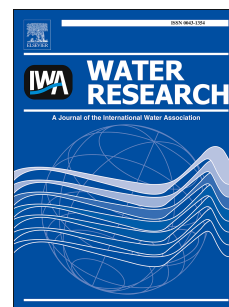


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Jennifer E. Schollée, Marc Bourgin, Urs von Gunten, Christa S. McArdell, Juliane Hollender



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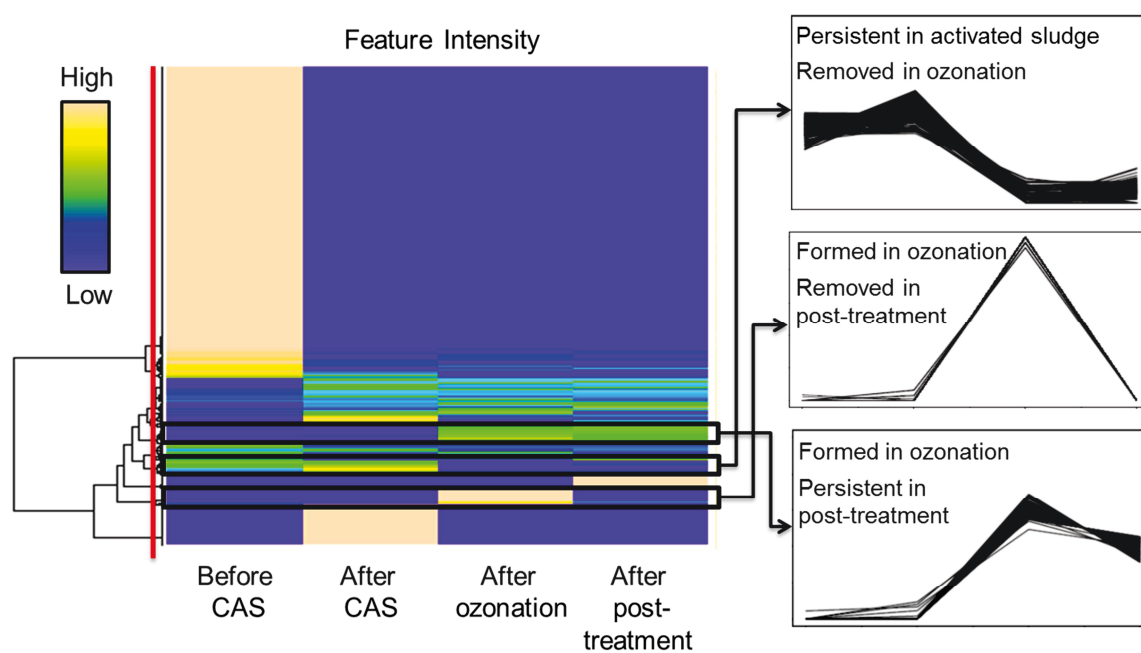
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# **Non-target Screening to Trace Ozonation Transformation Products in a Wastewater Treatment Train including different Post-treatments**

Jennifer E. Schollée,<sup>a,b</sup> Marc Bourgin,<sup>a</sup> Urs von Gunten,<sup>a,b,c</sup> Christa S. McArdell,<sup>a</sup> Juliane Hollender<sup>a,b,\*</sup>

<sup>a</sup>Eawag, Swiss Federal Institute of Aquatic Science and Technology, 8600 Dübendorf, Switzerland

<sup>b</sup>Institute of Biogeochemistry and Pollutant Dynamics, ETH Zurich, 8092 Zurich, Switzerland

<sup>c</sup>School of Architecture, Civil, and Environmental Engineering (ENAC), École Polytechnique Fédérale de Lausanne, 1015 Lausanne, Switzerland

\*Corresponding author: [juliane.hollender@eawag.ch](mailto:juliane.hollender@eawag.ch)

Additional author emails

Jennifer E. Schollée – [jennifer.schollee@eawag.ch](mailto:jennifer.schollee@eawag.ch)

Marc Bourgin – [marc.bourgin@eawag.ch](mailto:marc.bourgin@eawag.ch)

Urs von Gunten – [vongunten@eawag.ch](mailto:vongunten@eawag.ch)

Christa S. McArdell – [christa.mcardell@eawag.ch](mailto:christa.mcardell@eawag.ch)

## Abstract

Ozonation and subsequent post-treatments are increasingly implemented in wastewater treatment plants (WWTPs) for enhanced micropollutant abatement. While this technology is effective, micropollutant oxidation leads to the formation of ozonation transformation products (OTPs). Target and suspect screening provide information about known parent compounds and known OTPs, but for a more comprehensive picture, non-target screening is needed. Here, sampling was conducted at a full-scale WWTP to investigate OTP formation at four ozone doses (2, 3, 4, and 5 mg/L, ranging from 0.3-1.0 gO<sub>3</sub>/gDOC) and subsequent changes during five post-treatment steps (*i.e.*, sand filter, fixed bed bioreactor, moving bed bioreactor, and two granular activated carbon (GAC) filters, relatively fresh and pre-loaded). Samples were measured with online solid-phase extraction coupled to liquid chromatography high-resolution tandem mass spectrometry (LC-HRMS/MS) using electrospray ionization (ESI) in positive and negative mode. Existing non-target screening workflows were adapted to (1) examine the formation of potential OTPs at four ozone doses and (2) compare the removal of OTPs among five post-treatments. In (1), data processing included principal component analysis (PCA) and chemical knowledge on 31 possible oxidation reactions to prioritize non-target features likely to be OTPs. Between 394 and 1328 unique potential OTPs were detected in positive ESI for the four ozone doses tested; between 12 and 324 unique potential OTPs were detected in negative ESI. At a specific ozone dose of 0.5 gO<sub>3</sub>/gDOC, 27 parent compounds were identified and were related to 69 non-target features selected as potential OTPs. Two OTPs were confirmed with reference standards (venlafaxine *N*-oxide and chlorothiazide); 34 other potential OTPs were in agreement with literature data and/or reaction mechanisms. In (2), hierarchical cluster analysis (HCA) was applied on profiles detected in positive ESI mode across the WWTP and revealed 11 relevant trends. OTP removal was compared among the five post-treatments and 54-83% of the non-target features that appeared after ozonation were removed, with the two GAC filters performing the best. Overall, these data analysis strategies for non-target screening provide a useful tool to understand the behavior of unknown features during ozonation and post-treatment and to prioritize certain non-targets for further identification.

## 28 **Keywords**

29 Advanced wastewater treatment

30 Ozonation

31 Non-target screening

32 Transformation products

33 Micropollutants

34

35

## 1.1 Introduction

Thousands of organic compounds originating from domestic and industrial sources are discharged to municipal wastewater treatment plants (WWTPs). During conventional treatment in WWTPs, these compounds can be removed through sorption or degradation, or pass through treatment unchanged (Loos et al. 2013, Luo et al. 2014). The subsequent discharge of treated wastewater introduces compounds that are not removed during treatment into receiving waters, where they have the potential to affect downstream aquatic communities or contaminate drinking water resources (Kolpin et al. 2002, Reungoat et al. 2010). To enhance the abatement of micropollutants in wastewater treatment, advanced treatment, with either ozonation or activated carbon-processes, such as granular activated carbon (GAC) filtration or powdered activated carbon (PAC) treatment, is needed after conventional activated sludge (CAS) treatment. Both technologies have been shown to successfully remove a large number of compounds, including micropollutants (Boehler et al. 2012, Hernández-Leal et al. 2011, Hollender et al. 2009, Snyder et al. 2006). Ozonation in combination with biological post-treatment has been demonstrated to reduce the ecotoxicity of the effluent water (Escher et al. 2009, Margot et al. 2013, Prasse et al. 2015, Stalter et al. 2011).

During ozonation, organic compounds either react directly with ozone or the secondary oxidant hydroxyl radical (Staehelin and Hoigne 1985, von Gunten 2003a). These reactions can lead to the formation of ozonation transformation products (OTPs). Reactions with ozone involve electron-rich moieties (e.g., olefins, activated aromatic rings, neutral amines and thioethers), while reactions with hydroxyl radicals are much less specific (von Sonntag and von Gunten 2012). Many types of transformation reactions can occur and a variety of OTPs can be formed, e.g., *N*-oxides from tertiary amines, hydroxylated compounds from aromatic moieties, or aldehydes from olefins. Information on mechanisms has been used to predict possible OTPs from known parent compounds (Lee et al. 2017). In addition to OTP formation, oxidation by-products such as *N*-nitrosodimethylamine (NDMA) or bromate are (potentially) formed during ozonation (von Gunten 2003b, Wert et al. 2009, Zimmermann et al. 2011). It is assumed, and for some compounds has been confirmed, that the formation of oxygen-rich compounds such as aldehydes and carboxylic acids potentially leads to increased biodegradability (Hübner et al. 2015, Zimmermann et al. 2011). It is therefore recommended to apply a

biological post-treatment step after ozonation to remove biodegradable oxidation by-products formed from the organic matrix or OTPs (Hollender et al. 2009, Stalter et al. 2011, von Gunten 2003a).

