

Fate of emerging and priority micropollutants during the sewage sludge treatment - Part 2: Mass balances of organic contaminants on sludge treatments are challenging

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- Fate of emerging and priority micropollutants during the sewage sludge treatments: case study of
- 2 Paris conurbation part 2: mass balances of organic contaminants on sludge treatments are
- 3 challenging.
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Highlights

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- 71 OCs mass balances were performed in full-scale sludge treatment facilities
- Screw decanter was selected as reference as its action is only physical
- Several OCs mass balances are frequently not correct in screw decanter
- The frequency of correct mass balances is significantly correlated to logKow

1	•	Removal of the 71 OCs are reported in full-scale sludge digestion process			

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14 **Abbreviations**

- AP, alkylphenols (n=4 including t-OP, ter-octylphenol, NP1EO, nonylphenol
- monoethoxylate, NP2EO, nonylphenol diethoxylate, NP, nonylphenol)
- DMin, DMout, dry matter content of the sludge at the input and output of the process

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- LAS, linear alkylbenzene sulfonates (sum of the four congeners LAS-C10, LAS-C11, LAS-
- 21 C12, LAS-C13)
- 22 LQ/LD, limit of quantification/limit of detection
- OC, organic contaminants
- PAE, phthalates (n=4 including DEP, diethyl phthalate, DnBP Di-n-butyl phthalate, BBP,
- benzylbutyl phthalate, DEHP, diethylhexyl phthalate)
- PAH, polycyclic aromatic hydrocarbons (sum of 13 congeners including fluor, fluoranthene,
- BbF, benzo(b)fluoranthene, BaP, benzo(a)pyrene)
- PCB, polychlorobiphenyls [sum of 20 congeners including PCB28 (3Cl), PCB180 (7Cl),
- 29 PCB209 (10Cl)]
- 30 PFA, perfluorinated acids (n=2 including PFOA, perfluorooctanoic acids and PFOS,
- 31 perfluorooctanesulfonic acids
- 32 PHP, pharmaceutical products
- Qin, Qout, input and output flows of the process
- 34 SIAAP, Greater Paris sanitation authority
- 35 SRT, solid retention time
- 36 STP, sludge treatment plant
- 37 TSS, total suspended solid
- 38 WWTP, wastewater treatment plants

39 Abstract

This paper analyzes the fate of 71 priority and emerging organic contaminants all along the treatment trains of sewage sludge treatment facilities in Paris including dewatering by centrifugation, thermal drying and anaerobic digestion. It aimed at proposing and applying a mass balances calculation methodology to each process and pollutant. This data validation strategy demonstrated the complexity to perform representative inlet/outlet sampling and analysis campaigns at industrial scales regarding organic compounds and to propose options to overcome this issue. Centrifugation and drying processes only implied physical mechanisms as phase separation and water elimination. Hence, correct mass balance were expected observed for organic contaminants if sampling and analysis campaigns were representative. This was the case for hydrophobic and neutral compounds. For the other more hydrophilic and charged compounds, the mass balances were scarcely correct. Thus, the conventional sampling and analytical practices used with sludge should be questioned and adapted to better take into account the high heterogeneity of sludge and the evolution of matrix effect within sludge treatment processes on micropollutant determination. For the biological anaerobic digestion process where degradations can occur and removals can be observed, the mass balances were deeply interpreted for 60 contaminants. This process contributed to the elimination above 70% of 21 detected compounds including 16 pharmaceuticals, 2 phthalates, 2 hormones and 1 perfluorinated compound. Removals of domperidone, propranolol, escitalopram, lidocaine, verapamil and cefoperazone under this condition were reported for the first time.

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Key-words

Sewage sludge; organic compounds; mass balances; screw decanter; thermal drying;

anaerobic digestion

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1. Introduction

Wastewater treatment plants (WWTP) produce an important quantity of sewage sludge by

treating suspended solids, organic matter and nutrients from human activities. Actually, about

1 million ton dry matter (DM) of sludge are produced each year in France, against a total of

11 million tons DM of sludge in all Europe (Kelessidis and Stasinakis, 2012).

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Sludge are treated and conditioned before land application, incineration or landfilling (Fytili

and Zabaniotou, 2008; Mininni et al., 2015). In France (> 70% of the DM mass), as well as in

Europe in general (> 50%), the main pathway for sludge management is still land application.

The sewage sludge treatment aims to reduce fermentescibility, nuisances (particularly odour),

pathogens content and vector attraction potential as well as to reduce bulk and improve their

physical characteristics to aid handling, dewatering and acceptability for use on agricultural

land. Current practices in Europe are based on the requirements of the 1986 Directive on the

use of sewage sludge in agriculture (86/278/EEC) that was transposed to national laws with specific limits on heavy metals, organic contaminants and pathogens in various member states (Mininni et al., 2015). In the new EU regulation on fertilising products (2019/1009), sewage sludge composts and digestates are unauthorized Component Material Categories which could prevent their use in agriculture with the product status.

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A wide range of organic contaminants (OC) sorbs on sewage sludge during wastewater treatment (Clara et al., 2007; Mailler et al., 2014; Ruel et al., 2012) leading to their contamination by various compounds such as alkylphenols (AP), phthalates (PAE), polycyclic aromatic hydrocarbons (PAH), pharmaceuticals products (PHP), hormones or organotins (Bergé et al., 2012, 2013; Clarke and Smith, 2011; Mailler et al., 2017; Meng et al., 2016; Verlicchi and Zambello, 2015). Data on OC occurrence in sludge are nowadays extensive and diverse in the literature but the impact of sludge treatments on OC concentrations is difficult to assess considering that most studies do not distinguish different types of sludge from a given sludge treatment plant (STP); they mostly aimed at characterizing the contamination of raw and/or final treated sludge disregarding the type of sludge treatment. In addition, sludge sampling practises are still unsophisticated due to technical field constraints, they are mostly punctual samples taken from pipe or storage tank stitchings. Sludge being a very heterogeneous material, this lead to a sampling representativeness issue, in particular when assessing pollutants with highly variable properties such as hydrophilic and polar OC. Finally, sludge treatment includes physical, chemical and microbiological manipulation and may involve mechanical dewatering, heating, chemical reactions and microbiological transformation processes that may influence the loss, or potential formation, of OC by volatilisation, biological or abiotic transformation, bound residues formation or extraction with excess liquors that returned to the inlet of the WWTP (Mailler et al., 2017).

