in fish body with a bio-concentration factor larger than 16 and an in-vivo half-life larger than 117 h, indicating a potential ecological risk.
Because of its distinct advantages such as high inactivation efficiency, easy operation, and small space-occupancy, ultraviolet (UV) technologies have been increasingly applied for drinking water and wastewater disinfection (Parkinson et al., 2001; Li and Blatchley, 2009). Recently, UV photolysis of water-borne PPCPs during the WWTP effluent disinfection process has been more frequently reported (Sun et al., 2014; Wang et al., 2014); however, few studies have focused on the degradation kinetics and mechanism of CCBs under UV irradiation. As revealed by previous researches, most CCBs (e.g., ALP, DIL) are unstable against UVA, UVB (290–320 nm), or solar irradiation (Fasani et al., 2008; Kawabe et al., 2008), which indicates that the photo-degradation of CCBs under 254 nm UV is possible.

Moreover, co-existing matrix materials, such as dissolved organic matter (DOM) and NO\textsubscript{3}, may also affect the degradation kinetics of micro-pollutants under UV irradiation by altering UV light penetration and producing/scavenging hydroxyl radicals (\textOH). Mack and Bolton (1999) suggested that in the presence of NO\textsubscript{3} and NO\textsubscript{2} in aqueous solution, UV irradiation could yield \textOH with a quantum yield (\textphi_{\textOH}) of up to 9%. Wols and Hofman-Caris (2012) reported that DOM could not only act as important scavengers of \textOH with rate constants of about (2–8) \times 10^8 M\textsuperscript{–1} s\textsuperscript{–1} in surface water and wastewater, but also produce other radicals to indirectly degrade organic compounds. They also summarized the photochemical constants of over 100 organic compounds; however, none of the selected CCBs was included.

In the present study, three CCBs, including AML, DIL, and VER, were selected on behalf of the 1,4-dihydropyridine, 1,5-benzoazepine and phenylalkylamine classes, respectively. The degradation kinetics was investigated and the transformation byproducts (TBPs) were identified after exposing a relatively high concentration level of studied CCBs (i.e., 1.0 mg L\textsuperscript{–1}) to UV irradiation in Milli-Q water (MQ water). Then, the existence of functional pharmacophores in the TBP structures was examined and the degradation pathways were proposed accordingly. On that basis, a generic solid-phase extraction (SPE) method, which simultaneously used four different cartridges with different extraction modes, was adopted to extract and enrich TBPs formed in the photolytic experiments of target CCBs at a relatively low concentration level (i.e., 100 ng L\textsuperscript{–1}) under UV fluences typical for WWTP effluent disinfection. Moreover, the effects of co-existing matrix materials, including humic acid (HA), NO\textsubscript{3} and Cl\textsuperscript{–}, on the photo-degradation of AML were particularly clarified.

2. Materials and methods

2.1. Chemicals

The studied CCBs (>99% purity) including (+/-) verapamil hydrochloride (CAS No. 152-11-4), amlodipine besylate (CAS No. 111470-99-6) and (+)-cis-diltiazem hydrochloride (CAS No. 33286-22-5) were purchased from Sigma Aldrich (Schnelldorf, Germany), whose major physicochemical properties are shown in Table 1. Internal standards (amlodipine-d\textsubscript{4} maleic acid and verapamil-d\textsubscript{6} hydrochloride salt) were purchased from Toronto Research Chemicals (Ontario, Canada). Methanol and acetonitrile (ACN) were purchased from Fisher Scientific (Geel, Belgium), and ethyl acetate from Merck (Darmstadt, Germany). Formic acid (98%~100%, ACS grade) and HA sodium salt (CAS No. 68131-04-4, technical grade) were obtained from Sigma Aldrich, and KNO\textsubscript{3} and NaCl were from Panreac (Barcelona, Spain).

