

Supporting Information

Formation of transformation products during ozonation of secondary wastewater effluent and their fate in post-treatment: From laboratory- to full-scale

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Additionally to this SI pdf, an excel file with tree tables and a second pdf file are attached, which include the following information:

Excel Table S1: Information on the studied 87 parent micropollutants

Excel Table S2: OTP signals (777 in positive and 972 in negative ionization modes) of *O3bMix* experiments for 70 parent micropollutants (found in Approach 1)

Excel Table S3: OTP signals (totally 84) found in wastewater samples of four WWTPs which were matching signals in the *O3bAll* samples (Approach 2)

Figure S19: Figures with all 84 OTPs found in batch experiments and in wastewater treatment.

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Text S1. Micropollutant selection and separation into mixtures

Details on all parent micropollutants (MPs), including chemical structures, are provided in Table S1 in the supporting Excel file.

The MPs were spiked into 19 different mixtures, whereby each MP was present in two mixtures, while no other MP occurred twice in these two mixtures. The separation is visualized in Figure S1 and Figure S2. Figures S1 and S2 represent a 9 to 10 matrix, where each position is taken up by a MP. Each row was mixed together and these 9 mixtures were called MR0 – MR8. Accordingly, each column was mixed together and these 10 mixtures were called MC0 – MC9.

	MC0	MC1	MC2	MC3	MC4	MC5	MC6	MC7	MC8	MC9
MR0	Venlafaxine	Lidocaine	Flecainide	Atenolol-desisopropyl	Eprosartan	Lamotrigine	Sulfamethazine	Ketoprofen	Levamisole	Sucralose
MR1	Diclofenac	Mecoprop	Citalopram	Cetirizine	Propranolol	Sitagliptin	-	Prometon	Sulfapyridin	Metformin
MR2	Emtricitabine	Caffeine	Thiacloprid	Fenpropidin	Diphenhydramine	Norlidocaine	Mycophenolic acid	Sulfamethoxazole	2-Naphthalic sulfonic acid	Flufenamic acid
MR3	5-Methyl-benzotriazole	Cyprodinil	Atazanavir	Diuron	Norfenfluramine	Clindamycin	Atenolol acid	Amitriptyline	Rosuvastatin	2-7-naphthalic disulfonic acid
MR4	Etodolac	Benzotriazole	Trimethoprim	Methoxyfenozide	Ibuprofen	Tramadol	Clozapine	Atenolol	Oseltamivir	Hydrochlorothiazide
MR5	Acesulfame	Carbamazepine	Candesartan	Benzisothiazolone	Benzophenon 3	Bezafibrate	N-Bisdesmet. Tramadol	Sulpiride	Metoprolol	Pravastatin
MR6	Aliskiren	Cephalexin	Resveratrol	Valsartan acid	-	2,4-D	Phenazone	Clarithromycin	Amisulpride	N-Desmethyl-tramadol
MR7	Efavirenz	-	Progesterone	Fenfluramine	Valsartan	Naproxen	Atrazine	Oxcarbazepine	Diltiazem	Triclosan
MR8	Levetiracetam	Propyzamide	Gabapentin	Methyl-prednisolone	Codeine	Irbesartan	Napropamide	Oxazepam	Mefenamic acid	Ranitidine
	3° amines	2° amines	1° amines	olefins	ethynes	sulfonamide	aromatic compounds	Similar to benzotriazole	naphthalenes	N-hetero aromatics

Figure S1: Visualization of the micropollutant (MP) mixtures for the ozonation batch experiments (O3bMix). All MPs of a row were mixed together (called MR0 – MR8) and all MPs of a column were mixed together (called MC0 - MC9). The background colors indicate the ozone reactive functional groups of the micropollutants.

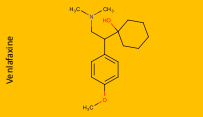
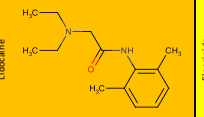
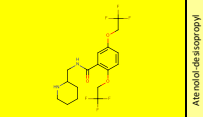
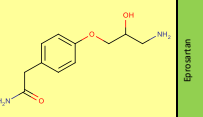
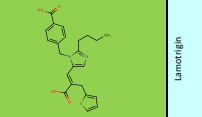
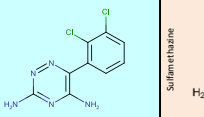
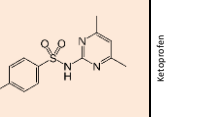
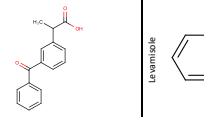
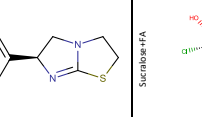
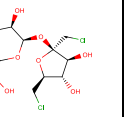
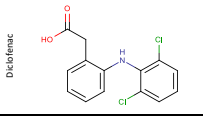
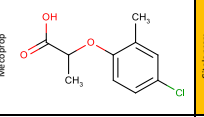
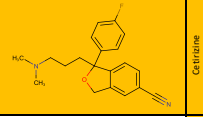
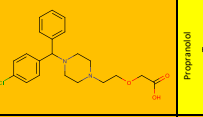
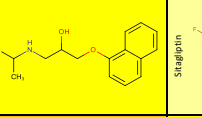
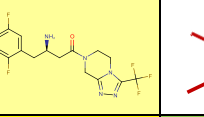
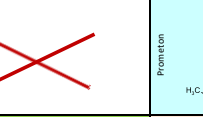
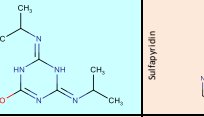
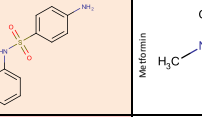
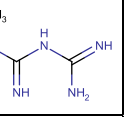
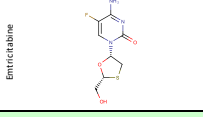
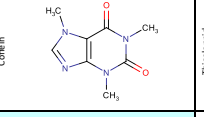
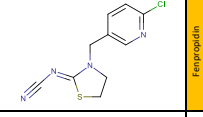
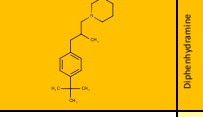
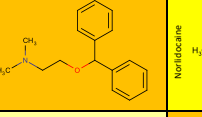
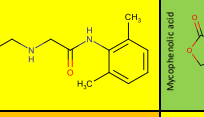
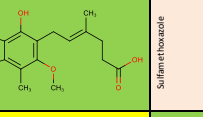
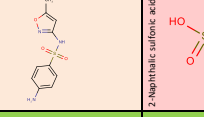

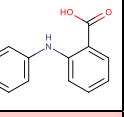
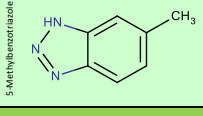
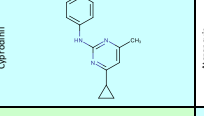
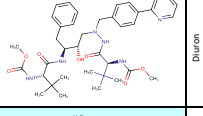
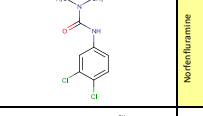
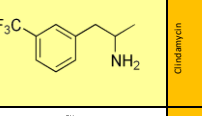

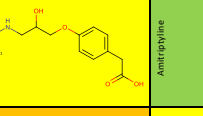
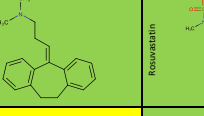
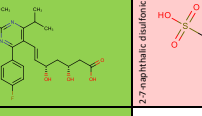
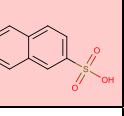
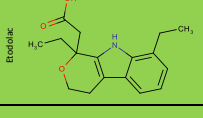
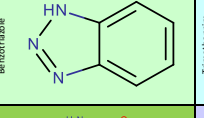
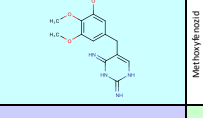
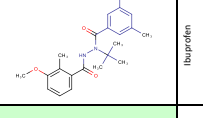
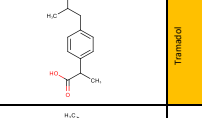
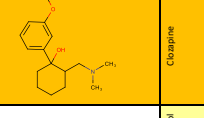
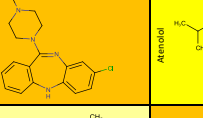
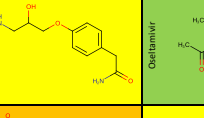
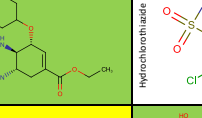
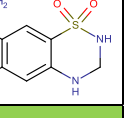
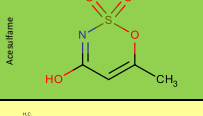
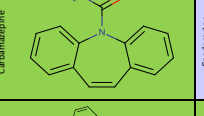
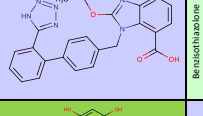
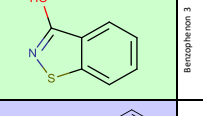
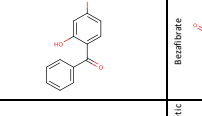
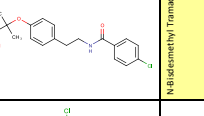
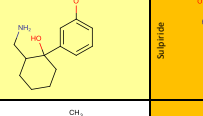

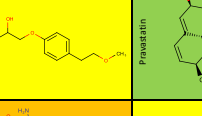
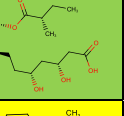
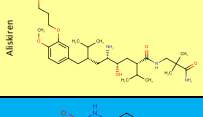
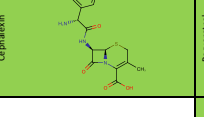
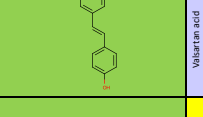
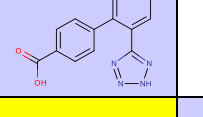
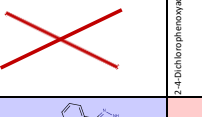
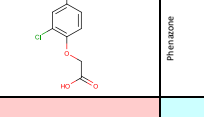
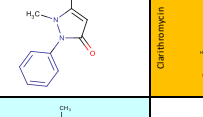
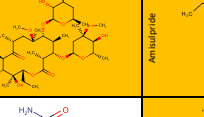
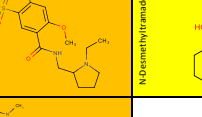
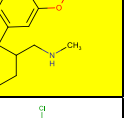
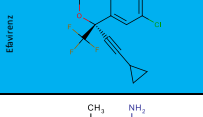
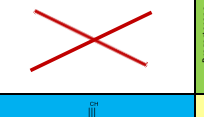
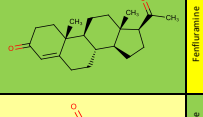
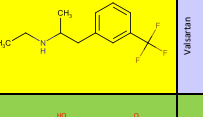
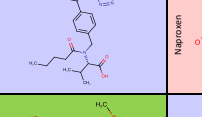
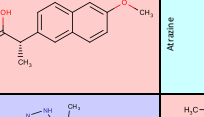
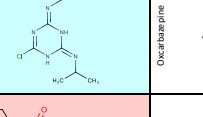
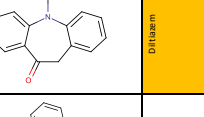
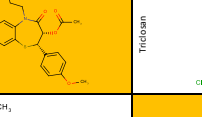
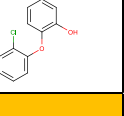
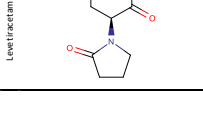
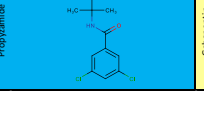
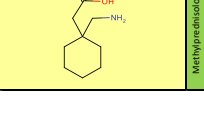
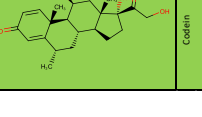
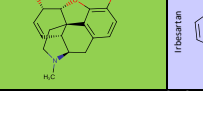
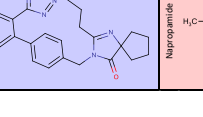
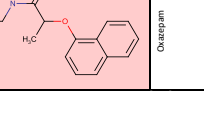
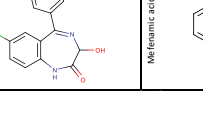
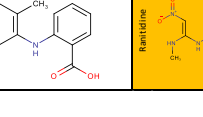
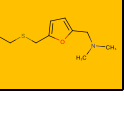
	MC0	MC1	MC2	MC3	MC4	MC5	MC6	MC7	MC8	MC9
MR0										
MR1										
MR2										
MR3										
MR4										
MR5										
MR6										
MR7										
MR8										