In this context, a study was conducted aiming at i) featuring OC pattern of the different types of sludge from Paris conurbation and the contamination of residual waters from thermal drying and centrifugation, ii) assessing the industrial scale efficiency of four widespread sludge treatment processes to remove/decrease these contaminations based on mass balances and iii) questioning the representativeness of conventional sampling and micropollutant analysis in sludge practises. The first objective was treated in the part 1 paper (Mailler et al., 2017) while this part 2 paper deals with the second and third objectives, based on the same database.

Therefore, centrifugation, anaerobic digestion, thermal drying and cooking + press filtration were considered. A total of 71 OC including Water Framework Directive priority pollutants, PHP, hormones, perfluorinated acids (PFA), linear alkylbenzene sulfonate (LAS), AP, PAE, PAH and polychlorobiphenyls (PCB) were investigated (SI Table 1). As the objective of this

study was also to apply mass balances calculations at industrial scale for each sludge

treatment process, OC in the residual water from thermal drying and centrifugation were monitored too.

The novelty of this paper is the of mass balances calculation at full-scale to this large spectrum of pollutants within three widespread sludge treatment processes, i.e. dewatering by centrifugation, anaerobic digestion and thermal drying, to question the sampling and analytical conventional practises and to analyse the behaviour of OC in those processes. Such mass balance analysis is an effective way to understand and model the fate of numerous OC in sludge treatment trains but to the best of our knowledge, its application to various treatments and OC is not available in the literature, especially based on sludge produced at real industrial scale STPs. In addition, sampling and analytical practises applied for micropollutant determination in sewage sludge were never questioned.

2. Materials and methods

The data presented in this study are calculated based on OC contents found in campaigns carried out in 2013-2014 and published in Mailler et al. (2017).

2.1. Sludge treatment plants monitored

STPs from three SIAAP WWTPs were sampled: Seine Aval, Seine Centre and Seine Gresillons. They treat sludge produced by the treatment of wastewater from Paris conurbation. The Seine Centre plant has a treatment capacity of 240,000 m³ of wastewater per day. Its sludge treatment consists in dewatering of mixed biological and physicochemical

settling sludge by centrifugation (Guinard D6LP30HBHF screw decanter) with cationic polymer addition (≈ 5.5 kg polyacrylamide polymer/ton dry matter (DM)) resulting in a production of almost 21,000 tons DM of centrifuged sludge per year (SIAAP source). The residence time in screw decanters is very short, estimated to be several minutes. The centrifuged sludge is then incinerated. The Seine Aval plant has a treatment capacity of 1,700,000 m³ of wastewater per day (biggest in Europe) and produces more than 55,000 tons DM of treated sludge per year (SIAAP source). After thickening, sludge is digested in twostage mesophilic (37°C) anaerobic digesters, which transforms an important part of organic matter (about 40%) into biogas, with a total residence time of 15-20 days. Then digested sludge is dewatered by high pressure and temperature cooking followed by press filtration (Porteous process). The Seine Gresillons plant has a treatment capacity of 300,000 m³ of wastewater per day. Sludge treatment consists in a thickening by centrifugation (GEA UCD-536-50-34 screw decanter) with cationic polymer addition (≈ 4.5 kg polyacrylamide polymer/ton DM), an anaerobic digestion (mesophilic or thermophilic digesters), a dewatering of digested sludge by centrifugation (GEA UCD-536-50-34 screw decanter) with cationic polymer addition (≈ 10 kg polyacrylamide polymer/ton DM) and then a thermal drying (thin film rotating dryer VOMM or belt dryer ANDRITZ) step. The residence time in thermal dryers is very short, estimated to be several minutes. A part of the thermal drying process used in this plant operates at a high temperature (255-280°C for thin film rotating dryer VOMM) compared to conventional dryers used for the other part of thermal drying (125°C)

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for belt dryer ANDRITZ). The characteristics of each WWTP and STP, and the full layout of each STP, are given in supporting material of Mailler et al. (2017).

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2.2. Data set and monitored pollutants

Different types of sludge have been sampled (punctual samples in the storage tank considering the estimated residence time in the processes) within the three studied STPs in April-May 2013 and April 2014 (3 to 4 consecutive weeks in 2 consecutive years in order to assess the OC content variability). The complete sampling strategy is described in Mailler et al. (2017). In this second paper, the OC concentration data on raw sludge and centrifuged sludge samples (n = 7), as well as residual water from centrifugation (n = 3), from Seine Centre WWTP are considered and treated together to evaluate the mass balance of OC in the dewatering centrifugation process. Sampling considered the very short residence time. Similarly, OC data on raw and digested sludge samples (n = 7) from Seine Aval, that were sampled considering a solid retention time of 16 days, are considered together to evaluate the mass balance of the mesophilic anaerobic digestion process. Finally, OC data on centrifuged sludge and thermal dried sludge samples (n = 6), as well as residual water samples from thermal drying (n = 3), from Seine Gresillons are considered together to evaluate the mass balance of the thermal drying process respectively. Similarly to centrifugation, the short residence time was considered during sampling.

71 emerging OC, including PHP (n = 18), hormones (n = 6), PFA (n = 2), LAS (n = 4), PAH

(n = 13), PCB (n = 20), AP (n = 4) and PAE (n = 4), have been analyzed in sludge samples (inlet and outlet of processes) and in residual water samples from centrifugation and from thermal drying. The 4 PAE and t-OP were only measured in 2014. The 66 other compounds were analyzed in 2013 and 2014 just after sampling. Due to its high DM content, the liquid samples from centrifugation were either extracted entirely or separation of dissolved and particulate phases was made before extraction of each sub-sample. The liquid samples from thermal drying were directly extracted. For sludge samples, the total OC concentration was measured after freeze drying and grinding of the raw samples. Briefly, analytical methods are based on specific extraction methods (QuEChERS, Accelerated Solvent Extraction, Solid Phase Extraction) coupled to Liquid Chromatography-tandem mass spectrometer detector for PHP and PFA, to Liquid Chromatography-fluorescence detector for PAH and nonylphenols, to Gas Chromatography-mass spectrometer detector for hormones, phthalates and octylphenols and to Gas Chromatography-high resolution mass spectrometer detector for PCB. The complete list and analytical procedures information are given in Mailler et al. (2017).