2.2. UV reaction system

Photo-degradation experiments were conducted in batch mode with a 2.0 L cylindrical glass reactor. The reaction solution was magnetically stirred and maintained at a constant temperature of 25 °C using a water bath (Fig. S1). A low-pressure mercury lamp (4 W, 30% UVC efficiency, Philips TUV G4T5) was used as the light source, which was initially warmed up for about 10 min to ensure a relatively stable output prior to each experiment. Afrazine actinometry was applied to measure the average UV fluence rate in the reactor (Canonica et al., 2008), which was equal to 0.77 mW cm\textsuperscript{–2} for MQ water and 0.52 mW cm\textsuperscript{–2} for the WWTP effluent (Text S1).

2.3. Experimental procedures and sample pretreatment

Photolytic experiments were conducted at two concentration levels of the target CCBs. Each experiment was run in triplicate and all glass containers were covered with aluminum foils to avoid

### Table 1

| Compound | Structure | Chemical formula | Molecular weight (g mol\textsuperscript{–1}) | Water solubility\textsuperscript{a} (mg L\textsuperscript{–1}) | pK\textsubscript{a} | Log K\textsubscript{ow} | Chemical properties of target CCBs. |
|----------|----------|-----------------|---------------------------------|-------------------|-----|----------------|----------------------------------|
| Amlodipine (AML) | ![AML structure](image1) | C\textsubscript{20}H\textsubscript{22}O\textsubscript{5}N\textsubscript{2}Cl | 408.1452 | 7.4 | 9.45 | 2.22 |
| Diltiazem (DIL) | ![DIL structure](image2) | C\textsubscript{22}H\textsubscript{26}O\textsubscript{4}N\textsubscript{2}S | 414.5178 | 16.8 | 8.18, 12.86 | 3.09 |
| Verapamil (VER) | ![VER structure](image3) | C\textsubscript{22}H\textsubscript{18}O\textsubscript{4}N\textsubscript{2} | 454.6016 | 3.94 | 9.68 | 3.79 |

\textsuperscript{a} Calculated by the ALOGPS 2.1 software (http://www.vcclab.org).
undesired photolysis. For the relatively high concentration level experiments, 1.0 mg L⁻¹ AML, DIL, and VER aqueous solutions (prepared in MQ water) were irradiated individually to determine their degradation kinetics and identify their TBPs. The solution pH was controlled at 7.0 with 10 mM phosphate buffer, so all the three CCBs were mainly present in the cationic form. At pre-selected time intervals, 1 mL aliquots of the irradiated sample were withdrawn and stored in 2 mL vials at −20 °C for later analysis.

For the relatively low concentration level experiments, the effluent collected from a local WWTP in Barcelona was used as matrix after being filtered through 0.7 μm glass fiber filters, whose major quality parameters are given in Table S1. The studied CCBs were spiked together into the reaction solution with an initial concentration of 100 ng L⁻¹ (each). At pre-selected time intervals, 400 mL aliquots of the irradiated sample were withdrawn, spiked with the internal standards (100 ng L⁻¹ each), and extracted by the generic SPE method as described elsewhere (Zonja et al., 2015). In brief, samples were adjusted to pH 6.5 ± 0.2 and extracted simultaneously with four different cartridges: Oasis HLB (500 mg/6 cc, Waters, Milford, USA), Bond Elut PPL (500 mg/6 cc, Agilent, Waghaeusel–Wiesental, Germany), Oasis MAX and Oasis MCX (both 200 mg/6 cc, Waters). The HLB cartridge was selected for its broad capacity of enriching multiple classes of organic compounds, while the PPL, MAX and MCX cartridges were selected for their speciality of enriching weak-polar, acidic and alkaline organic compounds, respectively. The four cartridges were connected in series, and conditioned sequentially with 5 mL of methanol/ethyl acetate (1:1, v/v) and 5 mL of MQ water. After an irradiated sample (400 mL) was extracted, the HLB and PPL cartridges were eluted with 3 × 3 mL of methanol/ethyl acetate (1:1), and the MAX and MCX cartridges were eluted sequentially with 3 mL of methanol/ethyl acetate (1:1) and 2 × 3 mL of methanol with 2% formic acid (for MAX) or methanol with 5% ammonia (for MCX). The resulting four extracts were combined together, evaporated to dryness under a gentle stream of nitrogen gas, and reconstituted with 400 μL of ACN/water (10:90) for later analysis. The generic SPE method could achieve acceptable recovery efficiencies for the target CCBs and identified TBPs (54.3%–119.9%, Table S2).