Figure S2: Visualization of the micropollutant (MP) mixtures for the ozonation batch experiments (O3bMix). All MPs of a row were mixed together (called MR0 – MR8) and all MPs of a column were mixed together (called MC0 - MC9). The background colors indicate the ozone reactive functional groups of the micropollutants. Color coding as in Figure S1: 3° amines (orange), 2° amines (yellow), 1° amines (light yellow), olefins (green), ethynes (blue), sulfonamide (light pink), aromatic compounds (white), similar to benzotriazole (light green), naphthalenes (pink), N-hetero aromatics - triazines (light blue), sartanes (purple).

Text S2. Laboratory ozonation batch experiments

Text S2.1 Simulation of organic matter containing water matrices and ozonation conditions

Before conducting the ozonation batch experiments with micropollutants, two water matrices with a DOC content of 1.3 and 5 were simulated by mimicking the organic matter with methanol and acetate. Methanol acts as promoter for the radical chain reaction with hydroxyl radicals ($\bullet\text{OH}$), which enhances the rate of ozone decomposition, while acetate is an inhibitor, which stops the radical chain reaction. Therefore, methanol and acetate represent sites in the dissolved organic matter (DOM), which control the ozone stability (Buffle et al. 2006, Elovitz and von Gunten 1999, Hoigné and Bader 1994, Staehelin and Hoigne 1985). The aim was to yield similar ozonation conditions as in real waters, i.e. ozone- and $\bullet\text{OH}$ exposure (and the concentration ratio of $\bullet\text{OH}/\text{O}_3$ known as R_{ct}), but with less interferences than DOM in the LC-HR-MS/MS screening for OTP signals. Details on the composition of the simulated water matrices as well as on the resulting ozonation conditions in the pre-experiments are provided in Table S4. Experiments were performed in a phosphate buffer at pH 7.5 (1-5 mM depending on the experiment), which does not affect the oxidation by ozone as long as the pH remains constant. Figure S3 shows the evolution of the ozone concentration and the R_{ct} values for various conditions. The specific ozone dose was chosen depending on the simulated DOC content to be 1 gO_3/gDOC and therefore the absolute ozone doses differed among the two water matrices. As can be seen from Table S4 and Figure S3, also the duration of the experiments differed, whereas the ozone exposure (0.020 - 0.035 Ms) at full ozone depletion and the R_{ct} values (1.3×10^{-8} - 2.4×10^{-8}) were in a similar range. By taking samples at different time points and quenching the ozone with indigo or sulfite, different ozonation conditions can be selected to best mimic real water conditions. The ozone exposures of 9 different municipal wastewaters were determined to be in the range of 0.0002-0.0028 Ms for a specific ozone dose of 0.5 gO_3/gDOC , and 0.004-0.013 Ms for a specific ozone dose of 1 gO_3/gDOC (Lee et al. 2013). For 5 lakes and 2 river waters, Elovitz et al. (2000) determined ozone exposures in the range of 0.0015-0.024 Ms for specific ozone doses of 0.67-1.0 gO_3/gDOC . The R_{ct} values for the seven surface waters were in the range of 0.95×10^{-8} - 5.8×10^{-8} and for the nine wastewaters in the range of 5.0×10^{-8} - 19×10^{-8} (for 0.5 gO_3/gDOC) and of 2.2×10^{-8} - 6.3×10^{-8} (for 1.0 gO_3/gDOC). By time-resolved experiments any water matrix can be simulated in terms of ozone and hydroxyl radical exposure and the R_{ct} values. This enables to carry out product studies under realistic ozonation conditions in absence of a cumbersome matrix, which may cause analytical problems.

Table S4 shows the water matrices simulating DOC concentrations of 1.3 mg/L and 5 mg/L which were used for the ozonation batch experiments with micropollutants (see Text S2.2 and S2.3).

The simulated DOC concentrations refer to a lake water and a wastewater effluent in Switzerland, respectively. The scavenging rates for the different water constituents are also shown in Table S4. They were calculated by the product of the second-order rate constants of the reactions with hydroxyl radicals and the concentration of the scavengers. For the simulation of the 1.3 mg/L DOC matrix, the relative scavenging rates of acetate and methanol are 30% and 70%, respectively. For 5 mg/L DOC matrix, the relative scavenging rates of acetate and methanol are 5% and 95%, respectively.

Table S4: Details on the simulated water matrices without addition of micropollutants.

Simulated DOC concentration in mg/L (scavenging rate) $k_{\text{DOC},\text{OH}} = 3 \times 10^4 \text{ L/mg/s}^a$	Acetate in mM (scavenging rate) $k_{\text{ac},\text{OH}} = 7.9 \times 10^7 \text{ M}^{-1}\text{s}^{-1}$ ^b	MeOH in mM (scavenging rate) $K_{\text{MeOH},\text{OH}} = 9.7 \times 10^8 \text{ M}^{-1}\text{s}^{-1}$ ^b	pH (Phosphate buffer 1-5 mM)	Targeted Ozone dose in mg/L	Ozone exposure in M·s (full-depletion)	R_{ct}
1.3 ($3.9 \times 10^4 \text{ s}^{-1}$)	0.15 ($1.19 \times 10^4 \text{ s}^{-1}$)	0.029 ($2.8 \times 10^4 \text{ s}^{-1}$)	7.5	1.3	0.035	1.3×10^{-8}
5 ($1.5 \times 10^5 \text{ s}^{-1}$)	0.10 ($7.9 \times 10^3 \text{ s}^{-1}$)	0.16 ($1.55 \times 10^5 \text{ s}^{-1}$)	7.5	5	0.020	2.4×10^{-8}

^{a)} (von Sonntag and von Gunten 2012), ^{b)} (Elovitz and von Gunten 1999)

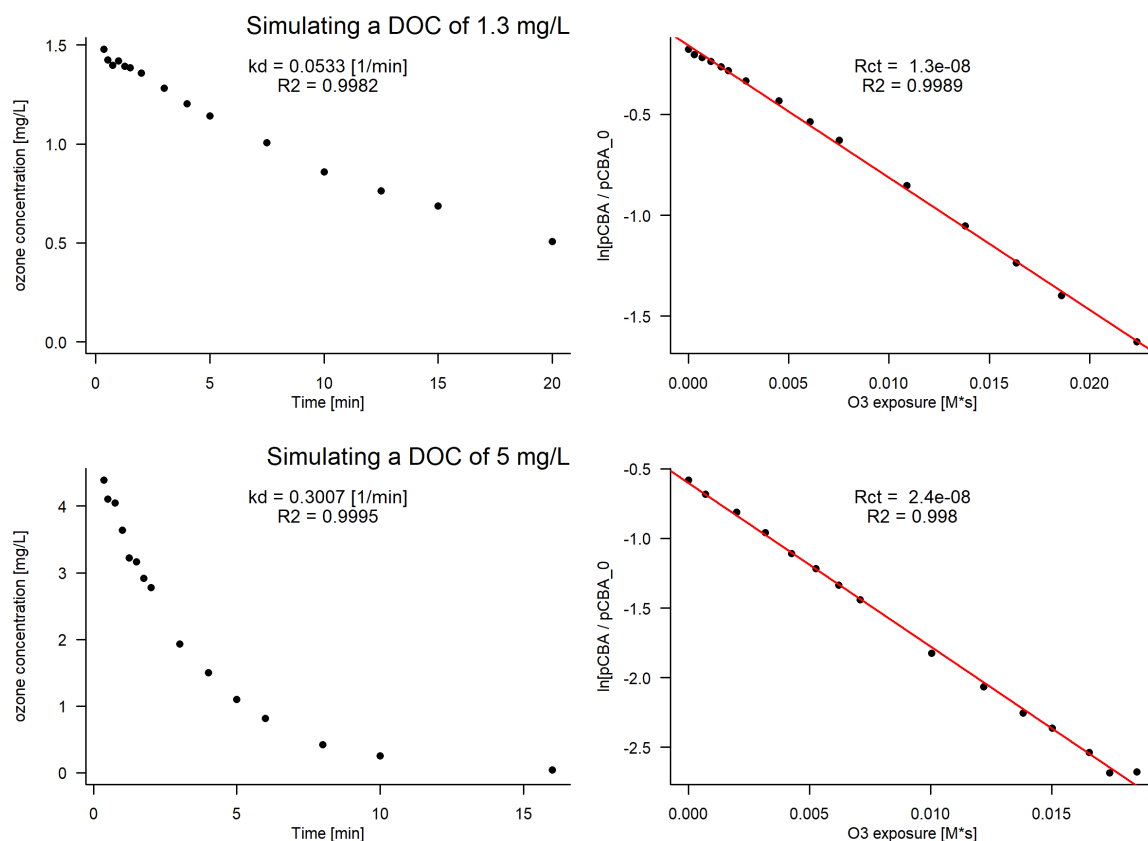


Figure S3: Ozonation conditions for the two different simulated water matrices without addition of micropollutants. Left: plots of ozone concentrations as a function of the reaction time. Right: plots of $\ln(pCBA/pCBA_0)$ as a function of the ozone exposure, the R_{ct} can be calculated from the slopes and is provided in the graphs.

Text S2.2 Ozonation conditions in the *O3bMix* experiments

In Figure S4 - Figure S8 the ozonation conditions of the 19 *O3bMix* experiments (spiked with 19 different MP mixtures as described in Section 2.1, main manuscript) are illustrated. These experiments were performed with 0.15 mM acetate and 0.029 mM methanol, simulating a DOC of 1.3 mg/L. The resulting average ozone exposure for complete ozone depletion was 0.011 ± 0.003 Ms and the estimated R_{ct} from the two-point fit of the 19 experiments (note that ozone quenching with sulfite did not work in the *O3bMix* samples) was $(2.3 \pm 0.7) \times 10^{-8}$. These experiments had a lower ozone exposure than expected from the experiments defining the simulation conditions (0.035 Ms) described above. The reason might be the different handling due to the addition of MPs. Nevertheless, the ozone exposure of the *O3bMix* experiments at full depletion of 0.011 ± 0.003 Ms fitted better to conditions with real water matrices (wastewaters and surface waters, see previous section) compared to the conditions of the preliminary experiments (Table S4).

