



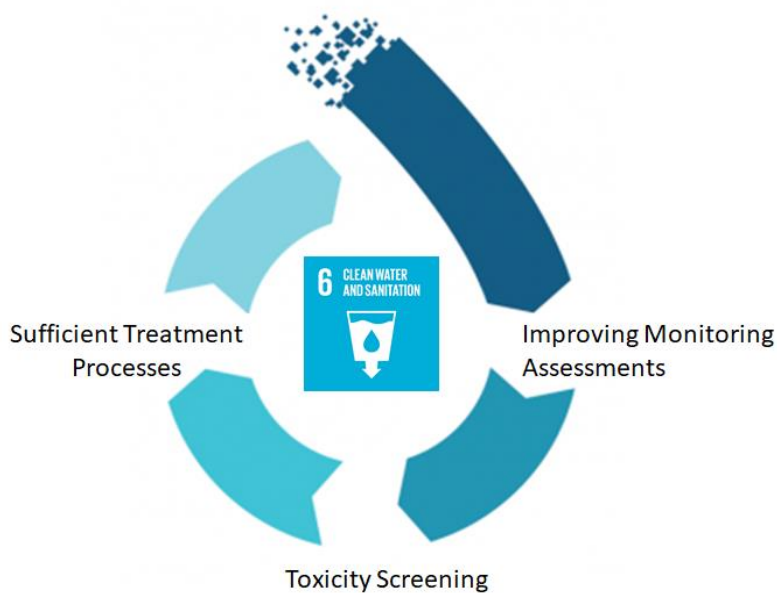
UNIVERSITÀ
DEGLI STUDI
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Università degli Studi di Torino

Doctoral School of Sciences and Innovative Technologies

PhD Programme in Innovation for the Circular Economy XXXIII Cycle

**Development of analytical methods using LC-MS/MS technique for CECs
detection and assessment of CECs in surface and drinking water samples
including toxicological screening**



Dimitra Papagiannaki

Supervisors:

Dr. Rita Binetti

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*To my parents Thanasis and Katerina.
For everything they gave up to make me who I am today.*

*Στους γονείς μου,
Θανάση και Κατερίνα*

Thesis Abstract

Water scarcity is one of the biggest threats society is facing around the globe and has been on the rise worldwide. With the rapidly increasing global population, the demands for clean and safe drinking water are increasing as well. However, the widespread pollution caused from both natural and anthropogenic factors is one of the main challenges that water companies have to face. For this reason, a more sustainable water management, adopting the Circular Economy concept, is necessary in order to improve water bodies quality, prevent waste and pollution and find efficient remediation technologies.

The main objective of this thesis was to develop advanced and accurate analytical tools for quantifying different classes of Contaminants of Emerging Concern (CECs) in different water bodies. For this purpose, two analytical procedures using Ultra High Performance Liquid Chromatography tandem with Mass Spectrometry were developed and fully validated. Both methods were applied successfully to two original explanatory studies that sought to quantify the levels of per- and polyfluoroalkyl substances, and pharmaceuticals and hormones in the water bodies of the Metropolitan Area of Turin, in Italy. Both monitoring assessments followed the principles of Green Analytical Chemistry, using fast and cost-efficient methods, and took into account the risk-approach by identifying the potential pollution sources. However, even if target monitoring assessments can provide fundamental information about the levels of pollution in an area, useful also for making treatment decisions, they are not sufficient in evaluating the holistic quality status of water bodies. This thesis is highlighting the importance of combining effect-based tools and non-target screening with conventional screening methods in order to better assess the water quality and better manage water. Hence, a non-target screening assessment using High Resolution Mass Spectrometry was done in order to reveal pollution patterns in the aquatic environment and identify potential novel contaminants.

Finally, taking into account the CECs' occurrence results after the monitoring assessments, the removal efficiency of different degradation methods was studied. More specifically, cost-efficient and environmental friendly techniques employing Advanced Oxidation Processes (AOPs) - followed by identification of byproducts in order to understand the degradation pathways and toxicological screening - as well as conventional treatment methods used in the Drinking Water Treatment Plant (DWTP) of Società Metropolitana Acque Torino (SMAT), were examined.

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List of Publications

P1. R. Binetti, P. Calza, G. Costantino, S. Morgillo, **D. Papagiannaki*** “Perfluoroalkyl Substance Assessment in Turin Metropolitan Area and Correlation with Potential Sources of Pollution According to the Water Safety Plan Risk Management Approach” *Separations* 2019, 6, 17.