2.4. Analytical methods

The parent CCBs and their TBPs formed in photolysis experiments were detected using an Acquity ultra performance liquid chromatograph (UPLC, Waters) coupled to a Q-Exactive mass spectrometer (Thermo Scientific, Bremen, Germany), with an injection volume of 10 μL. Chromatographic separation was performed with an Acquity C18 column (100 × 2.1 mm, 1.7 μm) that was guarded by a pre-column of the same packing material (5 × 2.1 mm, 1.7 μm). Elution gradient with two mobile phases, including ACN with 0.1% formic acid (A) and MQ water with 0.1% formic acid (B), were programmed as follows (min, %A): (0, 10), (1, 10), (5, 90), (6, 90), (6.5, 10), and (8, 10). The flow rate was 0.3 mL min⁻¹ and the column temperature was held constant at 40 °C.

Electrospray ionization interface was operated in the positive ion mode with spray voltage of +3.0 kV, heater temperature of 250 °C, and capillary temperature of 350 °C. Data-dependent scan mode was applied for identification of TBPs with the following operation parameters: resolution of both full and MS² scans = 35000 (full width at half maximum), scan range = 50–600 m/z, isolation window = 2 m/z, and collision energy = 35 eV. The MS² scan could confirm the structures of the parent compounds and TBPs.

3. Results and discussion

3.1. Degradation kinetics of target CCBs in MQ water

The UV degradation of three CCBs, with an initial concentration of 1.0 mg L⁻¹, was conducted individually in MQ water at pH 7.0. All experiments lasted for 60 min, which corresponds to a UV fluence of 2772 mJ cm⁻² (UV fluence rate = 0.77 mW cm⁻², Text S1). The photolysis rate constants, photochemical parameters and removal efficiencies of target CCBs are shown in Table 2. The degradation of AML, DIL and VER followed the pseudo-first-order kinetics with time-based rate constants (k_d) of 0.031, 0.044, and 0.011 min⁻¹, respectively. The molar absorption coefficients (ε₂₅₄) of target CCBs were determined from their absorption spectra (Fig. S3), and then the quantum yields at 254 nm (φ₂₅₄) could be calculated with Eq. (1) (Baenza and Knappe, 2011):

\[
φ_{254} = \frac{k_d \times U_{254}}{60 \times 0.77 \times \ln(10) \times ε_{254}}
\]

where U₂₅₄ is the molar photon energy at 254 nm (4.72 × 10² J E⁻¹), and 60 (min) and 0.77 (mW cm⁻²) convert the time-based k_d to its fluence-based value. As shown in Table 2, the ε₂₅₄ of AML, DIL and VER was 37266, 39153 and 8042 M⁻¹ cm⁻¹, respectively; and the φ₂₅₄ was 0.0037, 0.0050 and 0.0061 mol E⁻¹, respectively. After UV irradiation for 60 min, the removal efficiencies of AML and DIL could reach 84.7% and 93.6%, respectively; but VER was only removed by 51.2%. According to the acquired k_d values, the UV fluences required to remove 90% of AML, DIL and VER are thus calculated to be 3335, 2091 and 9315 mJ cm⁻², respectively, which are much higher than applied for wastewater disinfection (40–100 mg cm⁻²).

3.2. Identification of TBPs and proposed degradation pathways

The major TBPs were also identified in the relatively high concentration level experiments. The full scan of Q-Exactive MS was first used to register the exact mass of each TBP and determine their evolution (i.e., formation or decay) trends by calculating their relative concentrations. Afterward, the MS² scan was performed to obtain the fragmentation patterns of identified TBPs to elucidate their structures. This analytical approach is considered efficient in elucidating TBP structures when authentic standards are not available (Schymanski et al., 2014). Moreover, whether or not the functional pharmacophores existed in the TBP structures was also examined to assess the potential ecological hazards.