These experiments were performed also with another matrix (0.10 mM acetate and 0.16 mM methanol, simulating a DOC of 5 mg/L, average ozone exposure at full ozone consumption was 0.016 ± 0.002 Ms (data not shown). However, the ozone exposure of this matrix was higher than the values for real wastewater effluents and is therefore less representative for wastewater than the matrix simulating a DOC of 1.3 mg/L. Therefore, it is reasonable that only few additional OTP signals could be identified in the examined WWTP samples from the evaluation of this matrix. Based on these observations, we decided to focus our evaluation and presentation of the results only on the 1.3 mg/L DOC matrix. However, due to failure of one reactor in the first matrix (MC8), the MPs 2NS, ASP, MTO, OSE, SPD were evaluated in the other matrix (simulated DOC 5.0 mg/L).

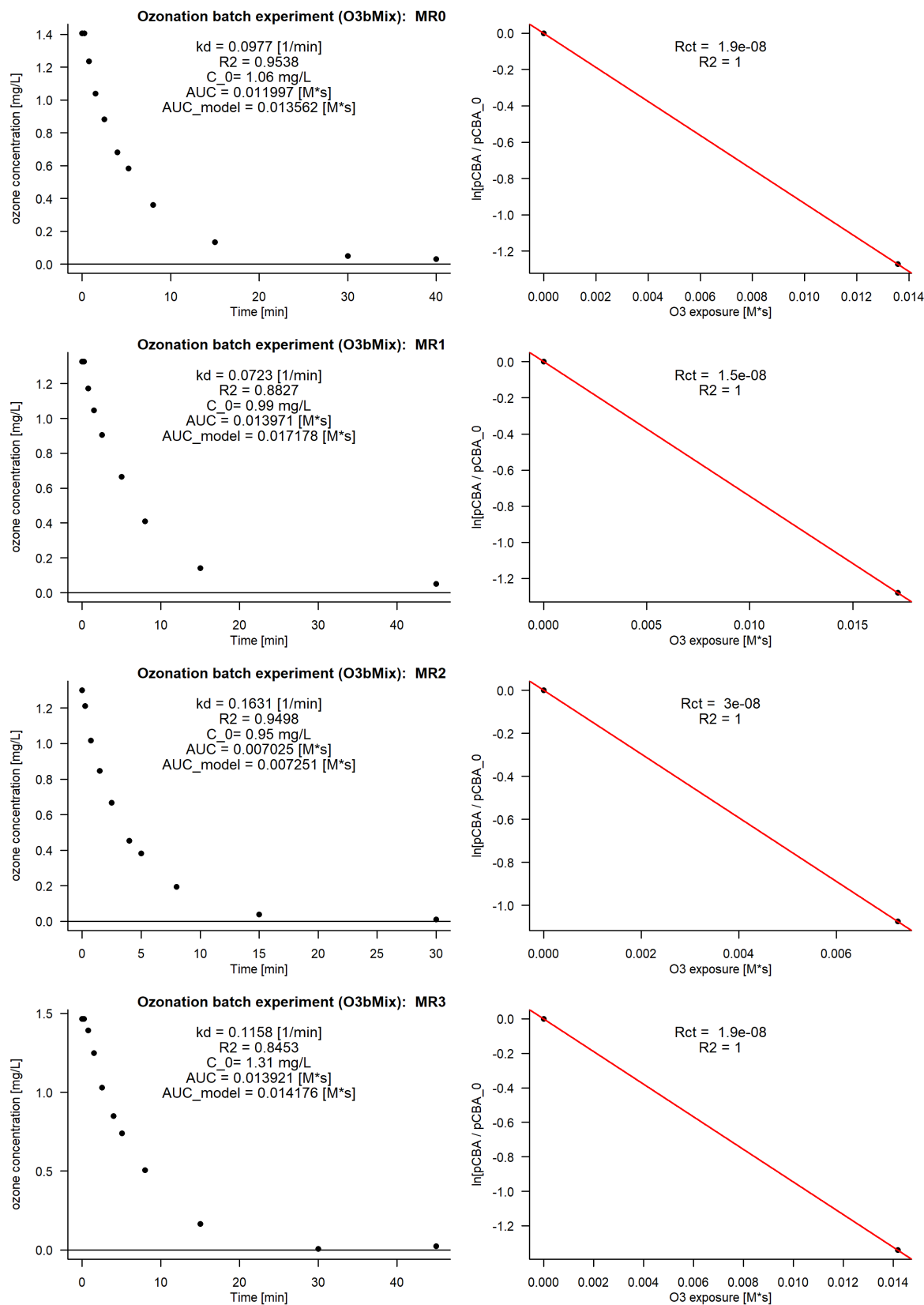


Figure S4: Ozonation conditions of O3bMix experiments with mixtures MR0-MR3 (1.3 mg/L DOC matrix). Left: Evolution of the ozone concentration, right: Calculation of R_{ct} .

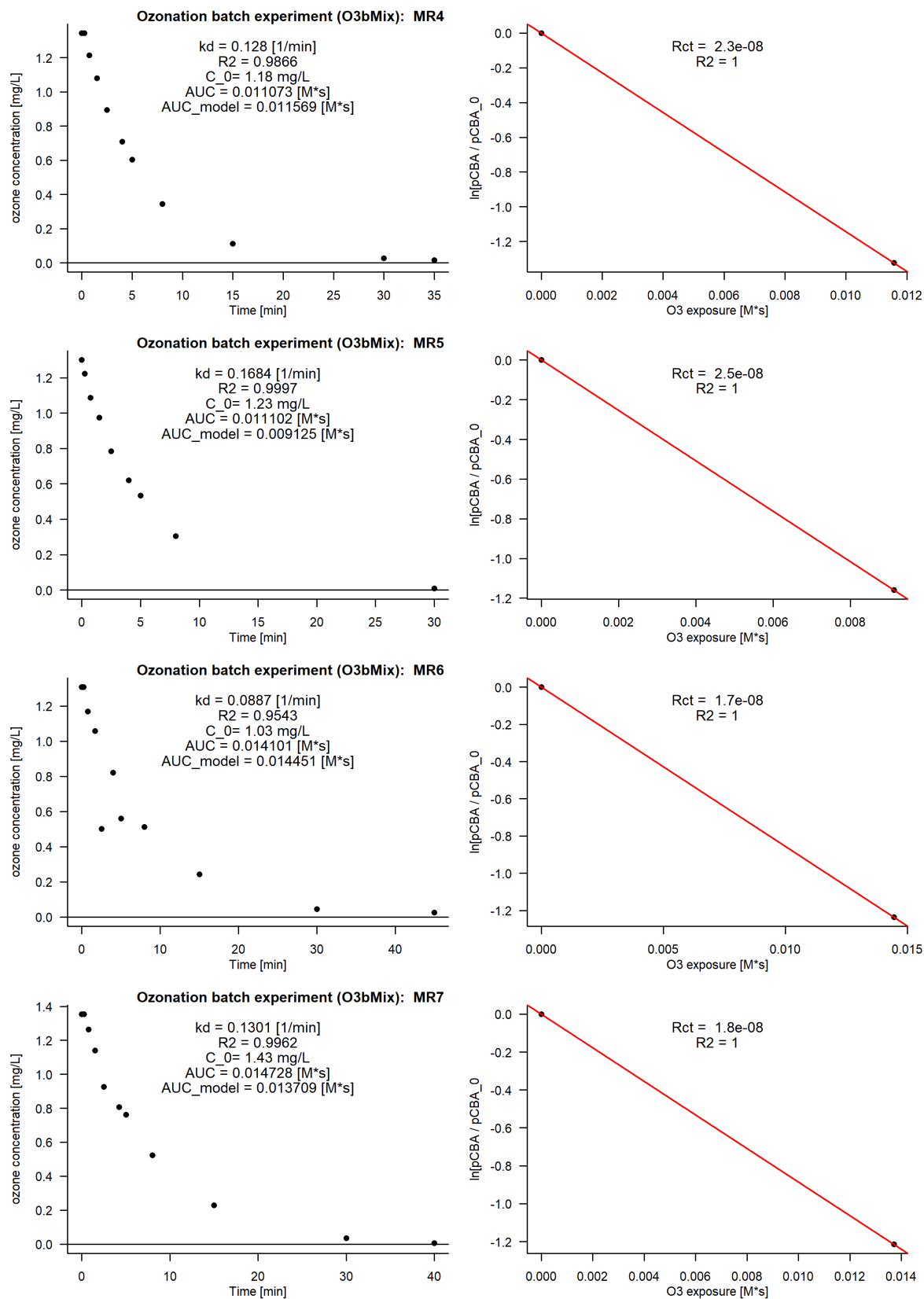


Figure S5: Ozonation conditions of O3bMix experiments with mixtures MR4-MR7 (1.3 mg/L DOC matrix). Left: Evolution of the ozone concentration, right: Calculation of R_{ct} .