Sand filtration has been used after ozonation as a biological post-treatment, where it has been shown to be effective for removal of NDMA and reduction of potential ozonation-induced toxicity in full-scale wastewater treatment (Hollender et al. 2009, Prasse et al. 2015). Bioreactors such as moving bed or fixed bed reactors have been proposed as alternatives for biological post-treatment of wastewater (Accinelli et al. 2012, Falås et al. 2012, Luo et al. 2014). Treatment with GAC can remove micropollutants or OTPs through sorption, depending on the compound properties or, after a period of adaptation through biological degradation (Grover et al. 2011, Simpson 2008). However, as previously reported, biological post-treatments are often only moderately effective at eliminating known OTPs (Margot et al. 2013) and ozonation may not always lead to an expected increase of OTPs biodegradability compared to the parent compounds (Hübner et al. 2015). In the case of known OTPs, it is possible to monitor their presence and removal in full-scale WWTPs (Borowska et al. 2016, Deeb et al. 2017, Zimmermann et al. 2011). However, the elucidation of OTPs of single substances is very labor-intensive and therefore data remain scarce. Consequently, many OTPs remain unknown, making evaluation of different treatment types difficult and biased toward known compounds. Therefore, application of non-target screening is promising to characterize unknown peaks in WWTPs and to determine if unknown, non-target compounds are either abated or formed during wastewater treatment.

Liquid chromatography coupled to high-resolution tandem mass spectrometry (LC-HRMS/MS) is increasingly applied for the analysis of known and unknown emerging contaminants in water (Hernández et al. 2014, Richardson and Kimura 2016). Numerous studies have shown that reverse-phase LC-HRMS/MS can be used for the simultaneous analysis of hundreds of known compounds including TPs (Bletsou et al. 2015, Cotton et al. 2016, Ferrer and Thurman 2012). Additionally, information from HRMS/MS measurement (such as exact mass and isotope information) can be acquired for suspect or non-target compounds and can then be used for structure elucidation of parent compounds or TPs without reference standards (Krauss et al. 2010, Müller et al. 2012, Singh et al. 2015). However, the breadth of information produced from LC-HRMS/MS can no longer be analyzed manually and requires the addition of data analysis strategies to prioritize unknown peaks of interest.

Recently, differential/statistical analysis has been used to compare removal of non-target compounds in advanced oxidation processes (Müller et al. 2012, Parry and Young 2016) and to understand the fate of unknown compounds in a conventional WWTP by prioritizing features for identification based on intensity profiles (Getzinger 2016).

To study the impact of ozonation and five post-treatments on the fate of micropollutants and OTPs, a large sampling campaign was conducted in a full-scale WWTP in 2014-2015. The WWTP is the first in Switzerland upgraded with ozonation to comply with the new Swiss water protection act that aims to reduce the load of micropollutants to receiving waters (Eggen et al. 2014). The aims of this study were to expand on the target and suspect screening approaches used in Bourgin et al. (2018) through the application and further development of existing non-target screening workflows (Chiaia-Hernández et al. 2017, Schollée et al. 2015) to the expansive data set, considering four different ozone doses and five post-treatments. First, non-target peaks detected either before or after ozonation were characterized with principal component analysis (PCA) and potential parent compounds and potential OTPs were linked together using expected ozone and/or OH radical reaction mechanisms. Second, hierarchical cluster analysis (HCA) was applied to compare the efficiencies of five post-treatment technologies in removing non-target features, focusing on OTPs that were detected.

## 2.1 Methods

### *2.1 Sampling*

Samples were collected at the full-scale WWTP Neugut (Dübendorf, Switzerland). Wastewater includes mostly residential sources as well as a number of food- and fragrance-related industries. The original treatment consists of primary treatment and CAS (both aerobic and anaerobic) treatment, including biological phosphorus removal. Effluent from the CAS treatment step is passed to an ozonation reactor that has been in full-scale operation since March 2014, followed by treatment with a sand filter. The ozonation system includes a multi-chamber reactor with a total volume of 530 m<sup>3</sup> and an average retention time of 43 minutes during dry weather. Ozone was injected in the first reaction chamber and was completely depleted at the effluent. After ozonation, a variety of post-treatment technologies were applied in pilot-scale and sampled for target and non-target analyses (Bourgin et al. 2018).



For the investigation of the four ozone doses, sampling was carried out in September/October 2014 (Supporting Information (SI), Table S1) during dry weather conditions. For each ozone dose, three 24-h flow-proportional composite samples were collected on consecutive days at the influent and effluent of the ozone reactor, for a total of 24 samples. Low variability was observed for target compounds and DOC among the three sampling days, and they were therefore considered as replicates for the PCA. Dissolved organic carbon (DOC) content ranged from 3.5 to 6 mg/L after CAS treatment. The following four ozone doses were tested: 2, 3, 4, and 5 mg/L ozone, corresponding to specific ozone doses of  $0.35 \pm 0.02$ ,  $0.54 \pm 0.05$ ,  $0.67 \pm 0.03$ , and  $0.97 \pm 0.07$  gO<sub>3</sub>/gDOC, respectively, for the three replicates. Exact values for each sampling date are provided in Table S1.

For sampling of the complete treatment train (including post-treatment systems), samples were collected as 24-h or 48-h composites at the following four locations: (1) before CAS treatment; (2) after CAS treatment, at the influent of the ozone reactor; (3) after the ozone reactor, before post-treatment; and (4) after one of the five post-treatments at the effluent of the treatment plant (SI, Figure S1 and Table S2). Four samples were collected for each location (total 32 samples) from February to May 2015. Ozone dosing was set around 2.7 mg/L ozone (ranging between 2.13 and 2.89 mg/L ozone or 0.45-0.57 gO<sub>3</sub>/gDOC). Five post-treatment setups were operated in parallel. Two GAC filters were used, one relatively fresh (GAC-A – sampled at 14,000–24,000 bed volumes (BV)) and one already pre-loaded prior to implementation with reduced sorption capacity but potentially stimulated biological activity (GAC-B – sampled at 36,000–46,000 BV). Additionally, three biologically active systems were tested – a moving bed bioreactor (MB), a fixed bed bioreactor (FB), and a full-scale sand filter (SF). The post-treatment reactors were fed with the effluent of the ozonation reactor to condition them and ensure the formation of a biofilm, with the exception of GAC-B which was conditioned up to 16,000 BV with the effluent of a secondary treatment. Details on each of the post-treatments is available in the SI. Samples were collected in 1L glass bottles and stored at 4°C until analysis. In the case of storage >7 days, samples were frozen at -20°C until analysis.

## 2.2 Measurement by online SPE-LC-HRMS/MS

Measurement details are provided in the SI, Section S2; quality control and target screening results are reported in Bourgin et al. (2018). Briefly, samples were filtered, diluted with nanopure water