The major physicochemical properties and MS² fragmentation patterns of the target CCBs and identified TBPs are shown in Table S3.

### Table 2
Photolysis rate constants, photochemical parameters, and removal efficiencies of target CCBs (UV fluence = 2772 mJ cm⁻², pH = 7.0).

| Compound | Photolysis rate constant, k_d × 10² (min⁻¹) | R² | Molar absorption coefficient, ε₂₅₄ (M⁻¹ cm⁻¹) | Quantum yield, φ₂₅₄ × 10² (mol E⁻¹) | Removal (%) |
|----------|---------------------------------------------|----|---------------------------------------------|---------------------------------|------------|
| AML      | 3.07 ± 0.04                                 | 0.990 | 37266 ± 84                                  | 3.68 ± 0.11                     | 84.7       |
| DIL      | 4.43 ± 0.31                                 | 0.950 | 39153 ± 42                                  | 4.98 ± 0.34                     | 93.6       |
| VER      | 4.11 ± 0.03                                 | 0.942 | 8042 ± 40                                   | 6.08 ± 0.10                     | 51.2       |

* Mean ± standard deviation (n = 3).
Based on the evolution trends and chemical structures of identified TBPs, two degradation pathways of AML were proposed, which corresponded to two primary TBPs. One of the primary TBPs was amlodipine pyridine derivative (A-407, m/z 407.1371), which was probably formed through a radical cation intermediate (Fasani et al., 2008). Under UV irradiation, the A-407 could lose the chlorophenyl group to form A-297 or lose the aminoethoxyl group to form A-348(Cl) (Fig. 1a). The Cl atom of A-348(Cl) could be further attacked by -OH to form A-330. Jakimska et al. (2014) also detected the three TBPs (i.e., A-297, A-348(Cl), and A-330) during photolysis of AML with a xenon lamp (1000 W, 250–1000 nm) and classified them as the secondary byproducts. However, we found that A-330 only appeared after 15 min of UV irradiation, while A-348(Cl) appeared after 3 min and reached a peak concentration at 30 min. Therefore, A-330 tended to originate from the degradation of A-348(Cl) and should be classified as a tertiary byproduct. The second degradation pathway started with the formation of A-391, which was probably attributed to the attack of -OH on the phenol ring of AML. Then, A-391 could sequentially produce A-348 and A-332 by losing the aminoethyl group and further losing the hydroxyl group. It is noted that none of the TBPs in the second pathway has been reported previously. This is most likely because the former study was performed in river water (Jakimska et al., 2014), so the matrix materials could strongly compete for -OH and inhibit the formation of A-391. The inhibition on A-391 formation by co-existing matrix materials was also observed in this study (see Section 3.4).

Moreover, the two degradation pathways showed different elimination efficiencies of the functional pharmacophore (highlighted in red). In the first pathway, when A-407 was formed, the aromatization of the 1,4-dihydropiridine moiety could significantly alter the original steric structure of AML by rotating the chlorophenyl group around Site 4 (Fig. 1a) to form the delocalized π bond between the pyridine and phenol rings. As a result, the boat-form structure of 1,4-dihydropiridine, which is crucial for the receptor protein (Rojstaczer and Triggle, 1996), was destructed. By contrast, the pharmacophore persisted in the TBPs of the second (A-391) pathway because either the substitution of Cl with OH at Site 5 or the rupture of the aliphatic chain at Sites 7–10 could hardly reduce the therapeutic activity of 1,4-dihydropiridine pharmaceuticals (Yamamoto et al., 2006; Reimao et al., 2010). Therefore, the first pathway is more efficient in eliminating the ecological hazards related to AML.