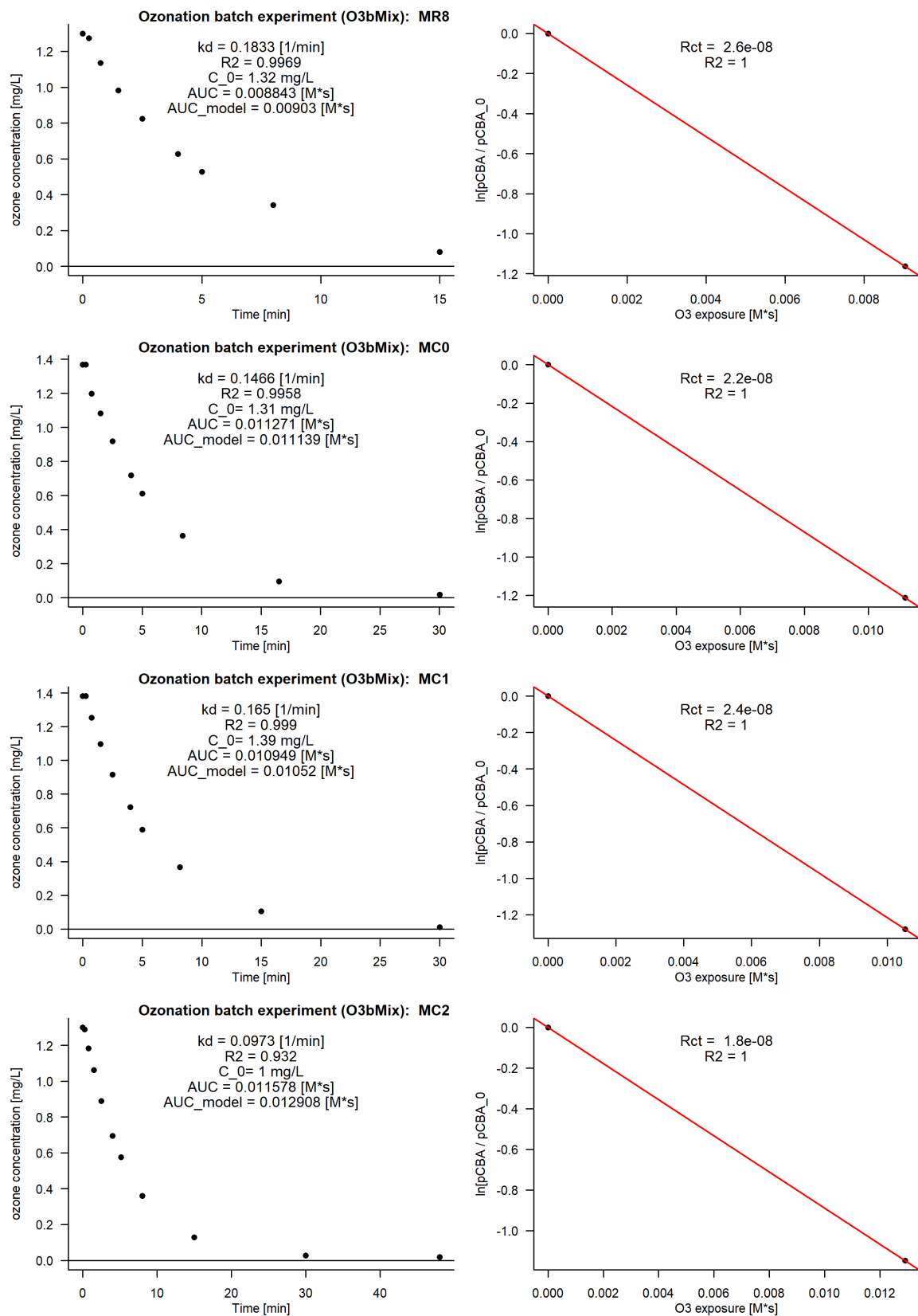


Figure S6: Ozonation conditions of O3bMix experiments with mixtures MR8 and MC0-MR2 (1.3 mg/L DOC matrix). Left: Evolution of the ozone concentration, right: Calculation of R_{ct} .

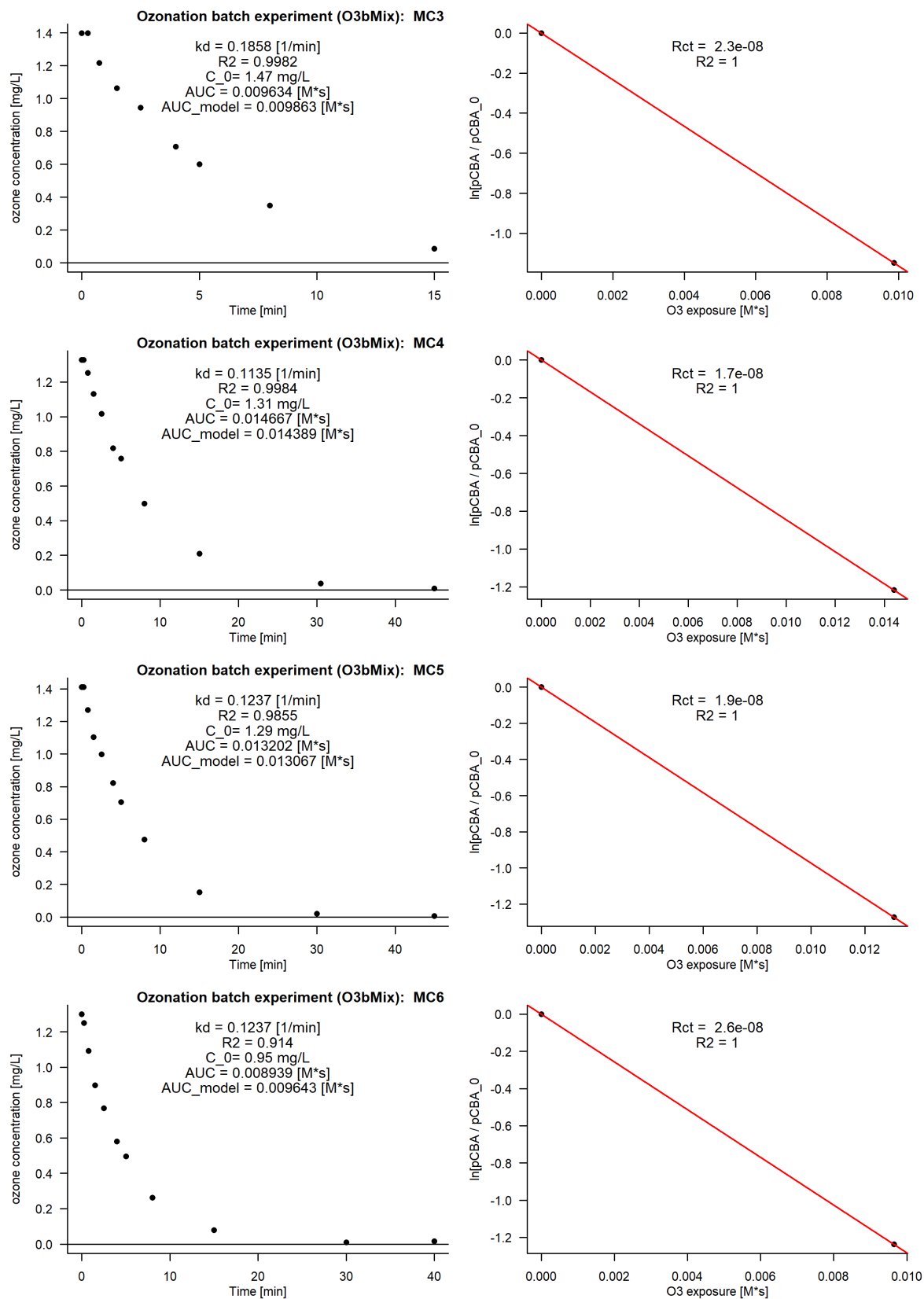


Figure S7: Ozonation conditions of O3bMix experiments with mixtures MC3-MC6 (1.3 mg/L DOC matrix). Left: Evolution of the ozone concentration, right: Calculation of R_{ct} .

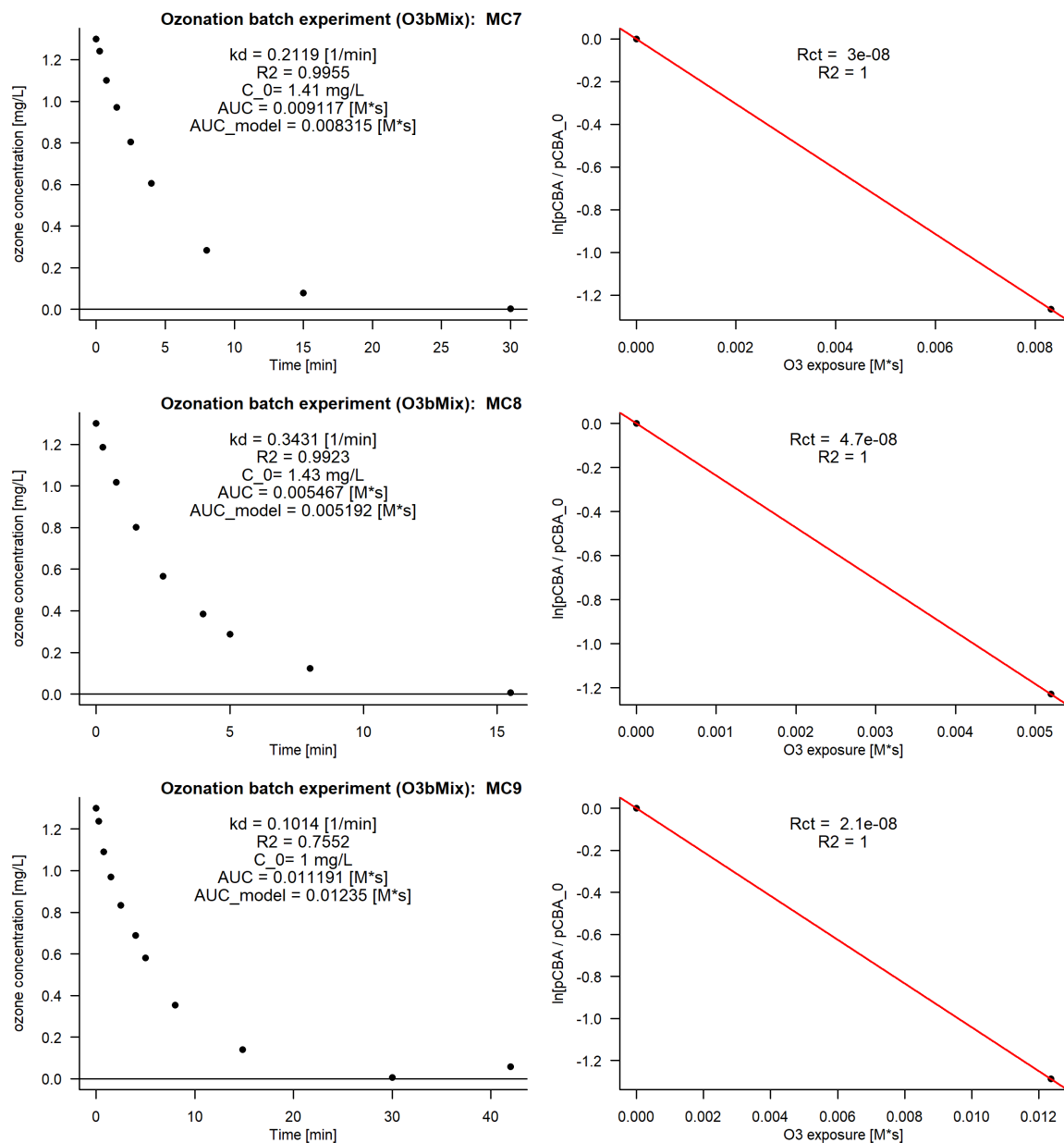


Figure S8: Ozonation conditions of O3bMix experiments with mixtures MR7-MR9 (1.3 mg/L DOC matrix). Left: Evolution of the ozone concentration, right: Calculation of R_{ct} .

Text S2.3 Ozonation conditions in the O3bAll batch experiment

Figure S9 shows the results of ozonation conditions of the *O3bAll* experiment. This experiment was performed with 0.15 mM acetate and 0.029 mM methanol, simulating a DOC of 1.3 mg/L.