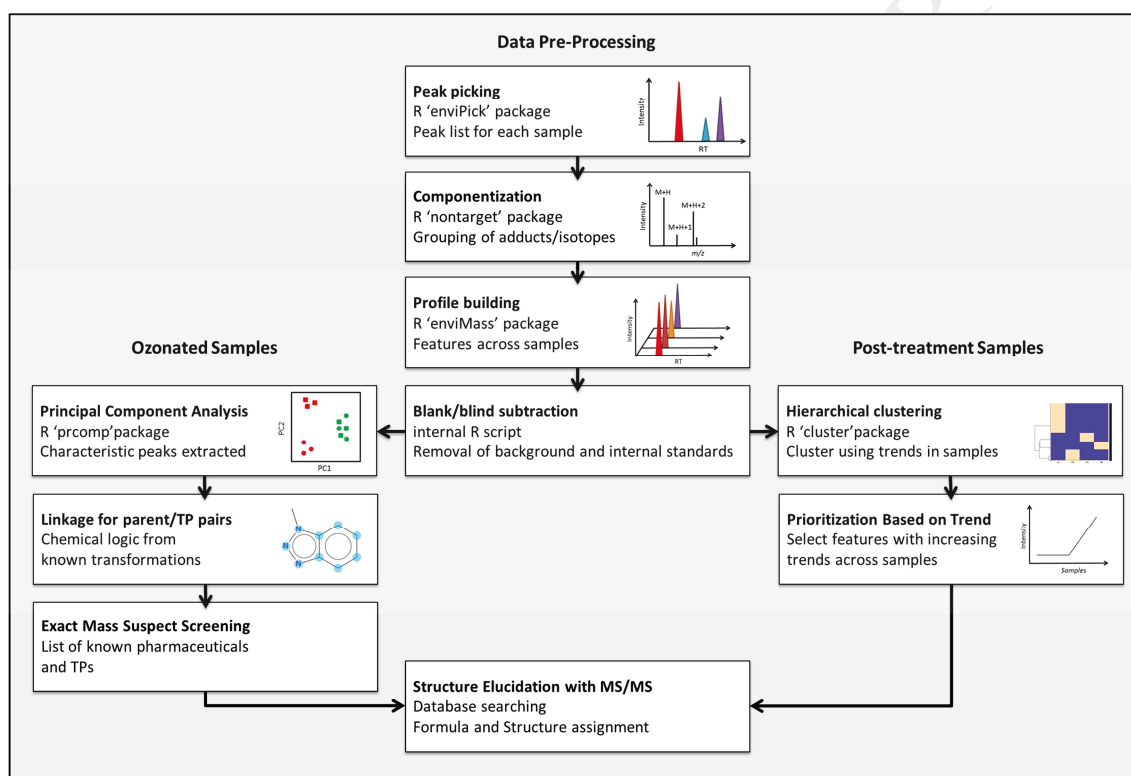
(Barnstead Nanopure, Thermo Scientific), and then spiked with 51 isotopically-labeled standards. A 20 mL sample was injected for measurement with online solid-phase extraction (SPE), using a multi-layer cartridge (Oases HLB (Waters), Strata X-AW and Strata X-CW (Phenomenex), Env+ (Biotage), followed by LC-HRMS/MS. Reverse-phase LC separation was done with an Atlantis T3 column (Waters). The enrichment and measurement methods were shown to be effective for a wide range of polar and medium polar organic micropollutants (Huntscha et al. 2012); however, highly polar OTPs that can be formed during ozonation may not be detected with this method.

Samples were measured on an Orbitrap mass spectrometer (Q Exactive Plus, Thermo Scientific; resolution 70,000 at 200  $m/z$ ) with electrospray ionization (ESI) in positive and negative modes separately. Settings are listed in SI, Table S4. An external mass calibration was performed and, using the spiked internal standards, the difference between measured accurate mass and expected exact mass was calculated to estimate mass accuracy of the measurements. For the majority of compounds and samples, mass accuracy was greater than 5 ppm, but in rare cases, deviations of up to 15 ppm were recorded. Consequently, a mass recalibration was performed (see below). A Top-5 data dependent acquisition method was used to trigger MS/MS measurements, using an inclusion list of 67 target compounds, a dynamic exclusion window of 3.0 seconds, and the “pick others” feature enabled. In this way, MS/MS measurements were triggered first on masses listed on the inclusion list, and second on the top 5 most abundant ions in the corresponding full scan. With the dynamic exclusion window, ions that were fragmented in one scan cycle were not fragmented again for 3.0 seconds, ensuring the greatest possible coverage of MS/MS fragmentation. A list of inclusion masses can be found in the SI, Table S5. Matrix effects of 43 target micropollutants were evaluated and intensities were found to vary by less than 10% between the different sample matrices for the individual substances (Bourgin et al. 2018).

### 2.3 Data Analysis

A flowchart of the general workflow is shown in Figure 1. Samples from ozonation and post-treatment analyses were processed separately because of the different sampling periods and study objectives. For evaluation of the four ozone doses, principal component analysis (PCA) and so-called linkage analysis were combined to investigate the possible OTPs resulting from the treatment. This workflow

has been demonstrated to be effective for prioritizing potential biological TPs resulting from CAS treatment (Schollée et al. 2015) and was adapted here for known oxidation reactions. Potential OTPs were prioritized with this workflow and their behavior in post-treatment samples was investigated. To probe possible trends of non-target features, especially the formation and elimination of OTPs across the entire wastewater treatment train, HCA was applied to samples from influent through CAS treatment and ozonation to the effluent of the five post-treatment setups. All data analysis was performed with the R software (R Development Core Team 2017). Specific information about each of the elements of the workflow is outlined below.



**Figure 1. Flowchart of data analysis with an illustration of the workflow. Pre-processing steps were similar for ozonated and post-treatment samples, apart from different replicate filters and a retention time alignment and mass-to-charge recalibration steps applied to the post-treatment samples to correct for multiple measurement runs. Subsequent multivariate analyses were different for the two sampling campaigns (ozonation (left), post-treatment (right)).**

### 2.3.1 Pre-Processing

All data files were converted and peaks centroided with the software ProteoWizard MSConvert (v.3.0.8725) (Chambers et al. 2012) to ".mzXML" files. Peak picking of each file was carried out with the R package 'enviPick' (Loos 2014b) (settings in SI, Table S6), whereby a feature (peak) list was

generated for each sample file, where each feature represents a unique  $m/z$  and retention time (RT) pair. Measurements for the post-treatment analysis were spread out over the course of the sampling campaign; therefore,  $m/z$  recalibration and RT alignment was performed using internal standards with the R package 'enviMass' (Loos 2014a), to account for batch deviations. Results of the  $m/z$  recalibration indicated that in one file and then in a negligible number of features were mass shifts of up to 15 ppm observed. These feature lists were then analyzed with the R 'nontarget' package (Loos 2015), where features belonging to the same compound, such as those related to isotopes, were grouped together. Thereafter, features were selected that were either classified as the monoisotopic peak of a compound or not related to other features (*i.e.*, unpaired), with the goal to reduce the amount of data through the removal of isotopes.

The reduced feature lists from all samples were then merged with the R package 'enviMass' (Loos 2014a) using the settings in SI, Table S7. The resulting data sets generated from the enviMass package included a row for each feature, a column for each sample, and was populated with the feature intensities. Then a blank (*i.e.*, nanopure water) and blind (*i.e.*, nanopure water spiked with internal standards) subtraction was performed, where any feature detected in a blank/blind sample was removed from the feature list. A blind subtraction was included to remove any potential noise or contamination resulting from the measurement, as well as the spiked internal standards. Finally, a restriction was applied such that a feature had to be detected in at least three samples. For the ozonation analysis, the feature had to be detected in any three samples; for the post-treatment analysis the restriction was applied on each sample type separately. Applying this restriction allowed the removal of noise peaks from the peak list, and ensured that non-target features were relevant for further investigation due to their continued presence in the WWTP.

### 2.3.2 Principal Component Analysis

Each feature was standardized (mean zero, unit variance) and PCA was performed with the R function 'prcomp'. The PCA results showed which samples grouped together and in which sample type features were most often detected, allowing for feature characterization. Using expected trends (*i.e.*, decreasing intensities for parent compounds or increasing intensities for OTPs), features could be tentatively classified as potential parent compounds or potential OTPs (or neither).

### 2.3.3 Linkage Analysis

After classification with PCA, a so-called linkage analysis was done. To find possible OTPs, the masses of likely OTPs were calculated using the exact mass of the potential parent compounds (obtained with the LC-HRMS/MS and classified with PCA) and the mass difference resulting from a list of proposed transformation reactions (see below). Proposed transformations stem from either reactions of the parent compound with ozone or hydroxyl radicals (or both) since both possibilities need to be considered during ozonation of real wastewaters. A summary of the reactions that were considered, with example reactions, is provided in the SI, Table S8 and the necessary R functions are available at <https://github.com/dutchjes/TPPrioritizR>. The resulting list of calculated OTP masses was then screened against the masses of potential OTPs classified by the PCA, using a 15 ppm mass tolerance, the maximum mass error observed in the samples. In the case of a match, a “link” was established between the potential parent compound and a potential OTP. The linkage analysis was carried out separately for each ozone dose so that the numbers and types of transformation reactions resulting from the four ozone doses could be compared.

Thirty-one transformation reactions (SI, Table S8) were considered and included additions of 1-3 atoms of oxygen, losses of various alkyl groups, and dehydrogenation. Six (de)conjugation reactions, as well as a number of reductive reactions, that were not expected during ozonation, were also included to provide an assessment of the potential number of false positive links. The linkage analysis was run through one cycle only; therefore, second- or higher-generation OTPs would not be detected, although in reality many of the transformations included are likely the result of multiple reaction steps (*i.e.*, addition of multiple oxygen atoms). Not all potential OTPs detected are expected to arise from reactions with micropollutants, some might result as by-products from the organic matrix.