In the degradation of DIL, a total of four TBPs were identified, including D-373 (by losing the acetyl group), D-355 (by losing the...
The degradation pathway of VER started with the bond cleavage between the methyl and the cationic amino groups (Fig. 1c), leading to the formation of two primary TBPs (V-151 and V-291) whose concentrations continuously increased during the reaction. Furthermore, V-167 could be produced when the phenol ring of V-151 was attacked by OH and V-277 could be produced when V-291 lost a methyl group attached to the cationic amino group. In UV photolysis, V-277 only appeared after 10 min and had similar MS2 fragments to V-291. Because the cationic amino group and the two phenol rings all account for the therapeutic activity (Mannhold et al., 1978; Toffoli et al., 1995), the four TBPs are free of ecological hazards.

3.3. Degradation of target CCBs in WWTP effluent

To simulate wastewater disinfection process, a low concentration level of target CCBs (100 ng L⁻¹ each) and a typical UV fluence range (0–230 mJ cm⁻²) were selected for photolytic experiments in the WWTP effluent. The degradation kinetics of target CCBs in MQ water and WWTP effluent are compared in Fig. 2. In the WWTP effluent, the degradation of AML and VER was obviously promoted. Specifically, the removal efficiency of VER was about 4 times that in MQ water at the end of the reaction (UV fluence = 230 mJ cm⁻²), while the concentration of AML was below its detection limit after 40 mJ cm⁻². This result agrees with other previous works. Jakimska et al. (2014) reported that the degradation of AML was promoted in river water and WWTP influent/effluent under the irradiation of a xenon lamp, compared to that in MQ water. Giri et al. (2014) suggested that the organic materials in the WWTP effluent should have triggered indirect photolysis to enhance the degradation of AML under UV irradiation. However, Kim et al. (2009) reported that the WWTP effluent could have different impacts (either promotion or inhibition) on the UV photolysis of different PPCPs. In the present study, we also found that the degradation of DIL in the WWTP effluent was significantly inhibited in the later phase of the reaction.

The background concentrations of three studied CCBs and their TBPs in the raw effluent were also determined. Table 3 shows that AML, DIL and VER had a background concentration of 2.5, 93.2 and 13.6 ng L⁻¹, respectively. Previous researches have also reported the occurrence of these CCBs in WWTP effluents (Batt et al., 2008; Spongberg and Witter, 2008; Tarconnicu et al., 2011). In addition, the two primary TBPs (A-407 and A-391) of AML were also detected in the raw effluent. As mentioned above, A-407 could be generated from the degradation of AML under UV or solar irradiation (Fasani et al., 2008; Jakimska et al., 2014), and A-391 could be generated by an attack of OH on the Cl atom. Therefore, their occurrence in the raw effluent might result from solar irradiation and/or biological transformation.

The relative concentration was used to depict the evolution (formation or decay) trends of the identified TBPs (Table 3). For a target TBP, the relative concentration was calculated by dividing a selected peak area by the maximum peak area detected during the whole reaction course (the background peak area was subtracted if this TBP was already detected in the raw effluent). Results indicate that although AML was completely removed after 40 mJ cm⁻², the two primary TBPs (i.e., A-391 and A-407) reached their maximum concentrations at 40 and 80 mJ cm⁻², respectively. Moreover, the formation of A-391 reflected the generation of OH in the UV irradiated solution. At the UV fluence of 160 mJ cm⁻², A-407 achieved a total removal, but 58.3% of A-391 remained in the solution. Because A-391 contains the functional pharmacophore (Fig. 1), the treated solution was still ecologically hazardous. As for DIL, the three primary TBPs (D-355, D-373 and D-401) reached their maximum concentrations at 40, 40 and 80 mJ cm⁻², respectively. Afterward, because the degradation of DIL was inhibited, no more TBPs were formed and their concentrations started to decrease. At the UV fluence of 160 mJ cm⁻², D-355 and D-401 were completely removed, while 68.4% of D-373, which contains the functional pharmacophore (Fig. 1), still remained in the treated solution. As for VR, the two primary TBPs (V-151 and V-291) exhibited a different...
evolution trend in the WWTP effluent, as compared to their unceasing accumulation in MQ water (Fig. 1). Both V-291 and V-151 reached their peak concentrations at 80 mJ cm\(^{-2}\) and then started to decrease, implying that the matrix materials may induce some further degradation pathways. Although 38.6% of V-151 remained at the UV fluence of 160 mJ cm\(^{-2}\), the treated solution was free of ecological hazards because V-151 does not contain the functional pharmacophore. In summary, all of the identified primary TBPs reached their maximum concentrations at a UV fluence ranging from 40 to 100 mJ cm\(^{-2}\), which is typical for disinfection of wastewater and reclaimed water (according to the EU guideline, 40–80 mJ cm\(^{-2}\) for wastewater and 100 mJ cm\(^{-2}\) for reclaimed water). It is seen that the ecological hazards of AML and DIL cannot be completely eliminated in the UV disinfection process.