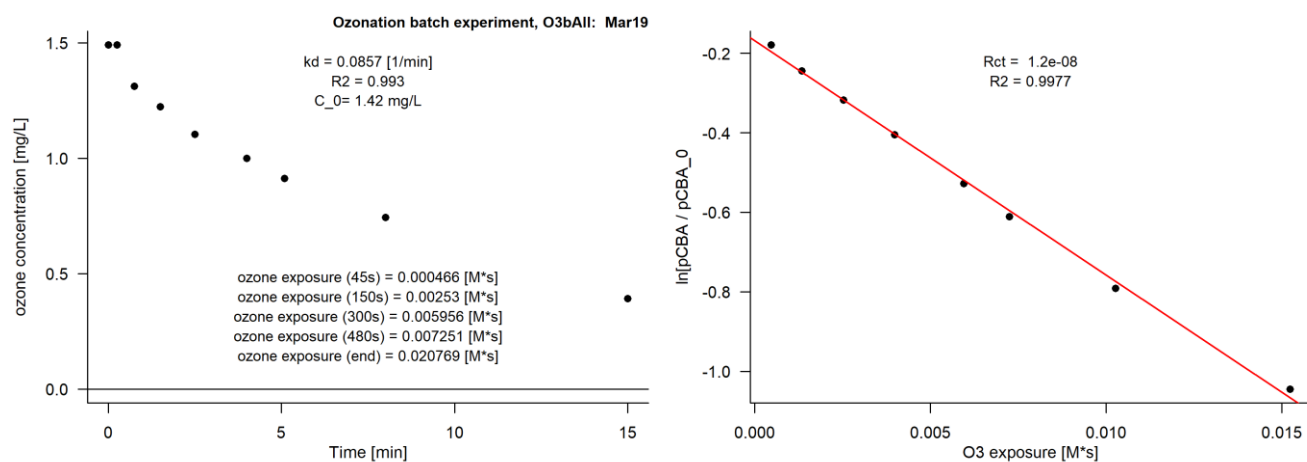


Figure S9: Ozonation conditions of the *O3bAll* experiment. Left: Evolution of the ozone concentration, right: Calculation of R_{ct} .

Text S3. Chemical analyses

Text S3.1 Method for the O3bMix samples

The LC gradient at a flow rate of 300 $\mu\text{L}/\text{min}$ and a column temperature of 30°C was:

Time in min	% (Nanopure water +0.1% formic acid)	% (Methanol + 0.1% formic acid)
0	100	0
4	100	0
19	5	95
25	5	95
25.1	100	0
30	100	0

The Electrospray ionization was triggered at a capillary temperature of 320°C and a spray voltage of 4 kV and 3kV for positive and negative ionization, respectively. The sheath gas flow rate was adjusted to 40, the auxiliary gas flow rate to 10, and the sweep gas flow rate to 0. Mass calibration and mass accuracy checks were carried out prior to the measurement with an in-house amino acid solution, which enhanced calibration for small masses. Full-scan acquisition was conducted at a resolution of 140'000 at m/z 200 and a scan range of m/z 50-750 with an injection time of 50 ms and an AGC target value of 5×10^5 . Ten data-dependent MS/MS scans were conducted after each full-scan with a resolution of 17'500 at m/z 200, an injection time of 100 ms, an AGC target value of 1×10^5 , and an isolation width of 1.0 Da. The dynamic exclusion was set to 3 sec. The data-dependent MS/MS were triggered after each full-scan on the $[M+H]^+$ and if negative ionization mode acquisition was conducted on the $[M-H]^-$ on masses predicted by O3-PPD (Lee et al. 2017, Schollée et al. 2021). The collision energy with which the MS/MS scans were triggered were calculated as $(-0.41) \times \text{exact mass} + 160$ for masses below m/z 350 and were set to 15 for masses above 350 m/z . Quantification of the parent micropollutants was done with the Tracefinder 4.1 software (Thermo Scientific). The smallest standard of the calibration row with a reasonable peak was treated as limit of quantification (LOQ) as given in Table S1.

Text S3.2 Method for the O3bAll samples

The LC gradient for the separation of the sample on the chromatographic column at a flow rate of 300 $\mu\text{L}/\text{min}$ and a column temperature of 30°C was:

Time in min	Nanopure water +0.1% FA in $\mu\text{L}/\text{min}$	Methanol + 0.1% FA in $\mu\text{L}/\text{min}$
0	260	40
5	260	40
20	15	285
29	15	285
29.5	260	40
35	260	40

The Electrospray ionization was triggered at a capillary temperature of 350°C and a spray voltage of 4 kV and 3 kV for positive and negative ionization, respectively. The sheath gas flow rate was adjusted to 40, the auxiliary gas flow rate to 10, and the sweep gas flow rate to 0. Mass calibration and mass accuracy checks were carried out prior to the measurement with an in-house amino acid solution, which enhanced calibration for small masses. Full-scan acquisition was conducted at a resolution of

140'000 at m/z 200 and a scan range of m/z 60-900 with an injection time of 100 ms and an AGC target value of 5×10^5 . Seven data-dependent MS/MS scans were conducted after each full-scan with a resolution of 17'500 at m/z 200, an injection time of 65 ms, an AGC target value of 1×10^5 , and an isolation width of 1.0 Da. The dynamic exclusion was set to 5 sec. The data-dependent MS/MS were triggered after each full-scan on the [M+H]⁺ and if negative ionization mode acquisition was conducted on the [M-H]⁻ on masses that were identified from the O3bMix experiments. The collision energy with which the MS/MS scans were triggered were calculated as $(-0.41) \times \text{exact mass} + 160$ for masses below m/z 350 and were set to 15 for masses above 350 m/z. Quantification of the parent micropollutants was done with the Tracefinder 4.1 software (Thermo Scientific). The smallest standard of the calibration row with a reasonable peak was treated as limit of quantification (LOQ) as given in Table S1.

Text S4. Identification of ozonation transformation products

Please note for 5 parent MPs (2-naphthalic-sulfonic-acid 2NS, amisulpride ASP, metoprolol MTO, oseltamivir OSE, and sulfapyridin SPD), for which an O3bMix reactor (MC8) failed, no OTP signals could be identified within the matrix simulating a DOC of 1.3 mg/L. However, the experiments were repeated within another matrix simulating a DOC of 5 mg/L (see Text S2.2). The results of these experiments are not shown. However, for these 5 MPs the result of the additional experiments were adopted.

Text S4.1 Generation of a list of suspect OTPs using laboratory ozonation batch experiments (O3bMix) with Compound Discoverer 2.1 (Approach 1)

For each parent MP, a separate Compound Discoverer 2.1 run in positive and negative mode was conducted to screen for corresponding OTP signals. The following details were chosen for the Compound Discoverer 2.1 run.

Two study factors were defined

- *Mix* with the levels: column mix, row mix, control mix
- *Time Points* with the levels: 0h (time zero sample, before ozone addition), S2 (2nd sample at 53s), S4 (4th sample at 158 s), S6 (6th sample at 308 s), S7 (7th sample at 488 s), Sc (sample of the control reactor)

For the uploaded measurement files, the study factors were selected. For the files of the control reactor, the *Time Points* Sc were selected instead of assigning different time point levels. In this way, the automatically calculated mean value of this group could be used. Figure S10 illustrates the selection.

At the Sample Groups and Ratios step, *Time points: 0h* was selected to generate Ratios as can be seen in Figure S11.

The applied Workflow is illustrated in Figure S12, the node settings are given in Table S5.

Input File Characterization

Manually define and assign the study variables for each input file. Or, to setup a regular expression that automatically extracts the study variables from each input file, click Advanced.

Delimiters: ☐ Underscore ☐ Hyphen ☐ Comma ☐ Space ☐ Plus ☐ Other

Assign Reset Advanced

Study Factors

Paste Copy Add

mix

MR4
MS7
MS9

Time points

0h
S2
S4
S6
S7
Sc

Samples

Sample	Sample Identifier	Sample Type	Time points	mix
1	20171027_neg_69	Sample	0h	MR4
2	20171027_neg_70	Sample	S2	MR4
3	20171027_neg_71	Sample	S4	MR4
4	20171027_neg_72	Sample	S6	MR4
5	20171027_neg_73	Sample	S7	MR4
6	20171027_neg_131	Sample	Sc	MS7
7	20171027_neg_132	Sample	Sc	MS7
8	20171027_neg_133	Sample	Sc	MS7
9	20171027_neg_134	Sample	Sc	MS7
10	20171027_neg_135	Sample	Sc	MS7
11	20171027_neg_142	Sample	0h	MS9
12	20171027_neg_143	Sample	S2	MS9
13	20171027_neg_144	Sample	S4	MS9
14	20171027_neg_145	Sample	S6	MS9
15	20171027_neg_146	Sample	S7	MS9

Cancel < Back Next > Finish

Figure S10: Input file assignment for Compound Discoverer 2.1 (Approach 1).

Sample Groups and Ratios

Select the study variables for sample grouping and add ratios for group comparisons.

Sample Group and Ratio Specification

Study Variables

☐ File

☒ Time points

☐ mix

☐ Sample Type

Variables printed in *italics* contain only a single value.

Manual Ratio Generation

Numerator:

Denominator:

Add Ratio

Bulk Ratio Generation

Denominators to be used:

☒ Time points : 0h

☐ Time points : S2

☐ Time points : S4

☐ Time points : S6

☐ Time points : S7

☐ Time points : Sc

Add Ratios

Generated Sample Groups

S7

Sample S7 MR4 F5: 20171027_neg_73

Sample S7 MS9 F15: 20171027_neg_146

Sc

Sample Sc MS7 F6: 20171027_neg_131

Sample Sc MS7 F7: 20171027_neg_132

Sample Sc MS7 F8: 20171027_neg_133

Sample Sc MS7 F9: 20171027_neg_134

Sample Sc MS7 F10: 20171027_neg_135

Generated Ratios

Clear All

X S2 / 0h

X S4 / 0h

X S6 / 0h

X S7 / 0h

X Sc / 0h

Cancel < Back Next > Finish

Figure S11: Sample Groups and Ratios for Compound Discoverer 2.1 (Approach 1).

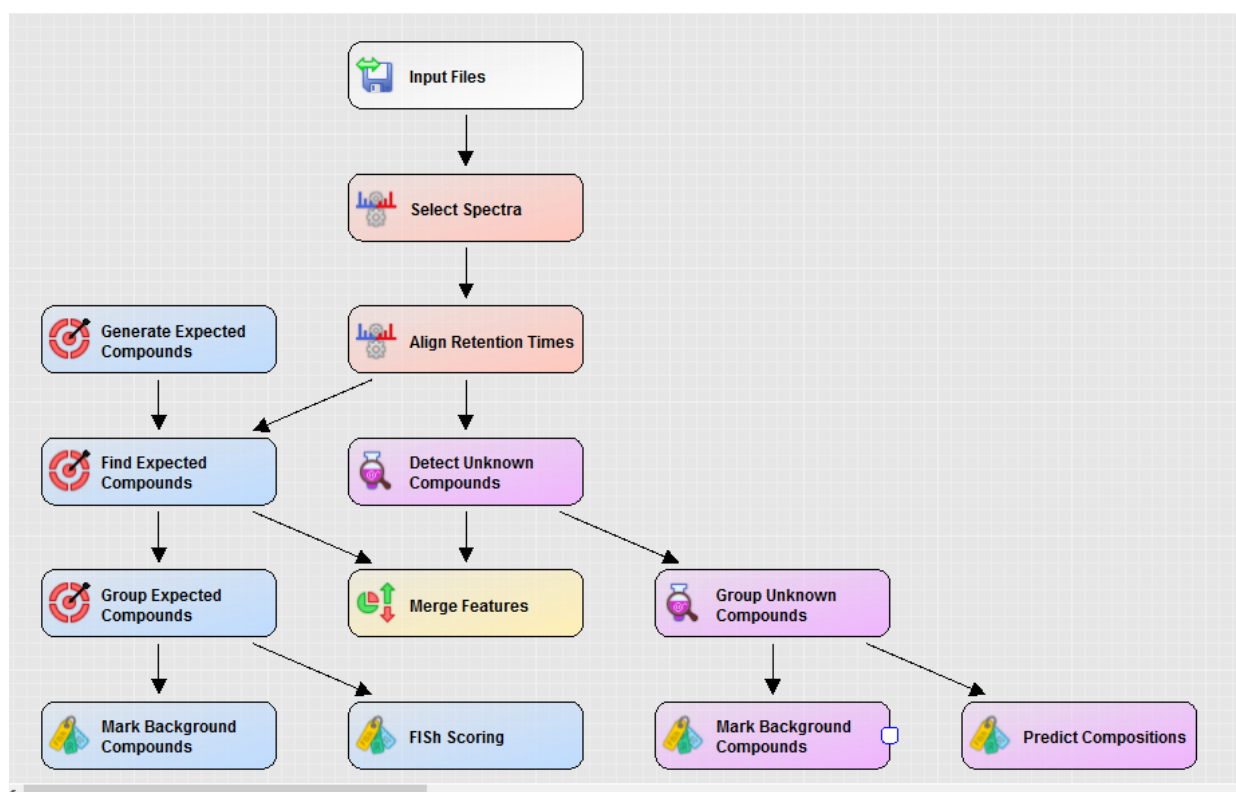


Figure S12: Applied workflow in Compound Discoverer 2.1 (Approach 1).