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2.3. Mass balance assessment

A data validation strategy based on mass balance assessment was performed on the three sludge treatment processes. First, the daily process DM performances during sampling campaigns were compared to 3 to 4 months of operation to check that days of sampling are

representative of a normal operation of processes. Then the DM mass balances were calculated in order to check the efficiency of processes and the representativeness of punctual samples. The verification of DM mass balance is a crucial prerequisite to validate the representativeness of samples, considering that getting reliable data on operational parameters was very challenging at industrial scale. If DM mass balance is validated, OC mass balance is calculated for compounds detected in both input and output sludge samples during at least one campaign considering the following rules. Centrifugation: The raw sludge mass load was calculated using the sludge daily volumetric flow (m³/d) and the average DM content (g DM/L) of the day measured by the WWTP in routine. The mass load of centrifuged sludge was estimated using the daily DM flow (t DM/d) of the campaign under consideration, the wet mass flow (t wet sludge/d) and the average DM content (g DM/L) of the day measured by the WWTP in routine. To quantify the mass load of the residual water, its daily volumetric flow (m³/d) was first calculated considering a DM balance in the centrifugation process (difference between DM mass flow in raw sludge and centrifuged sludge) and then multiplied by the average particulate content measured in the residual water by the WWTP in routine. SI Table 2 explains the calculations made to assess OC mass loads in the centrifugation process of Seine Centre. Once the OC mass loads were calculated, OC mass balances were established: considering the sampling and analytical uncertainties, a mass balance was considered correct / valid if the difference between input and output mass flows was +/- 30%.

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Thermal drying: the mass loads of centrifuged sludge were calculated using the sludge daily volumetric flow (m³/d) and the average DM content (g DM/L) of the day measured by the WWTP in routine. The residual water mass loads were calculated using the water daily volumetric flow (m³/d) estimated by considering that only water is removed from sludge and using the difference of DM content between dried and centrifuged sludge and the sludge daily volumetric flow. Regarding the extremely low suspended solids concentrations content in the residual water compared to sludge, this simplification is acceptable. The mass loads of thermal dried sludge were calculated using their daily volumetric flows (m³/d) estimated as the difference between the centrifuged sludge and residual water volumetric flows. SI Table 2 explains the calculations realized to evaluate OC mass loads in the thermal drying process of Seine Gresillons. Same criteria of +/- 30% was used to say if OC mass balance is correct or not. Anaerobic digestion: mass loads were not calculated for this process considering that the volumetric flow is equal in the inlet and outlet of the anaerobic digestion unit. Then, concentrations in ng/L were considered to evaluate mass balances and removals of OC directly. Removals were calculated for each sampling period because sampling took into account the HRT of the unit. If output concentrations were lower than the LD, this latter was used to calculate the removal.

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3. Results and discussion

Three different types of sludge treatment process were assessed in this work. The way to assess their role in the fate of OC was fully different. Indeed, centrifugation and thermal drying are physical processes implying respectively phase separation and water elimination. Except volatilization and thermal degradation that could theoretically occur in thermal drying, no removal of OC should be observed on both processes considering the very short residence time, only OC distribution between aqueous and solid phases (OC content was measured in both phases) should be observed with correct mass balances. For the biological process, as anaerobic conditions may favor OC transformation, removals can be calculated.

3.1. Mass balance of OC in dewatering by centrifugation

3.1.1. DM mass balance

Due to the lack of data on the volumetric flow of the residual water, we assumed a correct DM mass balance to calculate it. In this kind of process with very short residence time, loss of DM is assumed negligible which guarantee the feasibility of our calculation. Moreover, this sampling strategy and methodology of calculation were validated by Mailler et al. (2014) demonstrating a conservative mass balance for trace metals, i.e. for zinc with 731 \pm 133 and 729 \pm 142 mg/kg DM respectively for inlet and outlet sludge samples as no zinc was found in the aqueous phase. In order to validate the process performances, the conventional operation parameters measured during the sampling periods were compared to the average data

measured in routine by SIAAP in the corresponding three months (Table 1). The results confirmed the normal operation of the process during the sampling period and the representativeness of samples. This process is very repeatable in each period with a stable DM concentration factor of 5.4 ± 0.3 in 2013 and very efficient with an aqueous phase containing less than 4% of the initial DM in 2013 (Table 1). Similar stability and efficiency were observed in 2014. Mass balances were thus calculated for the detected OC.

Table 1 Performances of the centrifugation processes during the sampling period and comparison with average values from the daily operation monitoring of the facilities on the corresponding 3 or 4 months' period. CS for centrifuged sludge, RS for raw sludge, CW for residual water of centrifugation

		Proportion of DM		
Parameter	Concentration factor	lost in residual	Q_{out_CS}	Q_{out_CW}
		water		
		TSS _{out_CW} /DM _{in_RS}	Q _{in_RS} /concentration	
Equation	DM_{out_CS}/DM_{in_RS}	× 100	factor	Qin_RS - Qout_CS
Unit	-	in %	in m ³ /d	in m ³ /d
April, 9 th , 2013	5.4	2.0	245	1067
April, 30 th , 2013	5.3	3.5	317	1360
May, 14 th , 2013	5.3	5.3	285	1222
May, 22th, 2013	5.9	3.1	325	1583
June, 11 th , 2013	5.3	5.0	296	1281

April, 1st, 2014	6.0	2.7	213	1079
April, 8 th , 2014	4.8	1.9	453	1715
April, 15 th , 2014	4.5	3.2	405	1429
Average ± STD (5 d	5.4 ±0.25	3.7 ±1.3	293 ±31	1302 ±189
of sampling in 2013)	5.4 ±0.25	3.7 ±1.3	293 ±31	
$Average \pm STD \ of$				
daily operation data	5.2 ±0.4	3.2 ±1	264 ±64	1111 ±270
in	3.2 ±0.4			1111 ±2/0
april, may, june 2013				
Average ± STD (3 d	5.1 ±0.8	2.6 ±0.6	357 ±127	1407 ±318
of sampling in 2014)	3.1 ±0.8	2.0 ±0.0	33/ ±12/	1407 ±316
$Average \pm STD \ of$				
daily operation data	5.3 ±0.5	3.6 ±2.1	269 ±84	1131 ±318
in march, april, may,	3.3 ±0.3			1131 ±316
june 2014				