### 2.3.4 Suspect Screening and Structure Elucidation

Potential links from a specific ozone dose of  $0.54 \pm 0.05$  gO<sub>3</sub>/gDOC (2.7 mg/L ozone) were further prioritized for structure elucidation. This dose was selected for the following two reasons: (1) the highest number of links was detected at this dose and (2) it is the recommended specific ozone dose for the required abatement of micropollutants at the WWTP and was therefore used in the post-treatment analysis. Suspect screening was carried out on the parent masses present in the linkage results. The suspect list that was used included 195 target micropollutants (including OTPs) monitored

at this site (Bourgin et al. 2018) and an additional 1022 suspected pharmaceuticals (Singer et al. 2016). HRMS/MS spectra from any potential parent compound that matched to a suspect compound was then compared to reference standard spectra measured in-house (available at MassBank (Horai et al. 2010)). In the case of a positive identification of a suspect compound, structures were proposed for the associated potential OTP(s) based on the parent structure, expected transformation reaction as indicated by the linkage analysis, and knowledge on ozone reactivity (von Sonntag and von Gunten 2012). Confidence in structure assignment is reported using the levels proposed in Schymanski et al. (2014). Plausibility of OTP formation was further confirmed through prediction of OTPs with an ozone pathway prediction system (O<sub>3</sub>-PPS) (Lee et al. 2017), as well as literature searches.

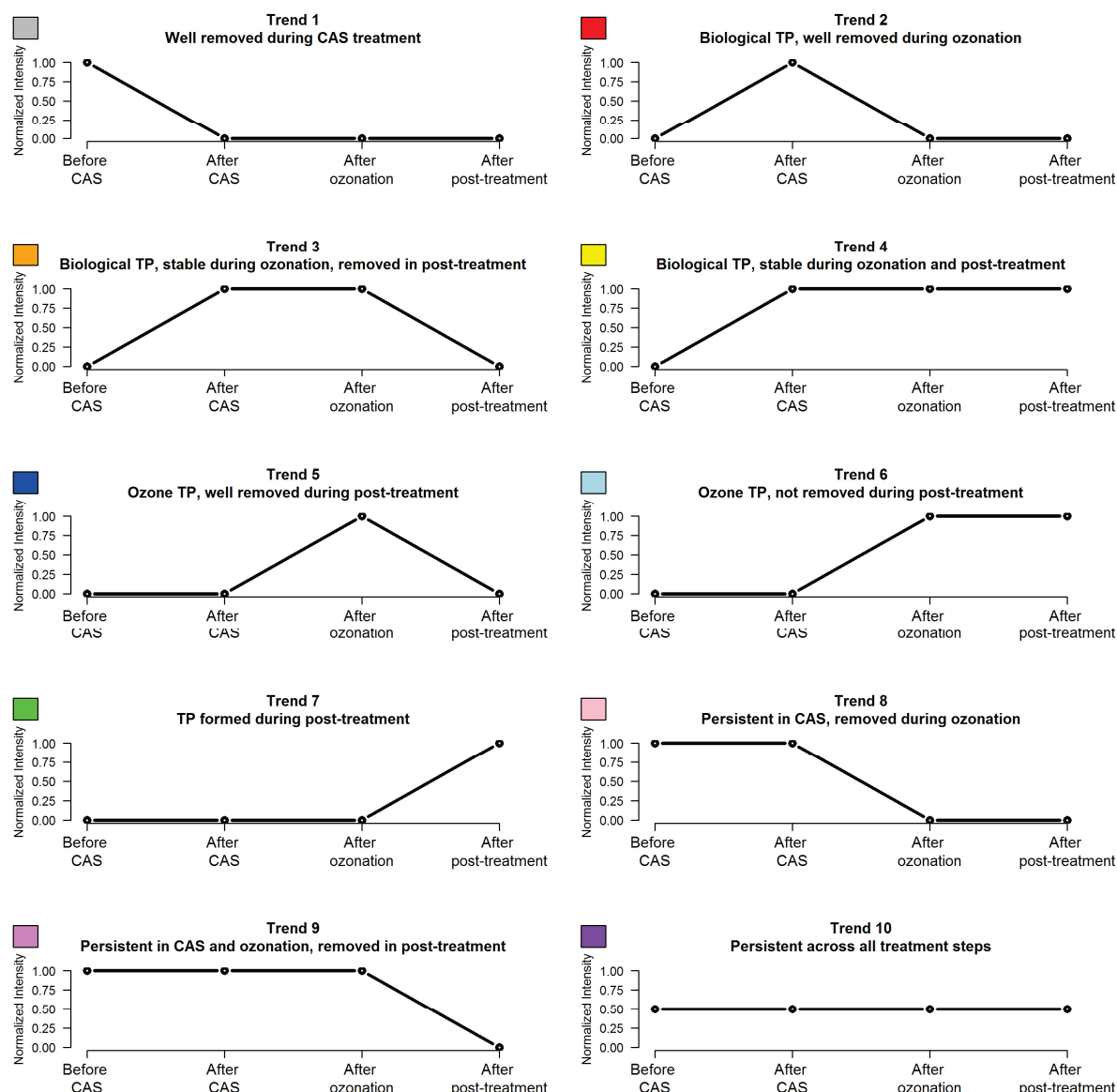
### 2.3.5 Hierarchical Cluster Analysis (HCA)

For analysis of the post-treatment systems, the trends of features across the WWTP were investigated, and similar feature profiles were clustered together with HCA. The majority of the links and potential OTPs were found in the positive ESI mode in the PCA and linkage analysis; therefore, we focused on this data for the HCA. To observe the behavior of features across the WWTP, first a feature list was generated for each of the four sample types (SI, Figure S1), where feature intensity corresponded to the maximum intensity detected in that sample type, multiplied by the corresponding dilution factor. The feature lists were then profiled together with the R package 'enviMass' (Loos 2014a) and a dissimilarity matrix was calculated using intensities normalized to the maximum intensity with the R package 'cluster' (Maechler et al. 2016). Finally a HCA was performed with the profiles using the Euclidian distance (which is the length of the line connecting them) and Ward method (designed to minimize intra-cluster variance, thereby producing compact clusters). Considering that the intensities of a peak could either increase, decrease, or remain the same from one sampling point to the next, there are  $3^{(n-1)}$  possible trends, corresponding to 27 possible trends in this case. Therefore, the resulting dendrogram was cut such that 30 clusters were identified and the profiles in each of these clusters visualized. Eleven trends were defined, corresponding to expected trends in the WWTP (Figure 2) and each of the 30 clusters was visually assigned to a trend that best described the behavior of the profiles in that cluster. Characteristics of features assigned to different trends were then compared. The R-script used for cluster analysis and characterization is available at <https://github.com/dutchjes/TPPrioritizR>. Validation of the HCA was performed with a set of eight

299 parent compounds (amisulpride, carbamazepine, cetirizine, clarithromycin, lamotrigine, ranitidine,  
300 tramadol, and venlafaxine) and their respective OTPs, known to be formed during ozonation.

301

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**Figure 2.** Examples of the ten trends that were defined to describe the intensity profiles in a cluster. Each trend can be associated with the specific behavior of a compound in the wastewater treatment plant, and is described in the title of each trend. A Trend 11 was also defined but is not shown here, including any cluster that did not fit into the other categories. Intensity profiles for each post-treatment were clustered separately and are therefore generally labeled as 'After post-treatment'. CAS refers to conventional activated sludge treatment.