Table S5: Compound Discoverer 2.1 settings for the evaluation of the O3bMix experiments (Approach 1). In blue are highlighted settings that were different for the evaluation of samples measured in positive and negative mode. The first settings refer to the evaluation in positive mode and the second to the evaluation in negative mode. In the Generate Expected Compounds note the respective parent MP was selected as highlighted in green.

Select Spectra 1. General Settings: - Precursor Selection: Use MS(n - 1) Precursor 2. Spectrum Properties Filter: - Lower RT Limit: 0 - Upper RT Limit: 0 - First Scan: 0 - Last Scan: 0 - Ignore Specified Scan: - Lowest Charge State: 0 - Highest Charge State: 0 - Min. Precursor Mass: 100 Da - Max. Precursor Mass: 5000 Da - Total Intensity Threshold: 0 - Minimum Peak Count: 1 3. Scan Event Filters: - Mass Analyzer: (not specified) - MS Order: Any - Activation Type: (not specified) - Min. Collision Energy: 0 - Max. Collision Energy: 1000 - Scan Type: Any - Polarity Mode: Is + / Is- 4. Peak Filters: - S/N Threshold (FT-only): 1.5	Detect Unknown Compounds 1. General Settings: - Mass Tolerance [ppm]: 5 ppm - Intensity Tolerance [%]: 30 - S/N Threshold: 3 - Min. Peak Intensity: 10000/1000 - Ions: [M+H]⁺/ [M-H]⁻ - Min. Element Counts: C H - Max. Element Counts: C90 H190 Br3 Cl4 K2 N10 Na2 O18 P3 S5	Generate Expected Compounds 1. Compound Selection - Compound: respective parent Micropollutant 2. Dealkylation - Apply Dealkylation: True - Apply Dearylation: True - Max. #Steps: 1 - Min. Mass [Da]: 200
	Group Unknown Compounds 1. Compound Consolidation - Mass Tolerance: 5 ppm - RT Tolerance [min]: 0.5 2. Fragment Data Selection: - Preferred Ions: [M+H]⁺ / [M-H]⁻	3. Transformations - Phase : some meaningful reactions - Phase : 0 - Others: manually entered reactions according to (Schollée et al. 2018) - Max. # Phase : 0 - Max.# All Steps: 3 4. Ionization - Ions: [M+H]⁺ / [M-H]⁻
	Mark Background Compounds 1. General Settings: - Max. Sample/Blank: 5 - Max. Blank/Sample: 0 - Hide Background: True	Find Expected Compounds 1. General Settings - Mass Tolerance: 5ppm - Intensity Tolerance [%]: 30 - Intensity Threshold [%]: 0.1 - SN Threshold: 3

5. Replacements for Unrecognized Properties: - Unrecognized Charge Replacements: 1 - Unrecognized Mass Analyzer Replacements: ITMS - Unrecognized MS Order Replacements: MS2 - Unrecognized Activation Type Replacements: CID		- Min. # Isotopes: 2 - Min. Peak. Intensity: 1000 - Average Peak Width [min]: 0
	Merge Features 1. Peak Consolidation - Mass Tolerance: 5 ppm - RT Tolerance [min]: 0.5	Group Expected Compounds 1.Compound Consolidation - RT Tolerance [min] : 0.5 2. Fragment Data Selection - Preferred Ions : [M+H]⁺ / [M-H]⁻
	Predict Compositions 1.Prediction Settings - Mass Tolerance: 5 ppm - Min. Element Counts: CH - Max. Element Counts: : C90 H190 Br3 Cl4 K2 N10 Na2 O18 P3 S5 - Min. RDBE: 0 - Max. RDBE: 40 - Min. H/C: 0.1 - Max. H/C: 3.5 - Max. # Candidates: 10 2. Pattern Matching - Intensity Tolerance [%]: 30 - Intensity Threshold [%]: 0.1 - S/N Threshold: 3 - Use Dynamic Recalibration: True 3.Fragments Matching - Use Fragments Matching: True - Mass Tolerance: 5ppm - S/N Threshold: 3	Mark Background Compounds 1. General Settings: - Max. Sample/Blank: 5 - Max. Blank/Sample: 0 - Hide Background: True
Align Retention Times 1. General Settings: - Alignment Model: Adaptive curve - Maximum Shift [min]: 2 - Mass Tolerance: 5 ppm		FISH Scoring 1.General Settings - Annotate Full Tree: True - Match Transformations: True - High Acc. Mass Tolerance: 2.5mmu - Low Acc. Mass Tolerance: 0.5 Da 2. Fragmented Prediction Settings - Use General Rules: True - Use Libraries: True - Max. Depth: 5 - Aromatic Cleavage: True - Min. Fragment m/z: 50

After the processing of Compound Discoverer 2.1, the following steps were conducted:

- Checking the presence of 3-5 internal standards and 1-3 parent compounds.
- Reducing the hits within the Merged Features tab by an automatic filtering with the following criteria as illustrated in Figure S13:
 1. the peak had to increase at least 5 times from the samples before to the samples after ozone addition
 2. presence in both mixtures containing the respective parent MP
 3. absence in the control mixture

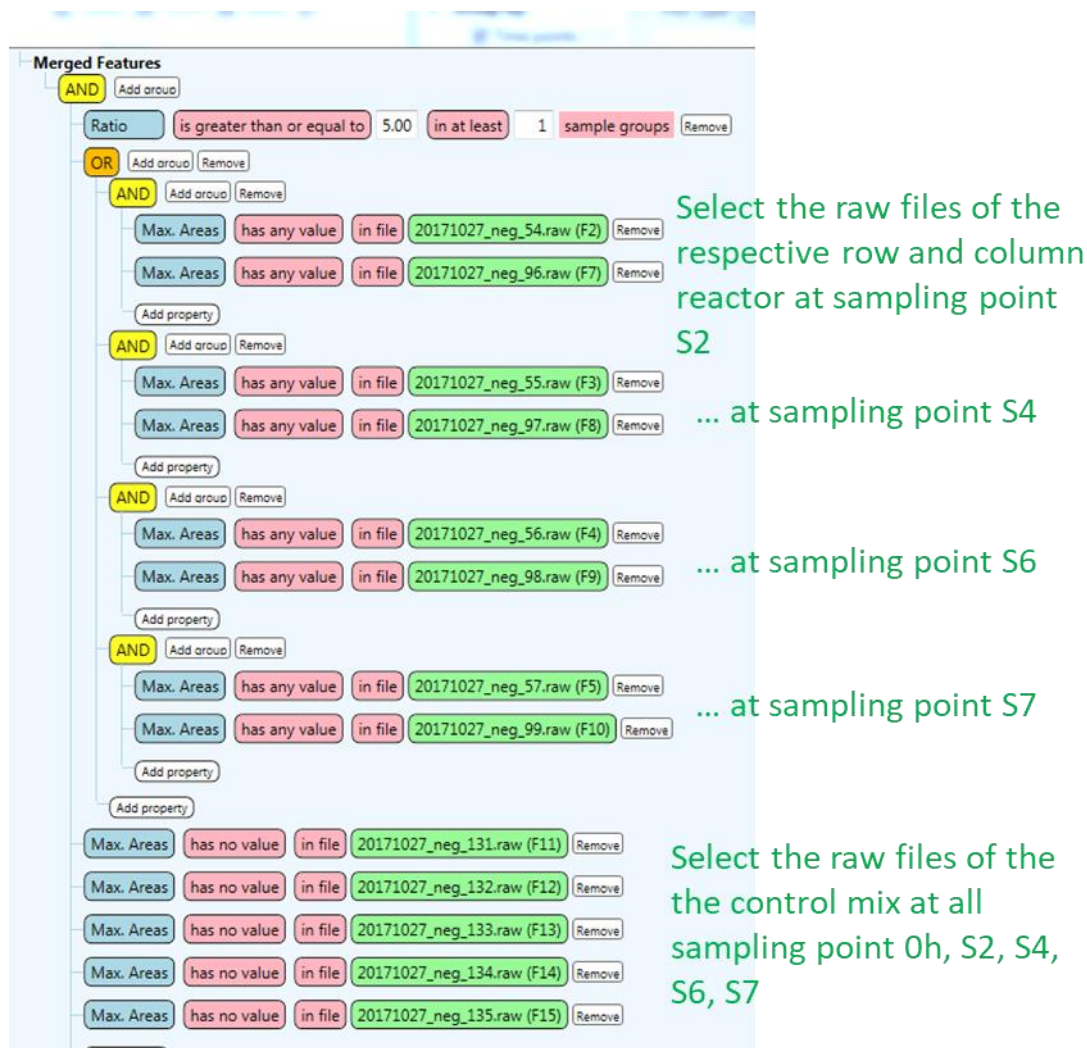


Figure S13: Filter criteria applied for Merged Features in Compound Discoverer 2.1 (Approach 1).

Afterwards, the profiles were manually investigated and selected based on the following criteria:

- reasonable peak shape by visual inspection
- meaningful presence / absence in the treated / control samples
- higher chance to be selected if questionable if present in the *Expected Compounds* tab

Text S4.2 Identification of OTP signals that are simultaneously forming in O3bAll and WWTP samples with Compound Discoverer 2.1 (Approach 2)

To identify OTP signals that are formed in our reference laboratory ozonation batch experiment (*O3bAll*) and in the WWTPs, we used Compound Discoverer 2.1 (Thermo Scientific) for a separate evaluation in positive and negative mode. We uploaded the samples from the *O3bAll* experiments and from the four different WWTPs. Since the *O3bMix* experiments were already performed and measured within Approach 1 (A-C), we uploaded these samples as well. By doing so, we performed Approach 2C and 2F at the same time.