3.1.2. OC mass balance

For several compounds (n=11), essentially PHP and PFA, no mass balance could be calculated because OC concentration in either raw sludge or centrifuged sludge or both was lower than the LQ. This was the case for ivermectine in both year and for azithromycin, cefoperazone, escitalopram, glybencyclamide, lidocaine, miconazole, PFOA, propranolol, sulfamethoxazole, verapamil, in 2014. Five of these compounds (azithromycin, lidocaine,

PFOA, propranolol, verapamil) were quantified in the residual water whereas they were not detected in the inlet or/and outlet of the centrifugation process. In this water matrix, the LQ was determined really low with a lower matrix effect than in the solid matrices, explaining their quantification.

An OC mass balance was calculated for all the sampling campaigns for 59 (2013) and 56

(2014) compounds, and for 6 (2013) and 4 (2014) more compounds for at least one campaign (SI Tables 3 and 4). The reason for no calculation was the same as mentioned above. Once validation of mass balances was assessed (SI table 4), the compounds were grouped according to the frequency of calculation of a correct mass balance (Table 2) and this parameter was plotted against physicochemical properties in order to explain the difficulties to get correct mass balance (Fig 1).

Table 2 Classification of the compounds according to the frequency of calculation of a correct mass balance and according to the range of concentration (if inferior or superior to 5 times the LQ) on the centrifugation and thermal drying processes. Underlined compounds are those found in the residual waters of each process. For families like PAH or PCB, calculation was also done on the sum (Σ) of all congeners. For compounds abbreviation see SI Table 1.

Centrifugation	Thermal drying

Frequency > 0.7 (group 1)

 μP_{in} and $\mu P_{out} > 5 \times LQ$ NP, BBP, LAS-C11, PFOS, NP, BaP, BbF, ΣPAH (13),

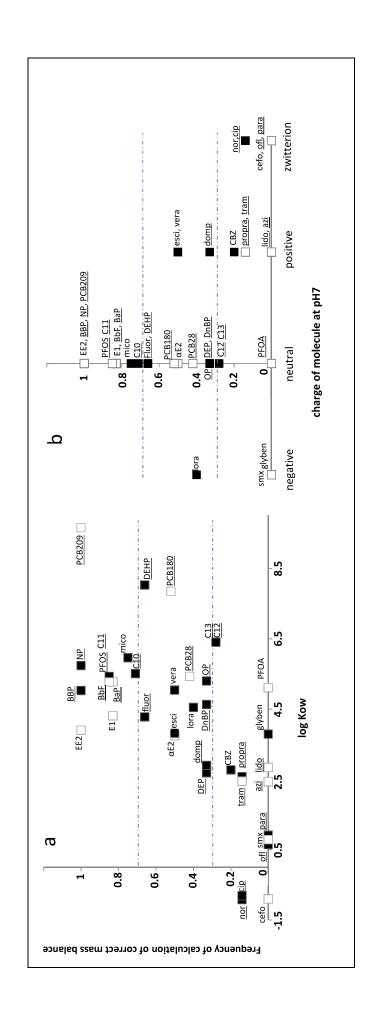
	mico, <u>LAS-C10</u>	Fluor, LAS-C10, LAS-C13	
<5 × LQ	EE2 DIE E1 D-D DCD200	E1, PCB180, PCB209,	
√J ^ LQ	EE2, <u>BbF</u> , E1, <u>BaP</u> , <u>PCB209</u>	∑PCB (20)	
	0.3 <frequency<0.7 (group="" 2)<="" th=""><th></th></frequency<0.7>		
	Fluor, DEHP, esci, vera, lora,	cbz, cip, nor, ofl, para, <u>azi</u> ,	
μP_{in} and $\mu P_{out}\!>\!5\times LQ$	t-OP, DEP, DnBP, domp	mico, LAS-C11, LAS-C12,	
		<u>tram</u>	
<5 × LQ	αE2, <u>PCB28, PCB180</u> ,	αE2, EE2, PCB28	
√J ∧ LQ	$(\underline{\sum}PCB(20), \underline{\sum}PAH(13))$		
	Frequency < 0.3 (group 3)		
	LAS-C12, LAS-C13, cbz,	nuonuo osoi alvihon lono	
μP_{in} and $\mu P_{out} > 5 \times LQ$	propra, cip, nor, para, glyben,	propra, esci, glyben, lora,	
	<u>ofl</u>	vera, domp	
25 v I O	tram, PFOA, lido, cefo, azi,	DEOC 1:1.	
<5 × LQ	smx	PFOS, lido	

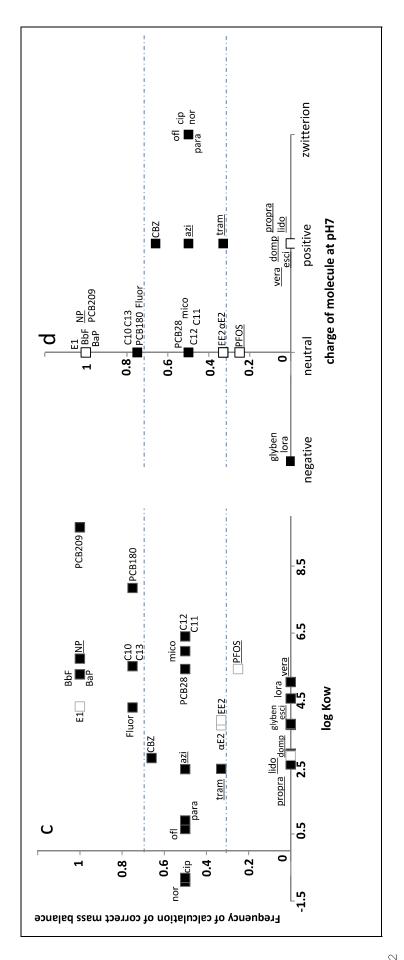
Fig 1 Impact of both logKow and charge at pH 7 of compounds on the frequency of calculation of correct mass balances in centrifugation (a and b) and thermal drying (c and d). Black square for compounds that are always quantified above 5 times the LQ, white square if one measure is 298 299

below 5 times LQ. Underlined compounds are those found in the residual water of centrifugation or thermal drying. For compounds abbreviation

see SI Table 1.