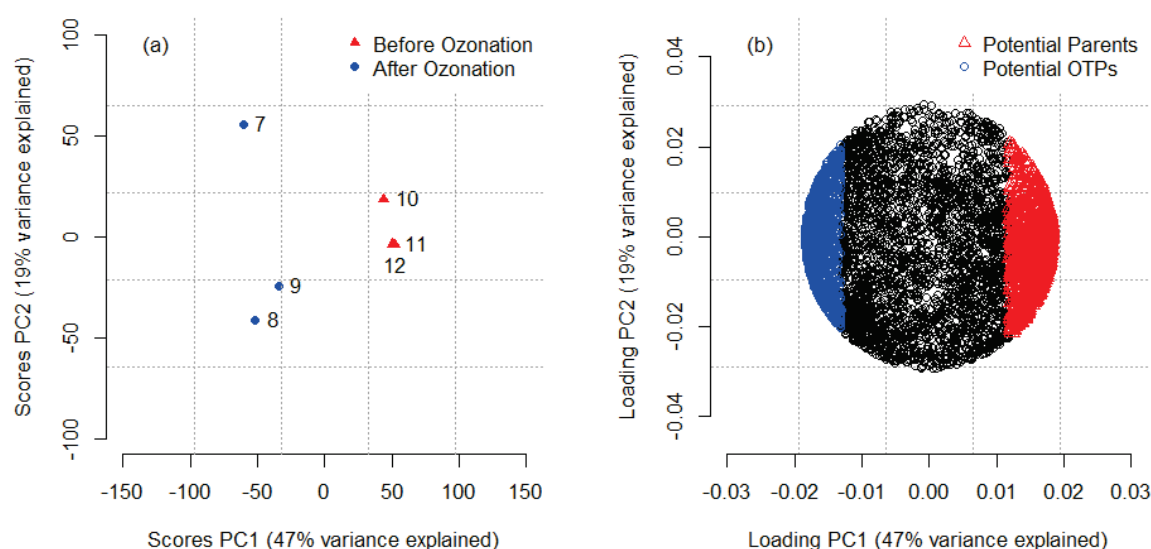
## 3.1 Results and Discussion

### 3.1 Analysis of Ozonated Samples

#### 3.1.1 Principal Component Analysis and Linkage Analysis



Classification of non-target features detected with LC-HRMS/MS was performed by interpretation of the PCA scores and loading plots on samples collected from the four specific ozone doses (0.4, 0.5, 0.7, and 1.0 gO<sub>3</sub>/gDOC). The total number of non-target features detected and included in the PCA for the specific ozone doses 0.4, 0.5, 0.7, and 1.0 gO<sub>3</sub>/gDOC were 4260, 6207, 3798, and 2279, respectively (Table 1). An example scores plot is shown for 0.5 gO<sub>3</sub>/gDOC in Figure 3a; scores plots from the other ozone doses for both ionization modes are shown in SI, Figures S2 and S3. In each case, the first principal component (PC1) could explain approximately 50% of the variance in the data sets (SI, Tables S9 and S10), which can be attributed to the difference in sample types (*i.e.*, before vs. after ozonation). PC2 (plotted along the y-axis) generally explained between 15 and 20% of the variance among the samples and can be attributed to daily changes in the wastewater composition, while additional PCs were related to differences in individual samples (SI, Figure S4).



**Figure 3. Principal component analysis (a) scores plot and (b) loading plot of WWTP effluent before and after ozonation for a specific ozone dose of 0.5 gO<sub>3</sub>/gDOC. The first and second principal components are shown, on the x- and y-axes, respectively. A good separation between the samples before ozonation (red triangles) and after ozonation (blue circles) was obtained along the x-axis and could explain 47% of the variance, allowing for classification of non-target features as potential parent compounds or potential OTPs (or neither, black circles). Similar results were obtained for the other ozone doses (SI, Figure S2). Numbers refer to the sample IDs (Table S1).**

Features were classified based on their loading in PC1 (Figure 3b; SI, Figures S6 and S7). The top 25% of features (*i.e.*, features with the highest PC1 loadings, and therefore characteristic of the samples before ozonation) were classified as potential parent compounds. Similarly, the bottom 25% of features (*i.e.*, features with the lowest PC1 loadings and therefore characteristic of the samples after ozonation) were classified as potential OTPs. This procedure resulted in equal numbers of features as potential parent compounds and potential OTPs. In the positive mode between 570 (1.0 gO<sub>3</sub>/gDOC) and 1552 (0.5 gO<sub>3</sub>/gDOC) features were selected as potential parents/OTPs (Table 1). In negative mode, between 68 (1.0 gO<sub>3</sub>/gDOC) and 430 (0.5 gO<sub>3</sub>/gDOC) features were classified as potential parents/OTPs (Table 1). Wastewater treated with 0.5 gO<sub>3</sub>/gDOC had the highest number of features, indicating that with this dose possibly the highest number of OTPs was detected (SI, Tables S11 and S12). This result is in agreement with previous work from Bourgin et al. (2018), which found that at this dose the highest number of suspected OTPs was detected. DOC removal was observed to be approximately 10%, independent of the specific ozone dose. The decrease in detected features at higher specific ozone doses is likely due to the fact that these higher generation OTPs were -too polar to be detected with online-SPE LC-HRMS/MS. This could be caused by either poor enrichment in the SPE cartridge or low retention on the reverse-phase LC column, and an alternative analytical method (*e.g.*, hydrophilic interaction liquid chromatography (HILIC)) would be required to detect them. Characteristics of features (*e.g.*, mass to charge ratios ( $m/z$ ), retention times, intensities, or mass defects (*i.e.*, the difference between the nominal mass and the exact mass)) detected after ozonation from the four ozone doses were compared and effluent features from the 0.5 gO<sub>3</sub>/gDOC were in some cases significantly different (SI, Figures S7 and S8).

From the linkage analysis with 31 transformation reactions, the number of unique features with links to potential OTPs ranged from 17–1416, depending on ionization mode and specific ozone dose, and represented between 4 to 21% of all detected features (Table 1). In positive ESI mode, the highest number of links were detected in samples with a specific ozone dose of 0.5 gO<sub>3</sub>/gDOC (Figure 4a). Consequently, the highest total number of links for both ESI modes (9636) was found in samples from 0.5 gO<sub>3</sub>/gDOC, followed by samples from 0.4 gO<sub>3</sub>/gDOC (5070) (SI, Table S13). Samples from specific ozone doses of 0.7 or 1.0 gO<sub>3</sub>/gDOC had the lowest number of links, namely 3983 or 1295, respectively.

Table 1. Summary of non-target features and links detected for the four specific ozone doses and two electrospray ionization (ESI) modes. Number of influent and effluent peaks were determined based on the characterization with PCA. The number of unique potential parent or OTP features refers to unique  $m/z$  and retention time features for which a link was detected through the linkage analysis.

	Positive ESI Mode				Negative ESI Mode			
	0.4 gO <sub>3</sub> /gDOC	0.5 gO <sub>3</sub> /gDOC	0.7 gO <sub>3</sub> /gDOC	1.0 gO <sub>3</sub> /gDOC	0.4 gO <sub>3</sub> /gDOC	0.5 gO <sub>3</sub> /gDOC	0.7 gO <sub>3</sub> /gDOC	1.0 gO <sub>3</sub> /gDOC
Total number of features	4260	6207	3798	2279	1048	1719	787	270
Number of influent peaks	1065	1552	950	570	262	430	197	68
Number of effluent peaks	1065	1552	950	570	262	430	197	68
Number of unique potential parent features	979	1416	874	482	184	343	100	17
Number of unique potential OTP features	869	1328	724	394	153	324	85	12
% features as potential parents with links	23	23	23	21	18	20	13	6
% features as potential OTPs with links	20	21	19	17	15	19	11	4
Average number of links for each parent	6	7	5	3	2	4	2	2
Average number of links for each OTP	2	8	6	4	2	3	2	1

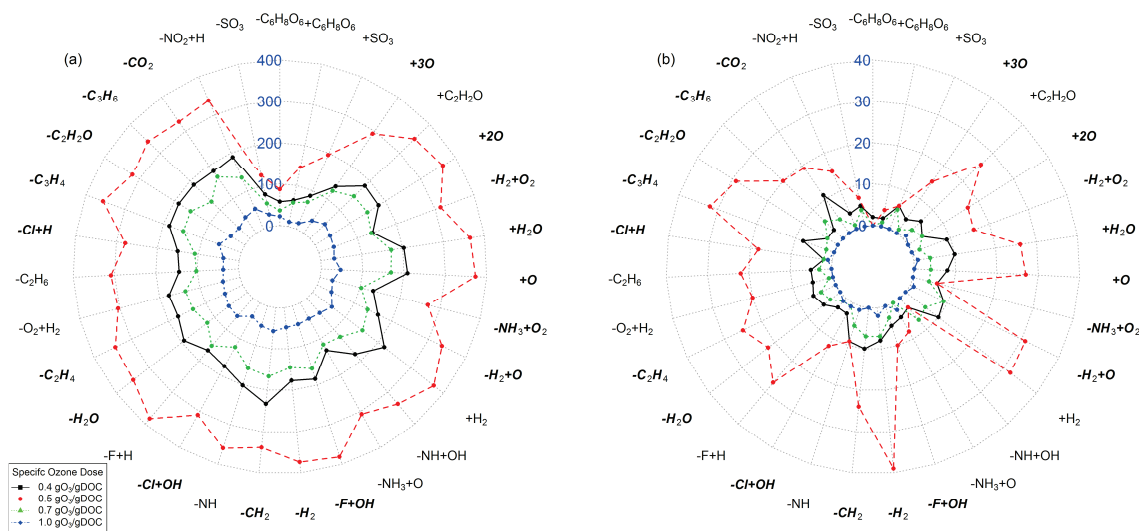


Figure 4. Radar plot of the different linkage types detected for each ozone dose in (a) positive ESI mode and (b) negative ESI mode. The scale, showing the number of times each transformation type was detected, is indicated on the grey concentric circles. The number of detections for each ozone dose is shown with the different colors and line styles. Mass differences are ordered starting at the top from the largest addition and continuing clockwise

to the largest loss. **Formula differences in bold and italic correspond to transformation types that are expected to occur either through reactions with ozone or hydroxyl radicals (further information in SI, Table S8).**