Three study factors (*Mix*, *Sample points*, *type*) with different levels were applied (see Figure S14). Most of them were only important for a neat arrangement in the resulting table. Relevant for the filtering were only the following three levels of the study factor *type*:

- *13_70_all* (assigned to all samples of the *O3bAll* experiment after ozone addition)
- *all_S0* (assigned to all samples of the *O3bAll* experiment before ozone addition)
- *Blind* (assigned to blind samples)

The applied Workflow is illustrated in Figure S15, the node settings are given in Table S6.

Study Definition Input Files Samples Analysis Results						
Sample	Sample Identifier	Sample Type	mix	Sample point	Type	
<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
⊕ S1	20181203_online_neg_017	Sample ▾	3_NG ▾	3_EFF ▾	3_EFF ▾	
⊕ S2	20181203_online_neg_018	Sample ▾	3_NG ▾	3_EFF ▾	3_EFF ▾	
⊕ S3	20181203_online_neg_019	Sample ▾	2_GL ▾	3_EFF ▾	3_EFF ▾	
⊕ S4	20181203_online_neg_020	Sample ▾	2_GL ▾	3_EFF ▾	3_EFF ▾	
⊕ S5	20181203_online_neg_021	Sample ▾	2_GL ▾	3_EFF ▾	3_EFF ▾	
⊕ S6	20181203_online_neg_022	Sample ▾	1_AR ▾	3_EFF ▾	3_EFF ▾	
⊕ S7	20181203_online_neg_023	Sample ▾	4_PR ▾	3_EFF ▾	3_EFF ▾	
⊕ S8	20181203_online_neg_025	Sample ▾	3_NG ▾	2_OZO ▾	2_OZO ▾	
⊕ S9	20181203_online_neg_026	Sample ▾	3_NG ▾	2_OZO ▾	2_OZO ▾	
⊕ S10	20181203_online_neg_027	Sample ▾	2_GL ▾	2_OZO ▾	2_OZO ▾	
⊕ S11	20181203_online_neg_028	Sample ▾	2_GL ▾	2_OZO ▾	2_OZO ▾	
⊕ S12	20181203_online_neg_029	Sample ▾	2_GL ▾	2_OZO ▾	2_OZO ▾	
⊕ S13	20181203_online_neg_030	Sample ▾	1_AR ▾	2_OZO ▾	2_OZO ▾	
⊕ S14	20181203_online_neg_031	Sample ▾	4_PR ▾	2_OZO ▾	2_OZO ▾	
⊕ S15	20181203_online_neg_033	Sample ▾	3_NG ▾	1_BIO ▾	1_BIO ▾	
⊕ S16	20181203_online_neg_034	Sample ▾	3_NG ▾	1_BIO ▾	1_BIO ▾	
⊕ S17	20181203_online_neg_035	Sample ▾	2_GL ▾	1_BIO ▾	1_BIO ▾	
⊕ S18	20181203_online_neg_036	Sample ▾	2_GL ▾	1_BIO ▾	1_BIO ▾	
⊕ S19	20181203_online_neg_037	Sample ▾	2_GL ▾	1_BIO ▾	1_BIO ▾	
⊕ S20	20181203_online_neg_038	Sample ▾	1_AR ▾	1_BIO ▾	1_BIO ▾	
⊕ S21	20181203_online_neg_039	Sample ▾	4_PR ▾	1_BIO ▾	1_BIO ▾	
⊕ S22	20181203_online_neg_042	Sample ▾	control ▾	Blind ▾	Blind ▾	
⊕ S23	20181203_online_neg_071	Sample ▾	control ▾	Blind ▾	Blind ▾	
⊕ S24	20181203_online_neg_086	Sample ▾	control ▾	Blind ▾	Blind ▾	
⊕ S25	20181203_online_neg_112	Sample ▾	control ▾	Blind ▾	Blind ▾	
⊕ S26	20181203_online_neg_113	Sample ▾	MRall ▾	S0 ▾	all_S0 ▾	
⊕ S27	20181203_online_neg_114	Sample ▾	MRall ▾	S2 ▾	_13_70all ▾	
⊕ S28	20181203_online_neg_115	Sample ▾	MRall ▾	S4 ▾	_13_70all ▾	
⊕ S29	20181203_online_neg_116	Sample ▾	MRall ▾	S6 ▾	_13_70all ▾	
⊕ S30	20181203_online_neg_117	Sample ▾	MRall ▾	S7 ▾	_13_70all ▾	
⊕ S31	20181203_online_neg_119	Sample ▾	MRall ▾	S0 ▾	all_S0 ▾	
⊕ S36	20181203_online_neg_124	Sample ▾	control ▾	Blind ▾	Blind ▾	
⊕ S37	20181203_online_neg_125	Sample ▾	MR0 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S38	20181203_online_neg_126	Sample ▾	MR0 ▾	S6 ▾	_13_70_S6 ▾	
⊕ S39	20181203_online_neg_127	Sample ▾	MR1 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S40	20181203_online_neg_128	Sample ▾	MR1 ▾	S6 ▾	_13_70_S6 ▾	
⊕ S41	20181203_online_neg_129	Sample ▾	MR2 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S42	20181203_online_neg_130	Sample ▾	MR2 ▾	S6 ▾	_13_70_S6 ▾	
⊕ S43	20181203_online_neg_131	Sample ▾	MR3 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S44	20181203_online_neg_132	Sample ▾	MR3 ▾	S6 ▾	_13_70_S6 ▾	
⊕ S45	20181203_online_neg_133	Sample ▾	MR4 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S46	20181203_online_neg_134	Sample ▾	MR4 ▾	S6 ▾	_13_70_S6 ▾	
⊕ S47	20181203_online_neg_136	Sample ▾	MR5 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S48	20181203_online_neg_137	Sample ▾	MR5 ▾	S6 ▾	_13_70_S6 ▾	
⊕ S49	20181203_online_neg_138	Sample ▾	MR6 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S50	20181203_online_neg_139	Sample ▾	MR6 ▾	S6 ▾	_13_70_S6 ▾	
⊕ S51	20181203_online_neg_140	Sample ▾	MR7 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S52	20181203_online_neg_141	Sample ▾	MR7 ▾	S6 ▾	_13_70_S6 ▾	
⊕ S53	20181203_online_neg_142	Sample ▾	MR8 ▾	S2 ▾	_13_70_S2 ▾	
⊕ S54	20181203_online_neg_143	Sample ▾	MR8 ▾	S6 ▾	_13_70_S6 ▾	

Figure S14: Input file assignment for Compound Discoverer 2.1 (Approach 2).

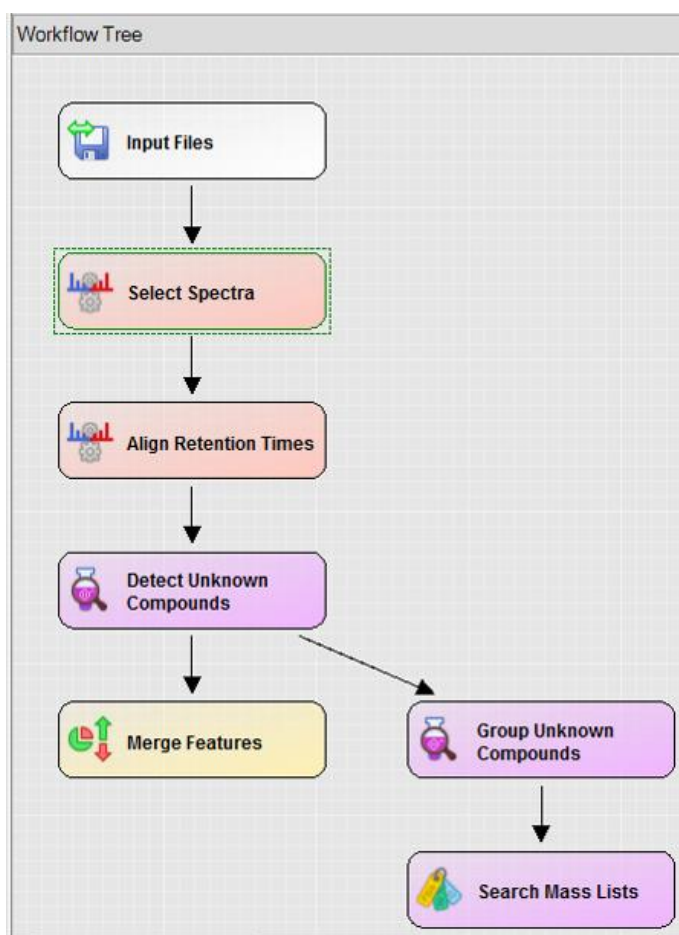


Figure S15: Applied workflow in Compound Discoverer 2.1 (Approach 2).

Table S6: Compound Discoverer 2.1 settings for the evaluation of the O3bAll experiments and WWTP samples (Approach 2). In blue are highlighted settings that were different for the evaluation of samples measured in positive and negative mode. The first settings refer to the evaluation in positive mode and the second to the evaluation in negative mode. In green are the two lists of suspected neutral OTP masses that were uploaded. The first list was generated through Approach 1 and the second list is a compilation of OTP masses known from literature.

Select Spectra 1. General Settings: - Precursor Selection: Use MS(n - 1) Precursor 2. Spectrum Properties Filter: - Lower RT Limit: 0 - Upper RT Limit: 0 - First Scan: 0 - Last Scan: 0 - Ignore Specified Scan: - Lowest Charge State: 0 - Highest Charge State: 0 - Min. Precursor Mass: 100 Da - Max. Precursor Mass: 5000 Da - Total Intensity Threshold: 0 - Minimum Peak Count: 1 3. Scan Event Filters: - Mass Analyzer: (not specified) - MS Order: Any - Activation Type: (not specified) - Min. Collision Energy: 0 - Max. Collision Energy: 1000	Align Retention Times 1. General Settings: - Alignment Model: Adaptive curve - Maximum Shift [min]: 2 - Mass Tolerance: 5 ppm	Group Unknown Compounds 1. Compound Consolidation - Mass Tolerance: 5 ppm - RT Tolerance [min]: 0.5 2. Fragment Data Selection: - Preferred Ions: [M+H] ⁺ / [M-H] ⁻
	Detect Unknown Compounds 1. General Settings: - Mass Tolerance [ppm]: 5 ppm - Intensity Tolerance [%]: 30 - S/N Threshold: 3 - Min. Peak Intensity: 10000/1000 - Ions: [M+H] ⁺ +H / [M-H] ⁻ -1 - Min. Element Counts: C H - Max. Element Counts: C90 H190 Br3 Cl4 K2 N10 Na2 O18 P3 S5	Search mass list Input file(s): CDin_allMPs_CDRPneg_neutral.csv, CDin_allMPs_litneg_neutral.csv - Mass Tolerance: 15 ppm - RT Tolerance: 1 Consider Retention Time: False

Merge Features

- Scan Type: Any - Polarity Mode: Is + / Is- 4. Peak Filters: - S/N Threshold (FT-only): 1.5 5. Replacements for Unrecognized Properties: - Unrecognized Charge Replacements: 1 - Unrecognized Mass Analyzer Replacements: ITMS - Unrecognized MS Order Replacements: MS2 - Unrecognized Activation Type Replacements: CID	1. Peak Consolidation - Mass Tolerance: 5 ppm - RT Tolerance [min]: 0.5	
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To identify signals in the WWTPs that were also formed in our *O3bAll* experiments, we used an automatic filter and a manual selection. The automatic filter had the following criteria (see Figure S16):

Signals had to

- be in at least one *O3bAll* sample after ozone addition
- increase at least 2 times in the *O3bAll* experiment from the sample before ozone addition to any sample at a later time point
- be 5 times higher in the *O3bAll* samples than in the blank samples (20 mL nanopure water augmented with 16 µL ISTD stock solution)
- have any value in WWTP samples after ozonation or post-treatment
- have a retention time between 4 and 28 min
- have a maximal area of at least 10000 / 1000 in positive / negative mode

Signals were manually selected, based on the pattern in the ozonation batch experiments over time and the occurrence in the WWTP samples. Since we had knowledge of suspect OTPs of 70 MPs from Approach 1, we used this information by uploading the list of all exact masses of identified OTP signals. That way, an easier selection of OTP signals was possible as MS² spectra were available. However, Approach 2 can be applied without this prior knowledge.