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As previously discussed, the centrifugation process was very conservative for DM and trace metals. As shown in figure 1 and table 3, mass balances were also frequently correct for 11 compounds including NP, some hormones, some PAH and PCB, BBP (one of the phthalates with the highest level in sludge), 2 LAS, one PHP (miconazole) and one PFA (PFOA). These compounds from group 1 are all neutral compounds together with log Kow above 3.5. They present concentration either lower than $5 \times LQ$ (hormones) or greater than $5 \times LQ$ (NP). They also present fluctuating concentrations in raw sludge (BbF load range between 2 and 28 g/d, SI table 3). Eight of these compounds were quantified in the residual water at less than 5% of the inlet mass of the centrifuge decanters, mainly because they were sorbed on the remaining suspended solids; this underlines the low contribution of this aqueous fraction to their mass balance assessment. However, if the inlet mass is high, this OC fraction present in the residual water may represent a non-negligible input on the total load of the WWTP; this point has to be considered to assess the global performances of WWTP. The group 2 was characterized by an intermediate frequency (between 30 and 70%) of calculation of a correct mass balance. Among the compounds, some persistent hydrophobic compounds like PCB, DEHP, fluor and t-OP, had sometimes incorrect mass balances (SI table 4) due to higher mass in the outlet. The other OCs of group 2 are 4 PHP (escitalopram, verapamil, loratidine, domperidone) with concentrations greater than 5 times the LQ and presenting positive and negative mass balances, the same was observed for aE2 whose concentrations were however close to the LQ. In this group 2, compounds have high logKow

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between 2.5 and 8 and only 4 compounds out of 12 are charged.

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The group 3 with mass balances frequently not correct includes compounds that are more hydrophilic (11/15 with logKow<3), mainly charged (12/15) and have concentrations close to the LQ (6/15) (figure 1). This unbalanced mass was mainly due to higher mass load in inlet than outlet. Moreover, in this group, compounds like tramadol, propranolol, lidocaine and azithromycin were quantified in the residual water at high proportion up to 90% of the outlet for azithromycin, meaning that it is essential in this case to measure the compounds in the residual water. It also includes 4 hydrophobic OCs: the two highly concentrated high molecular weight LAS and PFOA, which were very poorly quantified, and the negatively charged glybencyclamide (with quantification of higher mass loads in the outlet). The mass balance of the 3 hydrophilic well quantified fluoroquinolones was never correct even if they are easily sorbed to sludge thanks to their charge (cation exchange and bridging). Carbamazepine and propranolol, hydrophobic compounds and positively charged at pH7, accurately quantified at 100 times the LQ, are also present in the group 3 and presented higher mass load in the inlet than in the outlet of the centrifuge decanters. For the last 5 PHP and PFOA, their presence at concentration very close to the LQ may explain that the mass balance was never correct. Spearman (non-normal distributions) correlation tests (XL Stat software) were performed and indicated significant positive correlations (p-value < 0.05) between logKow and the frequency

of calculation of correct mass balances within screw decanter, both for compound detected

below (r = 0.673) and above 5 time the LQ (r = 0.675). For the compounds charge, a negative correlation is significant only for compounds detected above 5 time the LQ (r = -0.426). Other properties like molar mass or topological surface area were tested but they did not provide further explanation.

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These results demonstrated the complexity to establish OC mass balance in the centrifugation process for compounds with concentration near the LQ (due to uncertainties). The presence of positive or/and negative charges in the molecules could also explain this difficulty of mass balance calculation. Indeed, the addition of cationic polymers during centrifugation may change the interactions between OC and organic matter, that could induce changes in extracting yield and analysis (matrix effect) that are not compensated by the addition of deuterated compounds (no ageing). Analytical practices should then be adapted to consider this, with for example method validation for each type of matrix. In addition, the sampling strategy should be questioned, even if it was validated based on DM and by comparing to the average operation in a longer period (insufficient validations). Indeed, sludge is a very variable and heterogeneous material and OC concentrations in sludge are then very variable. The punctual samples considering the sludge residence time in the process could be sufficient for either highly concentrated or stable parameters such as DM, trace metals or some hydrophobic OC but insufficient for trace and less stable parameters such as hydrophilic and charged OC whose interactions with matrix would change according to the applied conditions.

The implementation of more thorough sampling strategy should be evaluated in order to reduce sample heterogeneity by (1) implementing 24h composite sampling systems adapted to sludge, (2) increasing the number of composite samples for one batch, , and (3) collecting multiple samples from a type of sludge taken from various sampling spots to get a more representative composite punctual sample.

3.2. Mass balance of OC in thermal drying

3.2.1. DM mass balance

The drying process was very repeatable in each sampling week of 2013 with a stable DM concentration factor of 3.5 ± 0.2 and its performances were very similar to those calculated on a three months' period (Table 3, SI Table 5). The calculated DM mass balance was correct with a difference between inlet and outlet always lower than 30% (SI fig 1). The TSS concentration in the residual water, only measured in 2014, underlined a very high efficiency of separation with residual water containing extremely low concentrations of DM. The measured and calculated operation data presented a higher variability in 2014 compared to 2013. In 2014, the first campaign could not be considered due to the impossibility to obtain the volumetric flow data for the centrifuged sludge inlet. Moreover, the DM mass balance calculated for the second campaign was not correct with a difference between inlet and outlet DM greater than 57%, highlighting a clear sampling representativeness issue. For these

Table 3 Performances of the thermal drying processes during the sampling period and comparison with average values from the daily operation monitoring of the facilities on the corresponding 3 or 4 months' period. nm, not measured; TDS for thermal dried sludge; CS for centrifuged sludge; TDW for residual water of thermal drying process.