It is reasonable that the samples with the highest number of features (0.5 gO<sub>3</sub>/gDOC) also had the highest number of links, because more links are possible when more features are detected (Table 1). Furthermore, each parent compound can be linked to multiple OTPs and each OTP can be linked to multiple parents. On average, more OTPs per parent compound were found in the linkage analysis of the 0.5 gO<sub>3</sub>/gDOC samples (7 OTPs per parent compound) compared to the linkage analysis of the 1.0 gO<sub>3</sub>/gDOC samples (3 OTPs per parent compound), perhaps also because fewer potential OTPs were detected at the higher doses. Additionally, the average number of links to parent compounds for each OTP, as well as the percentage of OTPs with links, was highest at intermediate specific ozone doses (8 links/OTPs or 21% at 0.5 gO<sub>3</sub>/gDOC versus 4 links/OTP or 17% at 1.0 gO<sub>3</sub>/gDOC). Overall, in the negative ESI mode, fewer links were detected, probably because fewer features were detected. However, also in the negative ESI mode, the average number of links, the average number of OTPs/parent compound, and the percent of OTPs with links were highest for the intermediate specific ozone dose (3.3 OTPs per parent compound at 0.5 gO<sub>3</sub>/gDOC vs. 1.2 OTPs per parent compound at 1.0 gO<sub>3</sub>/gDOC).

The most frequently detected transformation types were hydrogenation (885 links), addition of a single oxygen (864 links), and demethylation (861 links). Addition of a single oxygen can be related to multiple possible reactions with either ozone or OH radicals, including hydroxylation, formation of *N*- or *S*-oxides, or formation of an aldehyde from an olefin (Merel et al. 2017, von Sonntag and von Gunten 2012). Demethylation during ozonation, particularly at heteroatoms, has also been described for different micropollutants (Benner and Ternes 2009, Kuang et al. 2013). The prevalence of hydrogenation can currently not be explained and needs further investigation. Substitution reactions involving halogens are not favorable during ozonation, and were also rarely detected in the linkage analysis. The (de)conjugation transformations (*i.e.*, (de)acetylation, (de)sulfation, and (de)glucuronidation), that were included to provide an estimate of false positive links, were in general detected the least. Between 200-400 (de)conjugation links were detected in positive ESI mode across all ozone doses, compared to 500-900 for the other linkages, demonstrating the potential for mass-

related peaks to be erroneously linked together as parents and OTPs. Second- or third-generation OTPs incorporating multiple oxygen atoms were found less than expected, probably because they are not formed under the applied ozone doses or too polar to be detected with reverse-phase LC-HRMS/MS.

### 3.1.2 Characterization of OTPs resulting from Suspect Screening

The suspect screening was performed on the potential parent compounds detected at 0.5 gO<sub>3</sub>/gDOC (see above). From 159 suspect hits, 27 compounds were confirmed with HRMS/MS spectra match to a reference standard measured in-house (confidence level 1, SI, Table S14). Almost all 27 compounds have been described as persistent in CAS treatment, thereby making their presence in the ozone reactor and consequently the possibility of OTP formation plausible. In the linkage analysis, 182 potential OTPs were detected for these 27 parent compounds, an average of 6 potential OTPs per parent compound (SI, Table S15). Of the 182 potential OTPs, 69 were determined to be structurally possible by comparing the parent structure and proposed transformation, e.g., deethylation only possible for parent compounds with an ethyl group. Of these 69 structurally possible OTPs, 30 were predicted as possible ozonation products by a recently developed ozone pathway prediction system (O<sub>3</sub>-PPS) (SI, Table S15).

HRMS/MS spectra were triggered in data-dependent mode for 22 of the 69 structurally possible OTPs, and the structures of three OTPs were confirmed with reference standard matches (level 1). Hydrochlorothiazide was linked to chlorothiazide through a dehydrogenation (Table S16, TP structure 7), 2-aminobenzothiazole was linked to benzothiazole through a deamination reaction (Table S16, TP structure 1), and venlafaxine was linked to venlafaxine *N*-oxide (Table S16, TP structure 22). Chlorothiazide was recently found as the main OTP from hydrochlorothiazide oxidation (Borowska et al. 2016). Similarly, the formation of venlafaxine *N*-oxide during ozonation has been documented (Bourgin et al. 2018, Lajeunesse et al. 2013, Merel et al. 2017). The link between 2-aminobenzothiazole and benzothiazole would need to be confirmed through batch experiments. However, deamination on the heteroaromatic ring is unlikely during ozonation. For the other 19 potential OTPs with HRMS/MS spectra, probable structures (*i.e.*, level 3) were suggested for 13 of them, based on the structure of the parent compound and proposed transformation (SI, Table S16). Five of the suggested OTPs were also predicted by the software O<sub>3</sub>-PPS, and one OTP (oxidative

replacement of fluorine in citalopram; OTP structure 3) has been described previously in literature (Hörsing et al. 2012). HRMS/MS spectra were not informative enough for structure confirmation of the proposed OTPs, but in most pairs, two or more fragments were shared between parent compound and proposed OTP, possibly indicating related structures (Schollée et al. 2017).

For the 47 proposed OTPs where no HRMS/MS spectra were collected, 23 were also predicted by the software O<sub>3</sub>-PPS. Additionally, information about degradation during ozonation could be found in the literature for many of the parent micropollutants and circumstantial evidence was found for several of the proposed OTPs, as indicated in SI, Table S15. For instance, for 5-methyl-benzotriazole, an addition of two oxygen atoms was found to be in agreement with Müller *et al.* (2012). The structurally possible OTP suggested for DEET (*i.e.*, a deethylation) has also been reported (Tay *et al.* 2009). Two proposed OTPs (both oxygen additions) for phenazone were detected by Miao et al. (2015), two OTPs for metoprolol (*i.e.*, oxygen addition and dealkylation) were previously reported (Tay *et al.* 2012), and one OTP for trimethoprim was previously detected by Kuang *et al.* (2013). An interesting example presented itself in the case of atenolol. Molecular formulas of three structurally likely OTPs for atenolol corresponded to 7 different structures which were reported by Tay *et al.* (2011) since one formula was related to multiple OTPs (*e.g.*, oxygen addition at multiple locations on the molecule). Concerning the OTPs reported in the literature that were not detected here, the majority of these were resulting from transformations not predicted by the linkage analysis and can be incorporated in the future. For the OTPs, which were detected here but not reported in literature, most studies on micropollutant abatement during ozonation do not include identification of OTPs or only identify OTPs resulting from reaction with ozone but not with OH radicals, therefore, not all OTPs which might be formed during ozonation at a full-scale WWTP are known. Specifically, very little information is available on the fate of biological TPs from CAS treatment in ozonation, although a number of them (*i.e.*, desethyl-atrazine and *N*-desmethylvenlafaxine) were found to be possible parent compounds in the ozone analysis, highlighting the need to study their behavior and potential for OTP formation. Nevertheless, the numerous successful examples demonstrate that the combined PCA and linkage analysis approach to analyze full-scale WWTPs lead to the detection of OTPs of known micropollutants that had previously only been described in batch experiments.



For four of the 27 identified parent compounds (desethyl-atrazine, 4-acetamidoantipyrin (4-AAA), carbendazim, and irbesartan), no structurally possible OTPs were suggested. Low reactivity with ozone has been reported for two of these, desethyl-atrazine ( $k_{O_3}$  0.19 M<sup>-1</sup> s<sup>-1</sup> at pH 7 (Beltrán et al. 1998)), and irbesartan ( $k_{O_3}$  24 M<sup>-1</sup> s<sup>-1</sup> (Bourgin et al. 2018)). No second order rate constants for the reaction of ozone with 4-AAA or carbendazim are available in the literature. For compounds with low reactivity with ozone, the reaction with hydroxyl radicals dominates during ozonation of WWTP effluent water. Since reactions with hydroxyl radicals are less specific than those with ozone, the resulting transformations may not have been included in the linkage analysis and the resulting OTPs therefore not been found. Additionally, the following reasons can be suggested for why OTPs were not detected for these parent compounds: (i) it may be the case that first generation OTPs react further and the resulting second generation products were not predicted with the linkage analysis; (ii) it is possible that yields of OTPs were too low and therefore below the detection limit of the instrument; or (iii) that the OTPs were not detected with online-SPE LC-HRMS/MS (e.g., too polar to be retained in the SPE cartridge or on the reverse-phase LC column, or not ionized in the MS). In the future, incorporation of an additional LC method such as HILIC to detect highly polar OTPs would be advantageous.