Result Filters

☐ ON ☐ Compounds
☐ ON ☐ Compounds per File
☒ ON ☒ Merged Features
☐ ON ☐ Features
☐ ON ☐ Mass List Search Results
☐ ON ☐ Input Files

Merged Features

AND Add group Remove

OR Add group Remove

Max. Areas has any value in file 20181203_online_neg_114.raw (F27) Remove

Max. Areas has any value in file 20181203_online_neg_115.raw (F28) Remove

Max. Areas has any value in file 20181203_online_neg_116.raw (F29) Remove

Max. Areas has any value in file 20181203_online_neg_117.raw (F30) Remove

Add property

Max. Area is greater than or equal to 10000.00 Remove

OR Add group Remove

Ratio is greater than or equal to 2.00 in sample group (_L13_70all) / (all_S0) Remove

Ratio has no value in sample group (_L13_70all) / (all_S0) Remove

Add property

OR Add group Remove

Ratio is greater than or equal to 5.00 in sample group (_L13_70all) / (Blind) Remove

Add property

OR Add group Remove

Max. Areas has any value in file 20181203_online_neg_017.raw (F1) Remove

Max. Areas has any value in file 20181203_online_neg_018.raw (F2) Remove

Max. Areas has any value in file 20181203_online_neg_019.raw (F3) Remove

Max. Areas has any value in file 20181203_online_neg_020.raw (F4) Remove

Max. Areas has any value in file 20181203_online_neg_021.raw (F5) Remove

Max. Areas has any value in file 20181203_online_neg_022.raw (F6) Remove

Max. Areas has any value in file 20181203_online_neg_023.raw (F7) Remove

Max. Areas has any value in file 20181203_online_neg_025.raw (F8) Remove

Max. Areas has any value in file 20181203_online_neg_026.raw (F9) Remove

Max. Areas has any value in file 20181203_online_neg_033.raw (F15) Remove

Max. Areas has any value in file 20181203_online_neg_034.raw (F16) Remove

Max. Areas has any value in file 20181203_online_neg_035.raw (F17) Remove

Max. Areas has any value in file 20181203_online_neg_036.raw (F18) Remove

Max. Areas has any value in file 20181203_online_neg_037.raw (F19) Remove

Max. Areas has any value in file 20181203_online_neg_038.raw (F20) Remove

Max. Areas has any value in file 20181203_online_neg_039.raw (F21) Remove

Max. Areas has any value in file 20181203_online_neg_027.raw (F10) Remove

Max. Areas has any value in file 20181203_online_neg_028.raw (F11) Remove

Max. Areas has any value in file 20181203_online_neg_029.raw (F12) Remove

Max. Areas has any value in file 20181203_online_neg_030.raw (F13) Remove

Max. Areas has any value in file 20181203_online_neg_031.raw (F14) Remove

Add property

RT [min] is between 4.00 and 28.00 Remove

Add property

☐ Show all tables

All O3ball samples after ozone addition

All WWTP samples after ozonation or post-treatment

Figure S16: Filter criteria applied for Merged Features in Compound Discoverer 2.1 (Approach 2).

Text S5. Measured and calculated abatement of micropollutants in *O3bAll*

Figure S17 illustrates the observed abatement of the four parent MPs atrazine, bezafibrate, ibuprofen, and ketoprofen in the ozonation batch experiments (*O3bAll*) and the expected abatement based on second-order rate constants for their reactions with ozone and •OH (data derived from literature). The good match between experimental data and calculations shows that the ozonation batch experiments are well controlled in terms of ozone and hydroxyl radical exposures and the R_{ct} .

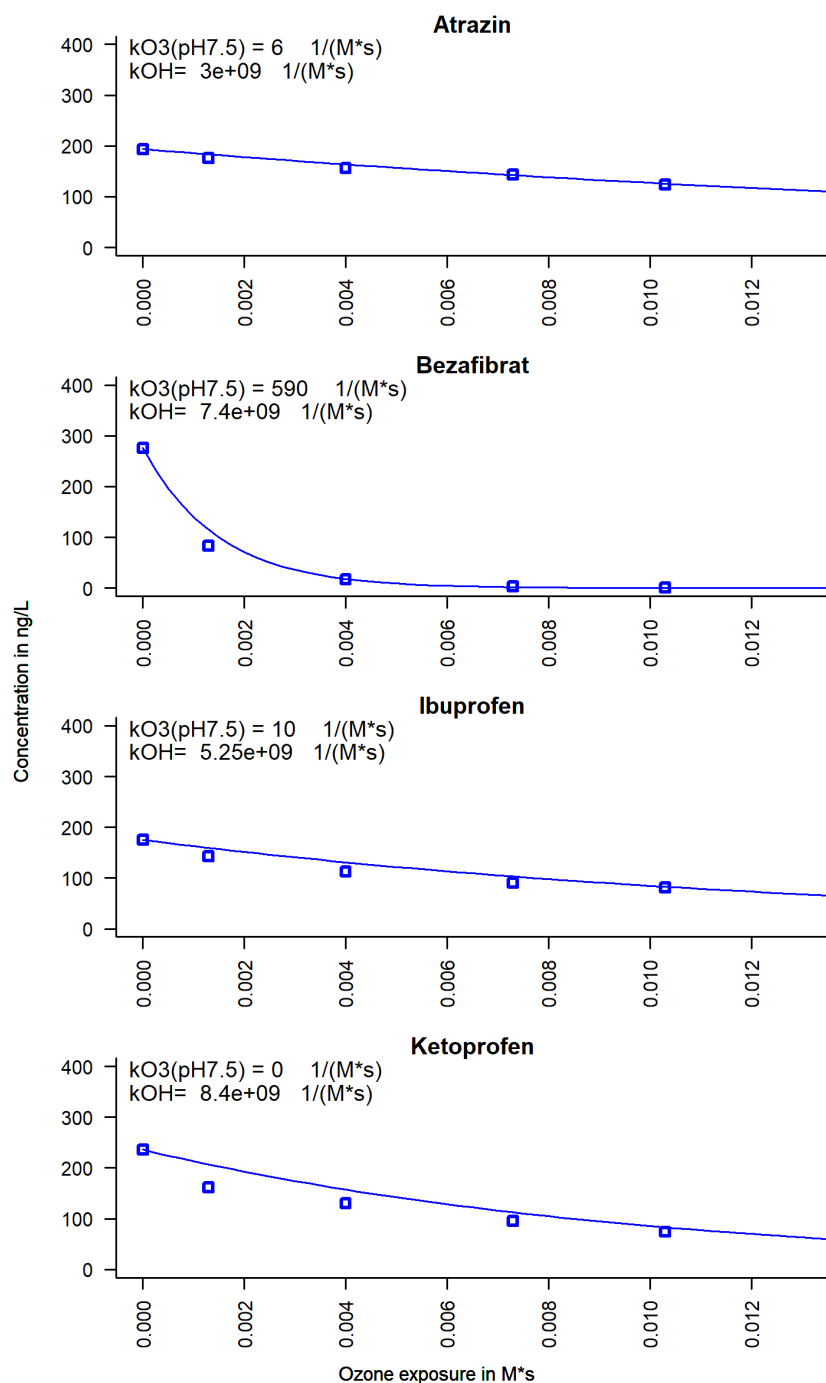


Figure S17: Abatement of the MPs atrazine, bezafibrate, ibuprofen, and ketoprofen (symbols) as a function of the ozone exposure in the *O3bAll* experiment. The line illustrates the calculated abatement based on the second order rate constants for the reaction of the MPs with ozone and hydroxyl radical from the literature (Acero et al. 2000, Huber et al. 2003, Real et al. 2009).

Text S6. Results of OTPs found in batch experiments and wastewater treatment

Overall, 84 OTPs could be detected, which originated from 40 of the 87 investigated MPs. For 47 MPs, no OTPs were assigned. Between 1 and 7 of the detected 84 OTPs were assigned to the same parent MPs (Figure S18). For most parents (22), only 1 OTP was found, for 17 MPs 2-4 OTPs, while for carbamazepine 6 OTPs and for sitagliptin 7 OTPs were observed.

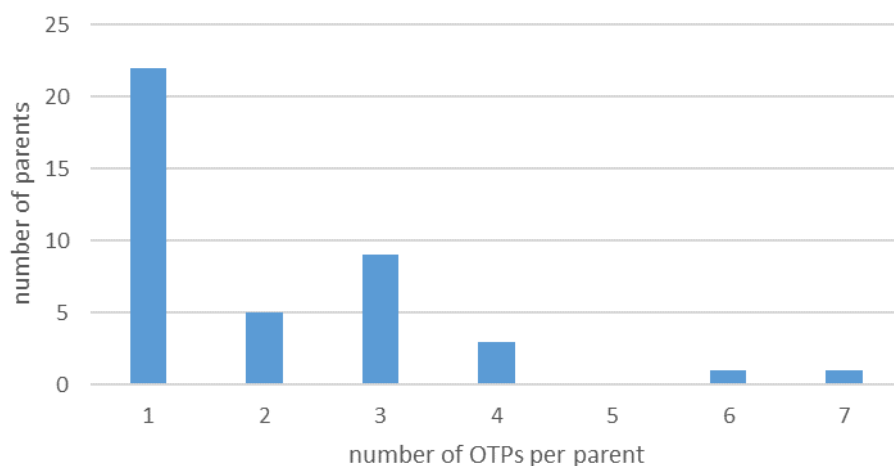


Figure S18: Amount of OTPs (from all 84 detected OTPs) assigned to a parent.

In the attached Excel Table S3, details on all the detected 84 OTPs are listed.

Figure S19:

In the attached pdf file, all the results for the 84 OTPs found in wastewater treatment are visually shown. The following information is given for each OTP:

- Top left: OTP formation in ozonation batch experiments
- Top right: Fate of parent MP in wastewater treatment
- Middle: (left) Fate of OTP in post-treatment (% formed or abated) and (right) OTP peak area in all wastewater samples in the four WWTPs. The score and evaluation of MS² spectra match of WWTP OZO samples to batch samples is given below.
- Bottom: Information on OTP identification (exact mass, formula, atomic modification, proposed structure, confidence level, Massbank identifier, MS² spectra and interpretation of it)

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