		Proportion of DM		·
Parameter	Concentration	lost in residual	Q_{out_TDS}	Q_{out_TDW}
	factor			
		water		
Equation	DM /DM	TSS_{out_TDW}/DM_{in_CS}	0 0	$Q_{in_CS} \times [DM_{out_TDS}]$
Equation	DM_{out_TDS}/DM_{in_CS}	× 100	$Q_{in_CS} - Q_{out_TDW}$	$-DM_{in_CS}$
Unit	-	in %	in m ³ /d	in m ³ /d
April, 9 th , 2013	3.4	nm	19.5	33.4
April, 30 th , 2013	3.6	nm	14.6	27.1
May, 14 th , 2013	3.3	nm	18.7	34.1
May, 22 th , 2013	-	nm	-	-
June, 11 th , 2013	3.6	nm	21.8	44.0
April, 1st, 2014	2.0	0.010	-	-
April, 8th, 2014	2.6	0.025	13.5	17.0
April, 15 th , 2014	3.6	0.003	32.5	41.8
Average ± STD (5				
d of sampling in	3.5 ± 0.2	-	18.6 ± 3.0	34.6 ± 7.0
2013)				

Average ± STD of

daily operation data

in

april, may, june

 3.5 ± 0.3

 2.7 ± 0.8

 3.5 ± 0.4

Average ± STD (3

2013

d of sampling in 2014) $Average \pm STD of$

daily operation data

in

march, april, may,

june 2014

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3.2.2. OC mass balance

For 5 compounds in 2013 (cefoperazone, ivermectine, sulfamethoxazole, PFOA, NP2EO) and 15 compounds in 2014 (azihromycin, carbamazepine, cefoperazone, domperidone, escitalopram, glybencyclamide, ivermectine, lidocaine, loratidine, miconazole, propranolol, sulfamethoxazole, tramadol, verapamil, PFOA), no OC mass balance could be calculated because OC concentration in either centrifuged sludge (inlet of thermal drying process) or thermal dried sludge or both were lower than the LQ. However, six of them (PFOA,

 0.014 ± 0.010

 $16.2 \pm 6.$

 28.5 ± 11.0

azithromycin, lidocaine, sulfamethoxazole, tramadol, verapamil) were quantified in the residual water extracted from the process even if not quantified in sludge, 4 of them were similarly quantified in the residual water of the centrifugation process. As previously stated, in such aqueous matrix, the LQ is very low with a lower matrix effect than in the solid matrix, allowing OC quantification at lower levels. This means they were present in the sludge but not quantified. An OC mass balance is calculated for all the considered sampling weeks for 51 (2013) and 56 (2014) compounds and for 10 more compounds in 2013 for at least one week (SI Table 3). The reason for no calculation is the same as previously mentioned. Once the validation of mass balances was assessed (SI Table 6), the compounds were grouped according to the frequency of quantification of a correct mass balance (Table 3). It is important to note that only one or two mass balances were calculated for some compounds due to data lower than LQ (azithromycin, estrone, glybencyclamide, loratidine, miconazole) or lack of data (phthalates). By comparison, more compounds presented correct mass balances (Table 3) in thermal drying process than in centrifugation process but trends observed by linking it to physicochemical properties were less interpretable (Fig 1). Spearman (non-normal distributions) correlation tests were performed and indicated a significant positive correlation (p-value < 0.05) only between logKow and the compounds detected above 5 time the LQ (r = 0.437). Due to possible volatilization during this thermal process, other properties like volatility (expressed by means of Henry constant or vapor pressure) were considered but it did

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not provide further explanation. The number of molecules quantified in the residual water is low (11/71) and their concentrations contribution to mass balances were low, the fraction being lower than 0.5% of the input mass except once for tramadol at 10% (SI Table 6). Compounds with correct mass balances included neutral persistent organic molecules such as all PAH, all PCB, surfactants, phthalates and one hormone. The group with intermediate frequency of correct mass balance gathered hydrophilic and hydrophobic compounds, either neutral, zwitterion or positively charged, mainly highly concentrated, except for hormones. For 7 negatively and positively charged PHP and PFOS, the mass balance was frequently not correct with mainly higher mass load measured in the output of the dryer (5/8), even for compounds with concentration largely above 5 time the LQ (6/8). In this group 3, 6 compounds were quantified in the residual water. It seems thus difficult to link the OC behavior during drying process to their physicochemical properties except for very hydrophobic and neutral compounds for which mass balances are frequently correct. Some heat sensitive molecules could be altered during this process resulting in a lower outlet mass, which is only observed for 3 compounds in the group 3. For compounds that interact with organic matter thanks to cation exchange and bridging, the change in organic matter structure that could occur during this high temperature process may modify the interactions with OC and thus interfere with the extraction and analysis efficiency. This could explain the behavior observed for the charged compounds in the group 3, which were also quantified in the residual water.

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3.3. Removal of OC in anaerobic digestion

As shown in SI Table 7, average removals of both DM and volatile solids calculated in the 3 or 4 weeks of sampling were similar to removals observed in digestion during the corresponding three months' period, attesting of the good performances of the anaerobic digestion and its time stability. Considering that the volumetric flow is equal in the inlet and outlet of the anaerobic digestion unit, concentrations in ng/L were directly considered to assess mass balances and calculate removals of OC. Two (in 2013) and 4 (in 2014) samples were considered to calculate OC removal on the anaerobic digestion unit. For 2 compounds, glybencyclamide and ivermectine, no removal was calculated due to inlet and outlet concentrations near and lower than the LQ (SI Table 8). In 2014, 5 more compounds, PFOA, sulfamethoxazole, azithromycin, cefoperazone and loratidine also presented inlet and outlet concentrations under the LQ, the same for carbamazepine but only for one week. In 2013, for these compounds, removals could be calculated but concentrations remained low which suggests to consider these removals with caution due to high uncertainty. One compound, miconazole, presented inlet concentrations lower than the LQ but outlet concentrations largely above the LQ for the 4 weeks in 2014 whereas it was the contrary for the 2 weeks in 2013. Thus, its high removal calculated in 2013 has to be also considered with caution (Table 4).

Table 4 Class of removals of OC during anaerobic digestion. Underlined molecules are only present in one class of removal. In brackets: number of weeks in which removal was calculated out of 6 available data. For PAH (sum 13), PCB (sum 20), LAS (sum 4). *data on miconazole have to be considered with caution due to high removals in 2013 and negative mass balances in 2014. For compounds abbreviation see SI Table 1.