3.4. Impacts of co-existing materials and pH on AML degradation

To explore the possible factors affecting the degradation of AML, a series of photolytic experiments was carried out in MQ water under typical pH conditions or with externally spiked Cl\(^{-}\) (1600 mg L\(^{-1}\)), HA (50 mg L\(^{-1}\) as TOC) and NO\(_3\) (48 mg L\(^{-1}\)) (Table S4). The degradation of AML and the evolution of detected TBPs (A-391, A-407 and A-330) were examined, as illustrated in Fig. 3.

**Table 3**

Relative concentrations (%) of target CCBs and their primary TBPs under UV irradiation in WWTP effluent.

| UV fluence (mJ cm\(^{-2}\)) | AML Primary TBPs | DIL Primary TBPs | VER Primary TBPs |
|-----------------------------|-----------------|-----------------|-----------------|
|                             | A-391 | A-407 | D-355 | D-373 | D-401 | V-151 | V-291 |
| 0                           | 100.0 | 0.0   | 100.0 | 0.0   | 100.0 | 0.0   | 100.0 |
| 40                          | 0.0   | 100.0 | 68.2  | 100.0 | 35.1  | 85.3  | 25.4  |
| 80                          | 0.0   | 88.8  | 100.0 | 79.0  | 95.4  | 85.5  | 100.0 |
| 100                         | 0.0   | 79.1  | 38.8  | 79.0  | 14.1  | 83.4  | 17.5  |
| 120                         | 0.0   | 76.3  | 1.7   | 73.5  | 0.0   | 76.3  | 0.0   |
| 160                         | 0.0   | 58.3  | 0.0   | 74.3  | 0.0   | 68.4  | 0.0   |
| 100%                        | 2.5   | 3.7   | 4.1   | 93.2  | 0.0   | 13.6  | 0.0   |

\(a\) The background concentration of a target CCB is calculated as \(C_{\text{BG}} = \frac{A_{\text{raw}}}{A_{\text{spiked}}} - \frac{A_{\text{raw}}}{A_{\text{raw}}} \times 100\), where \(A_{\text{spiked}}\) and \(A_{\text{raw}}\) denote the peak areas of a target CCB in the spiked and raw samples, respectively. At a spiked concentration of 100 ng L\(^{-1}\), the \(C_{\text{BG}}\) is mathematically equal to the concentration of a target CCB (ng L\(^{-1}\)) in the WWTP effluent. The background concentrations of TBPs are calculated in the same way, except that \(A_{\text{spiked}}\) is replaced by \(A_{100}\%\) (i.e., the largest detected peak area).

\(b\) Risk is assessed based on whether or not a target analyte contains the functional pharmacophore.