### 3.2 Analysis of Post-Treatment Sampling

#### 3.2.1 Non-target Peak Characterization including OTPs

Non-target screening was used to compare five post-treatment technologies (Figure 1). No large differences were observed among the number of features detected in the effluents of the five post-treatment step-ups (SI, Table S17). Approximately 3000-4000 features were detected in the effluent of each post-treatment, which is a slight reduction from the number of features in the effluent of the ozone reactor (4340) and in the effluent of the CAS treatment (5016). The characteristics of non-target features (*i.e.*,  $m/z$  and RT) were significantly lower after the CAS treatment (Tukey's HSD,  $p < 0.01$ , SI, Figure S9). The  $m/z$  values of non-target features in the GAC filters were significantly lower than those detected in the other steps of the WWTP (Tukey's HSD,  $p < 0.01$ ), likely because larger non-polar compounds are expected to be removed by sorption in the GAC filters.

Of the 69 structurally possible OTPs detected from the analysis of the specific ozone dose of 0.5 gO<sub>3</sub>/gDOC (section 3.1.2), 67 were also detected in the second set of samples analyzing the different post-treatments. No substantial differences between the intensity detected after ozonation and

intensity after post-treatment were observed, suggesting little to minor removal of these potential OTPs; however, reference standards would be needed for proper quantification.

### 3.2.2 Hierarchical Clustering across the WWTP

To further investigate the behavior of non-target features during wastewater treatment, hierarchical clustering was applied to group features including OTPs into clusters based on intensity profiles across the WWTP. For each feature, intensity profiles were constructed from the maximum intensities detected in the four sample types, *i.e.*, before CAS treatment, after CAS treatment, after ozonation, and after post-treatment.

The HCA was validated with a suspect screening of eight pharmaceuticals (amisulpride, carbamazepine, cetirizine, clarithromycin, lamotrigine, ranitidine, tramadol, and venlafaxine) and 12 corresponding OTPs in the WWTP with sand filtration as post-treatment. The correct trend was assigned to five of eight parent compounds and nine of twelve OTPs (SI, Table S18). For Of the falsely assigned parent compounds, two (tramadol and ranitidine) were incorrectly grouped during pre-processing and one (clarithromycin) was detected in the blind and therefore removed during pre-processing. For the three falsely assigned OTPs, two were not assigned to the expected trend because of their low intensities ( $<2E6$  for amisulpride *N*-oxide and cetirizine *N*-oxide) and one (clarithromycin *N*-oxide) because of large retention time shifts over the samples which were not completely corrected for with the RT alignment. Four OTPs were not detected in either the target or non-target screening, and were considered as true negatives (and as correctly classified) in the validation.

Clustering and trend assignment was done separately for each of the five post-treatment set-ups. The normalized clusters resulting from the HCA of each post-treatment are shown in SI, Figure S12. It is expected that the number of features with Trend 1 or Trend 2 would not change among the five set-ups, because the CAS treatment and ozonation were always the same although slight variations in the total number of features may occur as a result of the data pre-processing or the HCA. Table 2 illustrates that the number of features assigned to Trend 1 or Trend 2 indeed did not vary substantially (RSD 1.4% for Trend 1 and 5.0% for Trend 2). In all cases the largest number of features (59-61%) were removed during the CAS treatment (*i.e.*, Trend 1).



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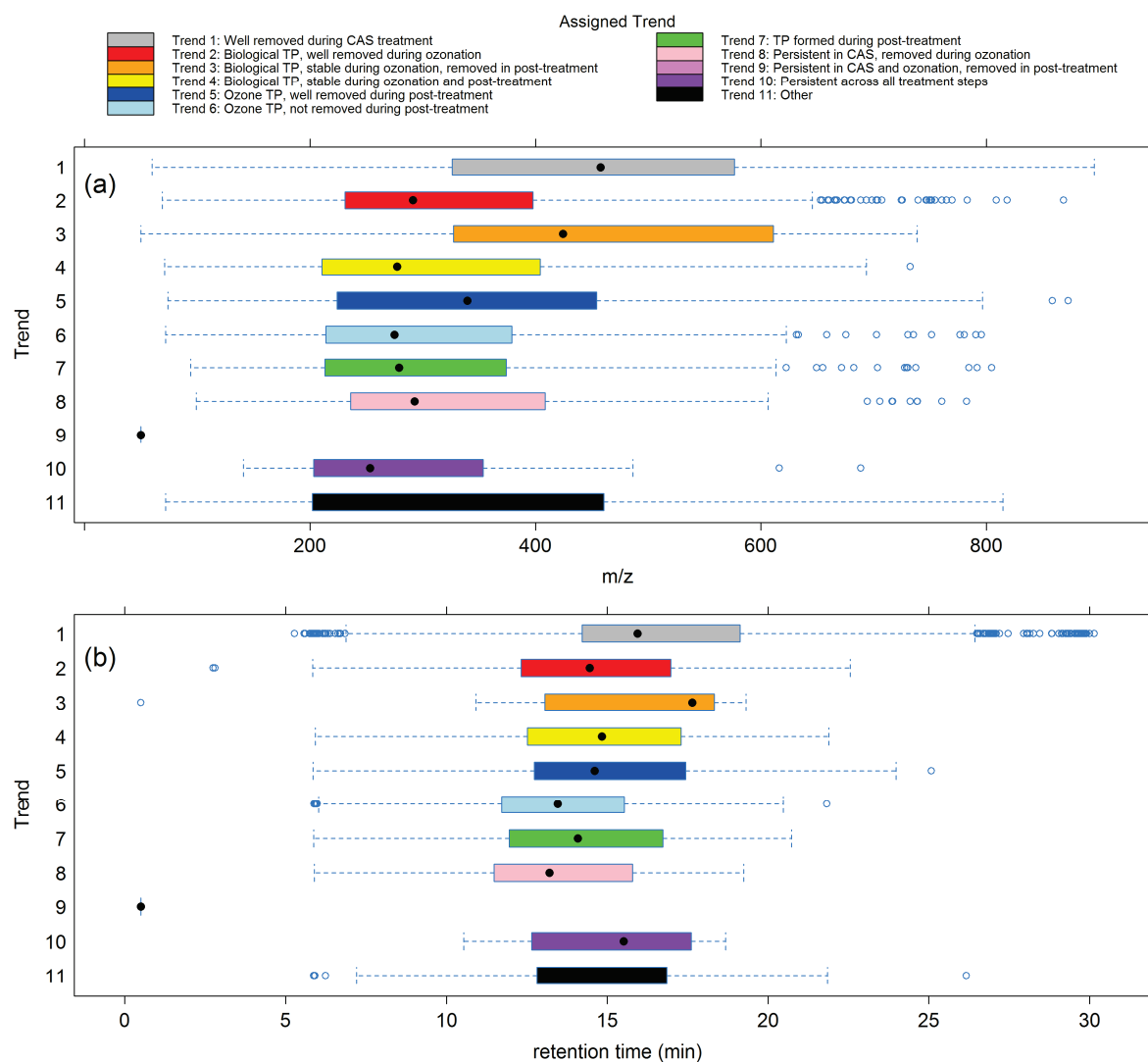
**Table 2. Summary of the number / percentage of non-target features removed or formed in each of the described trends in the five post-treatment set-ups. CAS treatment refers to conventional activated sludge treatment. GAC-A was relatively fresh (14,000–24,000 BV) and GAC-B pre-loaded (36,000–46,000 BV).**

<i>Trend</i>	<i>Intensity Profile Behavior</i>	<i>GAC-A</i>	<i>GAC-B</i>	<i>Moving bed</i>	<i>Fixed bed</i>	<i>Sand filter</i>	<i>RSD</i>
1	Well removed during CAS treatment	4890 / 61%	4758 / 59%	4866 / 61%	4804 / 60%	4903 / 61%	1.4%
2	Biological TP, well removed during ozonation	1442 / 18%	1304 / 16%	1395 / 17%	1302 / 16%	1419 / 18%	5.0%
3	Biological TP, stable during ozonation, removed in post-treatment	155 / 2%	125 / 2%	22 / 0.3%	26 / 0.3%	15 / 0.2%	97%
4	Biological TP, stable during ozonation and post-treatment	35 / 0.4%	200 / 3%	230 / 3%	304 / 4%	236 / 3%	50%
5	Ozone TP, well removed during post-treatment	597 / 8%	452 / 6%	385 / 5%	393 / 5%	374 / 5%	21%
6	Ozone TP, not removed during post-treatment	122 / 2%	231 / 3%	327 / 4%	305 / 4%	312 / 4%	33%
7	TP formed during post-treatment	269 / 3%	276 / 3%	303 / 4%	298 / 4%	266 / 3%	5.7%
8	Persistent in CAS, removed during ozonation	184 / 2%	123 / 2%	105 / 1%	181 / 2%	152 / 2%	23%
9	Persistent in CAS and ozonation, removed in post-treatment	0 / 0%	0 / 0%	0 / 0%	0 / 0%	0 / 0%	0.0%
10	Persistent across all treatment steps	13 / 0.2%	11 / 0.1%	45 / 0.6%	30 / 0.4%	39 / 0.5%	55%
11	Other	283 / 4%	524 / 7%	361 / 5%	394 / 5%	268 / 3%	28%
	Total number of features detected	7990	8004	8039	8037	7984	0.3%