	Compounds with low	Compounds with high
Class of removal	concentrations	concentrations
In %	[OC] < 5*LQ	[OC] > 5*LQ
Removal < 30	E3(3), T(1), αΕΕ2(1), αΕ2(2), βΕ2(2)	para(1), cbz(1), ofl(1), nor(2), cip(1), LAS(2), NP2EO(2), <u>PAH(2)</u> , PCB(3), DEHP(1), BBP(2), DEP(1)
30 < Removal < 70	PFOA(2), smx(1), cefo(1), E1(4), E3(3), T(5), αΕΕ2(1), αΕ2(2), βΕ2(1)	PFOS(2), para(1), cbz(1), propra(1), esci(1), lido(1), vera(1), ofl(2), nor(1), cip(3), LAS(1), NP2EO(3), PCB(1), DEHP(1), BBP(2), DEP(1), DnBP(3)
Removal > 70	smx(1), <u>azi(1)</u> , cefo(1), <u>lora(1)</u> , αΕΕ2(3), Ε1(2)	PFOS(4), para(4), cbz(2), propra(5), esci(5), lido(4), mico(3)*, trama(5), vera(4), domp(6), ofl(2), nor(2), cip(1), DEP(1), DnBP(1)

For several persistent compounds (NP, LAS, PAH, PCB, αE2, βE2), all quantified greatly

above the LQ (except hormones), the outlet concentrations were often greater than the inlet ones, preventing to calculate a removal (SI Table 8). The higher outlet concentration for these persistent compounds could also be due to the issues previously presented regarding sampling and modification of matrix effect within the process. This confirms the necessity to deeply assess those questions prior to consistently analyze their fate in anaerobic digestion at fullscale. However, usually the difference between outlet and inlet concentrations was lower than 30%. Considering measurement and propagation errors, the mass balance could be considered correct meaning that these compounds are not or poorly degraded under anaerobic condition.. In addition, a negative mass balance could also reflect the production of molecules due to biological transformation. This is the case for NP (6 weeks/6 weeks), NP1EO (5/6) and t-OP (2/4). For these compounds, the higher output concentration may reflect the anaerobic transformation of parent compounds (NPnEO, n≥2), which was already observed by Murdoch and Sanin (2016) and Janex-Habibi et al., (2009), even if these transformation products have been also observed to be removed anaerobically (Patureau et al., 2008; Paterakis et al., 2012). For PAH, the same reason could be argued for the low molecular weight PAH (fluorene, phenanthrene) which presented negative mass balances and may be produced from the anaerobic transformation of high molecular weight PAH. For these latter, we observed either low removals (<30%, Table 4, SI Table 9) or negative mass balances comprised between 0 and -30%. PAH are thus poorly degraded under anaerobic conditions which is contradictory to some results obtained in lab-scale studies with spiked mixed sludge (Barret et al., 2010).

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These discrepancies in removals could be due to higher sample homogeneity (lab-scale reactors were fed with the same homogenized frosted/defrosted sludge) and PAH availability (spiking of sludge) at lab-scale compared to full-scale. PCB and LAS were also poorly removed by anaerobic digestion with either negative mass balances (for respectively 2 and 4 weeks) or removals lower than 40% and 50% respectively in other weeks. These results are consistent with data obtained at lab-scale by Benabdallah El-Hadj et al. (2007) for PCB and De Oliveira et al. (2010) for LAS. For phthalates, only DEP and DEHP presented negative mass balances for one or two weeks respectively. For other weeks, removals were below 70%, except for DEP and DnBP that were already shown to be easily degraded by comparison to DEHP and BBP (Fountoulakis et al., 2006; Abdel daiem et al., 2012). Several hormones (α EE2, α E2, β E2) presented negative mass balances at -30% for one (α EE2) or three weeks (β E2) but were also removed between 9-58% (α E2, β E2) and 17-79% (α EE2) for the remaining weeks (SI Table 9). Similar results were obtained by Gonzales-Gil et al. (2016) in lab-scale digesters with production or low removal of βE2, and higher removal of αEE2. However, E1 was always removed above 30%, which was not observed by Gonzales-Gil et al. (2016) nor by Noguera-Oviedo and Aga (2016) in manure anaerobic digestion. Testosterone presented moderate removals whereas E3 low to moderate ones. Anaerobic transformation of testosterone was only reported under anoxic conditions with nitrate and iron as electron acceptors (Shih et al., 2017) whereas E3 was also reported to be poorly degraded under anaerobic conditions (Singh, 2015).

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For the fluoroquinolones, only one mass balance is negative, the same for carbamazepine (SI Table 8). For the 5 remaining weeks, removals were calculated and varied between 1 and 78% for fluoroquinolones and 22-99% for carbamazepine (Table 4, SI Table 9). Those removals are quite surprising compared to literature (Liu et al., 2013; Alvarino et al., 2014; Gonzales-Gil et al., 2016). Indeed, fluoroquinolones are fully recognized as recalcitrant compounds and have been shown to be degraded under aerobic conditions (Dorival-Garcia et al., 2013; Rusch et al., 2019) but degradation under anaerobic conditions is not fully documented. From our data, anaerobic removal of ciprofloxacin was in average 43 ±17% (average on 5/6 weeks, range 28-72%) which is in accordance with the recent work of Do and Stuckey (2019) who observed 50-78% anaerobic removal of ciprofloxacin from synthetic wastewater with few detected transformation products. The removal range for carbamazepine was wide ($66 \pm 39\%$ average on 4/6 weeks) by comparison to the data in literature presenting carbamazepine as a very recalcitrant compounds under both aerobic and anaerobic conditions (Alvarino et al., 2014; Yang et al., 2016). However, several transformation products were identified by König et al. (2016) suggesting possible anaerobic reduction of this compound. Another PHP presented a wide range of removal, paracetamol, from 3 to 99% with 4 weeks in 6 being above 70%, classifying it as a highly removed compound. This compound has been shown to be well removed under aerobic conditions (Wu et al., 2012; Margot et al., 2015) and also anaerobic ones by Narumiya et al. (2013), Yang et al. (2016) and more recently Gonzales-Gil et al. (2019) who reported its degradation by enzymes extracted from anaerobic sludge. For