Fig. 3 shows that AML was degraded more quickly at 8.5 than at 7.0. The \(\phi_{2254}\) of AML, which denotes the efficiency of photolysis, was calculated to be 0.0037 and 0.0098 mol E\(^{-1}\) at pH values of 7.0 and 8.5, respectively, based on its \(c_{2254}\) (Fig. S3). The change of pH could affect the speciation of AML in the reaction solution and subsequently alter its reactivity under UV irradiation. Baeeza and Knappe (2011) also found that the \(\phi_{2254}\) increased with increasing pH for the UV photolysis of trimethoprim. Correspondingly, A-391 was generated more quickly at pH 8.5 (Fig. 3b). However, the formation of A-407 was a little slower at pH 8.5 (Fig. 3c), probably because it was degraded faster to produce A-330 (Fig. 3d). A higher pH seemed to facilitate the substitution of Cl with OH on the phenol ring.

**Fig. 3.** Degradation of AML (a) and evolution of identified TBPs including A-391 (b), A-407 (c), and A-330 (d) under different experimental conditions.
In the presence of Cl⁻ and at pH 7.0, the degradation of AML was the fastest in the first 4 min and then slowed down remarkably (Fig. 3a). It is well known that UV irradiation of Cl⁻ produces Cl₂, which can easily react with organic compounds. As the evolution trends of A-391 and A-407 are similar to those in MQ water at pH 7.0 (Fig. 3b and c), the initially promoted degradation of AML was probably attributed to the formation of chlorinated compounds. Results also indicate that the concentration of A-330 started to increase slowly after 24 min and was much lower than that in MQ water (Fig. 3d). Considering that A-391 continuously accumulated in the solution over the whole reaction course, A-330 could only originate from the degradation of A-407. This result partly confirms the degradation pathways of AML as proposed in Fig. 1a.

In the presence of HA and at an initial pH of 5.9, the degradation of AML was inhibited to some extent (Fig. 3a). Because HA strongly competed for ·OH (Wols and Hofman-Caris, 2012), the formation of A-391 and A-330, which was triggered by an attack of ·OH on the Cl atom, was obviously inhibited (Fig. 3b and d). Consequently, the inhibited formation of A-330 led to the continuous accumulation of A-407 during most of the reaction course (Fig. 3c). Therefore, the A-407 pathway predominated in the degradation of AML when a high concentration of HA was available.

Among all tested influential factors, NO₃⁻ showed the greatest promotion on the degradation of AML (Fig. 3a). Along with the degradation of AML, A-391 and A-407 were formed quickly in the beginning of the reaction and then decayed gradually (Fig. 3b and c), while A-330 unceasingly accumulated over the whole reaction course. As UV irradiation of NO₃⁻ produces ·OH with φ₉OH of up to 9% (Mack and Bolton, 1999), the further degradation of A-391 and A-407 could be ascribed to the indirect oxidation of ·OH.

4. Conclusions

In this study, the degradation kinetics and pathways of three CCBs under UV irradiation were investigated in both MQ water and WWTP effluent. In addition, the elimination of functional pharmacophores from the TBP structures was also examined to assess the potential ecological hazards. Based on the experimental results, the following conclusions can be drawn:

- UV photolysis of three studied CCBs conform to the pseudo-first-order reaction kinetics, with rate constants of 0.031, 0.044, 0.011 min⁻¹ for AML, DIL, and VER, respectively.
- Several TBPs of AML and DIL still contained the functional pharmacophores, while those of VER were free of ecological hazards.
- The pharmacophore-containing TBPs of AML and DIL reached their peak concentrations under UV fluences typical for disinfection of wastewater and reclaimed water (i.e., 40–100 mJ cm⁻²), indicating potential ecological hazards.
- The degradation of VER and AML was promoted in the WWTP effluent, while the degradation of DIL was inhibited. The removal efficiencies of AML, DIL, and VER in the WWTP effluent were 100%, 21%, and 32% at 100 mJ cm⁻², respectively.
- An increase in pH (from 7.0 to 8.5) or the presence of NO₃⁻ promoted the degradation of AML, while the presence of HA exerted an inhibiting effect. In the presence of Cl⁻, the degradation of AML was the fastest in the beginning of the reaction and remarkably slowed down afterward.

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Appendix A. Supplementary data

Supplementary data related to this article can be found at http://dx.doi.org/10.1016/j.watres.2015.05.028.

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