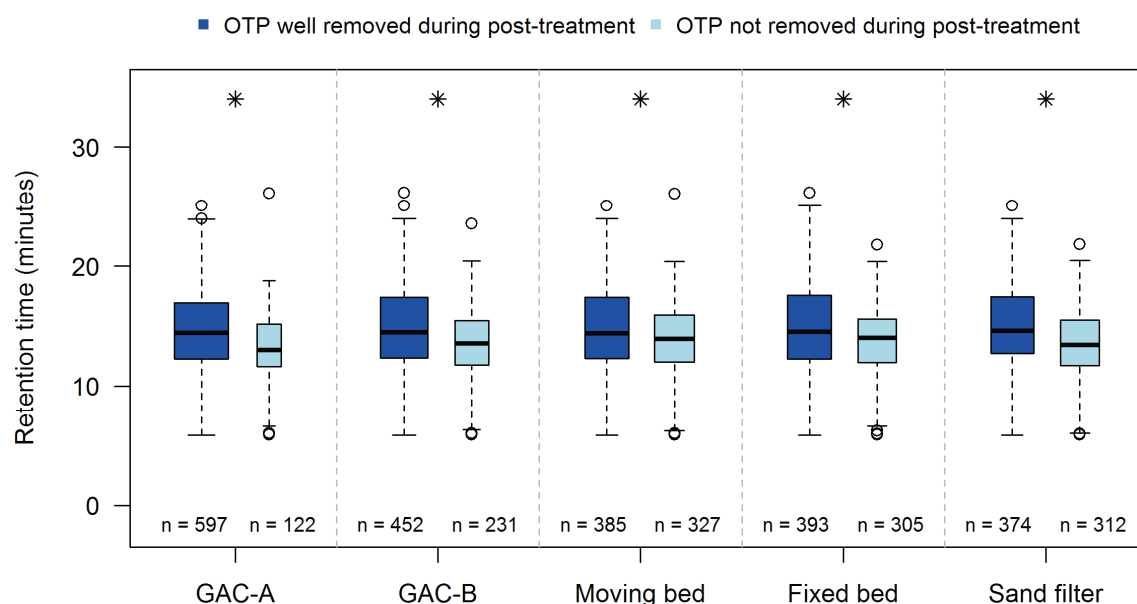
The number of features in the Trends 3 to 11 show differences in the post-treatments. Features with Trend 5 represent compounds that are likely formed during ozonation, and are subsequently removed in the post-treatment. The sand filter, moving bed, and fixed bed post-treatments each had relatively low percentages of features with this trend (5%), indicating possibly low removal of OTPs. The two GAC filters had slightly more features with this trend (8% in GAC-A and 6% in GAC-B). In addition to biological degradation, sorption can also occur on the GAC filters, possibly accounting for the increased removal potential in the GAC filters. The better performance of GAC-A compared to GAC-B may be explained by the fact that GAC-A was only run to approximately half of the BVs and therefore exhibits a higher sorption capacity, which was also shown in target analysis (Bourgin et al. 2018). DOC abatement was also higher in the two GAC filters (15-20%) compared to the fixed bed and moving bed reactors (5 and 6%, respectively). Conversely, the number of features assigned to Trend 6 (likely OTPs not removed in post-treatment) was highest in the sand filter and two bioreactors (4%), and lowest in the two GAC filters (2-3%). Only a minor portion of features detected appeared to be

formed during ozonation (Trend 5+6; -9-10%), and from these, 54% to 83% were removed in post-treatment, with highest removal efficiencies being observed in the two GAC filters. Other research based on target micropollutants has also shown that GAC may be more efficient at removing OTPs compared to solely biological filters (Knopp et al. 2016) but further systematic investigations would be needed to assess the fate of OTPs in GAC filters, e.g., over a longer run time. It has to be noted that many of the clustered observed peaks formed during ozonation arise from the reaction of ozone with matrix components and not with micropollutants and need to be further evaluated. Therefore, further evaluation is necessary to determine the efficiency of post-treatments on OTP removal.

The characteristics of non-target features in each trend type were compared for each post-treatment analysis (shown for the sand filter in Figure 5 and SI, Figure S13, and the other post-treatments in the SI, Figures S14-S17) to determine if specific feature characteristics were indicative of, e.g., better biodegradability. In almost all cases there was no significant difference (Tukey's HSD,  $p < 0.05$ ) in mass defect of non-target features in each of the 11 trends (SI, Figures S18-S22), indicating that, e.g., chlorinated compounds with negative mass defect were not less prone to transformation. In all five post-treatments, OTPs that were removed had significantly higher RTs ( $p < 0.05$ , Mann Whitney U test) than OTPs which were not well removed, indicating possible recalcitrance of some polar OTPs (Figure 6).



**Figure 5. Box-whisker plots of the (a)  $m/z$  and (b) retention time in each of the 11 defined trends with sand filtration as post-treatment step for positive ESI. Colors represent the trends, also indicated on the y-axis.**



**Figure 6. Box-whisker plots of the retention times of OTPs removed or not removed during five post-treatment steps, measured in positive ESI. OTPs removed were features classified with Trend 5 and OTPs not removed were features classified with Trend 6 in the cluster analysis. The number of features (n) in each box is also represented by the width of the box (i.e., large box indicates more features). Within one post-treatment sample set-up, the retention times of OTPs in Trend 5 and Trend 6 were in all cases significant (Mann Whitney U test,  $p < 0.05$ ), as indicated by a star.**

The clusters with trends relevant to OTP formation can be compared to the potential OTPs from the linkage analysis to prioritize possible OTP peaks for future structure elucidation. From the linkage analysis with an specific ozone dose of 0.5 gO<sub>3</sub>/gDOC, 1328 unique non-target features were suggested as potential OTPs (SI, Table S12). With the HCA, 686 non-target features were assigned either to Trend 5 or Trend 6 using the sand filter post-treatment, indicating possible formation during ozonation. With these two prioritization methods, 131 non-target features overlapped using a 15 ppm  $m/z$  tolerance and 1 minute RT window. While this overlap may appear to be quite small, wastewater inputs vary considerably, as can treatment efficiency. Taking into account that the samples were collected in some cases more than six months apart, these features may represent relevant OTPs that warrant further attention due to their frequent detection.

## 4.1 Conclusions

- The combination of PCA and chemical logic (*i.e.*, linkage analysis) made it possible to gain an overview of the types of non-target peaks present in wastewater effluent treated with different ozone doses. Results suggested that treatment with a specific ozone dose of 0.5 gO<sub>3</sub>/gDOC led to the formation of the highest number of observable OTPs (1552), supporting previous findings, although other enrichment or chromatography methods may be necessary to detect highly polar OTPs.
- Chemical logic was able to suggest 69 structurally plausible OTPs originating from 27 parent compounds, and three OTPs could be confirmed by reference standards. A number of the suggested OTPs had previously been detected in ozonation batch experiments and some were also predicted to be likely by mechanistic information using an ozone pathway prediction system (O<sub>3</sub>-PPS), providing further evidence of their presence in the full-scale WWTPs.
- The presented workflow, combining HRMS data with statistical methods such as PCA, chemical logic and mechanistic information, provides an efficient approach to select and prioritize potential parent compounds for further studies. The ability to monitor OTPs in a full-scale WWTP is critical, because it is not possible to conduct batch experiments on all chemicals. Intensities of the potential OTPs in the effluent of five post-treatment technologies suggest incomplete removal, although further studies are needed for quantification.
- From hierarchical cluster analysis across all treatment steps, a small portion of the non-target features (9%) was found to be formed during ozonation and 54% to 83% of these features were removed during post-treatment. Compared to the other post-treatment options, the two GAC filters evaluated appeared to perform slightly better in removing non-target features formed during ozonation.

## 5.1 Acknowledgements

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### *Appendix A – Supplementary data*

Supplementary data is available as indicated in the text.

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## 1 Highlights

- 2 • Non-target screening with HRMS was applied to a wastewater treatment train
- 3 • Statistics were used to prioritize potential ozonation transformation products OTPs
- 4 • 69 plausible OTPs were found; 34 agreed with literature data or reaction mechanisms
- 5 • 54 to 83% of potential OTPs were removed in the five post-treatments

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