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both carbamazepine and paracetamol, the presence of an amide electron withdrawing group may explain their anaerobic reduction as for atenolol and diatrizoate (Ghattas et al., 2017). Among PHP with input concentration above 5 times the LQ, propranolol, escitalopram, lidocaine and verapamil presented mainly high removals (5 or 4 weeks/6) with output concentrations below the LQ. The same was observed for PFOS. Very few information on their anaerobic degradability is available in literature. Other betablockers have been shown to be moderately (metoprolol) to greatly (atenolol) removed. The removal mechanism implies amide hydrolysis (Gonzales Gil et al., 2019) for atenolol, and possibly the methoxy moiety for metoprolol like tramadol, naproxen and venlaflaxin (Ghattas et al., 2017). However, none of these groups is present on the structure of propranolol (Table 6). Amide hydrolysis could explain lidocaine degradation whereas dehalogenation might be expected for escitalopram. Odemethylation would be possible for verapamil, which was shown to be poorly removed by Yang et al. (2016). For PFOS, it seems that the very high polarity of the C-F bound renders these compounds non-degradable by dehalogenation (Ghattas et al., 2017). The high removal observed may be due to the presence of the electron-withdrawing group sulfonyl. Sulfamethoxazole and cefoperazone presented also moderate to great removals but these results have to be confirmed, their concentrations being close to the LQ. Sulfamethoxazole has been already shown to be well removed during anaerobic digestion (Narumiya et al., 2013; Gonzales-Gil et al., 2016; Alvarino et al., 2018), may be due to the strong electronwithdrawing group sulfonyl.

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Tramadol and domperidone were the 2 compounds presenting always high removals. Azithromycin and loratidine behaved similarly but concentrations were very close to the LQ making the confirmation of these results necessary. Tramadol was already shown to be anaerobically degraded via the *O*-demethylation of the methoxy moiety (Falas et al., 2016; Ghattas et al., 2017). No information was found for domperidone, but dehalogenation could be hypothesized as possible reaction. The Table 5 synthetized all the confirmed and hypothetic removal mechanisms discussed.

Table 5 Possible reactions explaining the OC anaerobic removal (only compounds with a high frequency of above 70% removal)

		Reaction/functional
Compounds		4. 1
(nofonon and)	Structure	group (italic when
(references)		hypothesis)
Tramadol	O_CH ₃	
(Falas et al., 2016;	HO CH ₃	O-demethylation
Ghattas et al., 2017)	CH ₃	
Domperidone (No	HN O NH	Reductive dehalogenation
reference)	N N CI	

Propranolol (No Secondary amine reference) (No Escitalopram Reductive dehalogenation reference) (No Lidocaine Amide (EWG) reference) Verapamil (No H₃CO. O-demethylation reference) PFOS (No reference) Sulfonyl Paracetamol (Narumiya et al., 2013; Wijekoon et -Amide (EWG) al., 2015; Yang et al., -Hydroxyl (EDG) 2016)

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CONCLUSIONS

This paper dealt with organic contaminants mass balances assessment applied to

physicochemical sludge treatment processes from WWTP sludge treatment trains. For dewatering by centrifugation and thermal drying, the complexity to get correct mass balances for several OC was deeply shown despite prior validation of the campaign and data. Those mass balances calculations were possible as 1) the representativeness of the sampling campaigns was confirmed when analysing global performances, 2) the sampling strategy and the calculation method allowed calculating correct DM mass balances which is a prerequisite step before OC mass balance calculation, and 3) the OC quantification method used is accurately validated based on current analytical good practices. However, those verifications and insurances were not sufficient in the case of many OCs. The frequency of correct mass balances relied on three parameters: (1) the measured content compared to the analytical method limit of quantification, (2) the hydrophobicity and (3) the charge of the compounds. Thus, for centrifugation, mass balances were more frequently correct for hydrophobic and neutral compounds, and inversely for hydrophilic and charged compounds. For these latter, concentrations close to the LQ render difficult such calculation. For thermal drying, similar trends were observed, however correct mass balances were obtained frequently (>0.7) for 47 compounds, while they were frequently not correct for 8 compounds. Like centrifugation process, it seems to be mainly explained by the hydrophobicity, charge and OC content compared to LQ. This paper also highlighted negligible to limited transfer to residual water for most of the compounds. For centrifugation, only propranolol and DEP presented important mass flow that has to be considered for the overall mass balances onto WWTP. The OC

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transfer to the residual water is even lower for thermal drying as only 11 compounds were quantified and for most of them the mass flow was really negligible (<0.5%) compared to the input mass flow, except for tramadol.

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These differences of mass balances determination between compounds could be explained by 1) the sampling practices that do not guarantee the representativeness of samples for OC and 2) the analytical practices that do not fully take into account the variations of matrix effect between inlet and outlet of processes. Thus, when assessing the fate of OC in full-scale sludge treatment plants, authors would recommend increasing the number of composite samples for one campaign in order to reduce sample heterogeneity instead of increasing number of sampling campaign. The revision of the sampling practices by implementing 24h composite sampling systems adapted to sludge is also a solution that has to be investigated. This underlined that results coming from one-time punctual samples, that are mainly used in conventional sampling strategy for sludge, have to be considered with caution. The difficulty to compare results from one site to another could also be explained by this type of sampling strategy. Moreover, it is important for charged and hydrophilic compounds to measure their presence in the residual water as it may significantly contribute to the mass balance and assess the returned mass to the WWTP inlet, which could be non-negligible. Establishing and validating correct mass balances of such physicochemical processes are also a prerequisite for WWTP modelling purposes. Finally, thorough evaluation of the matrix effect in both inlet and outlet sludge from a process should be generalized to make quantification of OC more

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Regarding anaerobic digestion, even if the SRT was considered for sampling, some compounds presented high range of removals. Consequently, compounds were grouped in three classes of removal instead of giving a single value of removal. It underlined also the importance of developing more rigorous sampling strategies. Compounds like domperidone, propranolol, escitalopram, lidocaine, verapamil and cefoperazone were shown for the first time to have high removals under anaerobic conditions at full-scale, which could be explained by their physicochemical structure.

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