P2. **D. Papagiannaki**, C. Medana, R. Binetti, P. Calza, P. Roslev “Effect of UV-A, UV-B and UV-C irradiation of glyphosate on photolysis and mitigation of aquatic toxicity” *Scientific Reports* 2020, 10, 20247.

P3. D. Palma, **D. Papagiannaki**, M. Lai, R. Binetti, M. Sleiman, M. Minella, C. Richard “PFAS degradation in ultrapure and groundwater using non-thermal plasma” *Molecules* 2021, 26, 924.

P4. **D. Papagiannaki***, S. Morgillo, G. Bocina, P. Calza, R. Binetti “Occurrence and human health risk assessment of pharmaceuticals and hormones in drinking water sources in the Metropolitan Area of Turin in Italy” *Toxics* 2021, 9, 88.

P5. **D. Papagiannaki**, D. Palma, A. Cedrino, M. Lai, M. Minella, C. Richard, R. Binetti “Identification of novel by-products after PFAS degradation with non-thermal plasma” *Manuscript*.

P6. I. Sciscenko, **D. Papagiannaki**, R. Binetti, C. Escudero-Oñate, I. Oller, A. Arques “Dissolved organic matter monitoring along with Turin drinking water plants employing EEM-PARAFAC”, Submitted to *Water*.

Abbreviations

| | |
|------------|---|
| ADI | Acceptable Daily Intake |
| AOPs | Advanced Oxidation Processes |
| ASTM | American Society for Testing and Material |
| ARGs | Antibiotic Resistance Genes |
| AFFF | Aqueous Film-Forming Foams |
| MEC | Average Detected Concentration |
| BW | Body Weight |
| CE | Circular Economy |
| CA | Concentration Addition |
| CECs | Contaminants of Emerging Concern |
| DDA | Data Dependent Acquisition |
| DIA | Data Independent Acquisition |
| DWI | Drinking Water Intake |
| DWTP | Drinking Water Treatment Plant |
| ESI | Electrospray Ionization |
| EPs | Emerging Pollutants |
| EDCs | Endocrine Disrupting Compounds |
| EPA | Environmental Protection Agency |
| EQSs | Environmental Quality Standards |
| EEA | European Environmental Agency |
| GCPs | Green Chemistry Principles |
| HCA | Hierarchical Cluster Analysis |
| HPLC-MS/MS | High Performance Liquid Chromatography coupled to Mass Spectrometry |
| HPLC | High Performance Liquid Chromatography |
| HRMS | High Resolution Mass Spectrometry |
| HV | High Voltage |
| IS | Internal Standard |
| ICH | International Conference on Harmonisation |
| LOD | Limit of Detection |
| LOQ | Limit of Quantification |
| EC50 | Median Effective Concentration |
| LC50 | Median Lethal Concentration |
| MeOH | Methanol |
| MDGs | Millennium Development Goals |
| MVDA | Multivariate Data Analysis |

| | |
|---------------------|---|
| N(L)OAEI | No (Low) Observed Adverse Effect Level |
| log K _{ow} | n-octanol-water partition coefficient |
| OLS | Ordinary Least Squares |
| PLS-DA | Partial Least Squared Discriminant Analysis |
| ppb | Parts per billion |
| ppm | Parts per million |
| ppt | Parts per trillion |
| PFCs | Perfluorinated Compounds |
| PFSAs | Perfluoroalkane Sulfonates |
| PFCAs | Perfluoroalkyl Carboxylates |
| PFASs | Per- and polyfluoroalkyl Substances |
| POPs | Persistent Organic Pollutants |
| PhACs | Pharmaceuticals' and Hormones' Active Compounds |
| PDBEs | Polybrominated Diphenyl Ethers |
| PP | Polypropylene |
| PNEC | Predicted No-Effect Concentration |
| PC | Principal Component |
| PCA | Principal Component Analysis |
| PSs | Priority Substances |
| pGLV | Provisional Guideline Value |
| PG | Pyrex glass |
| Q1 | First Quadrupole |
| Q3 | Third Quadrupole |
| QC | Quality Control |
| QF | Quantification Frequency |
| RT | Retention Time |
| RQ | Risk Quotient |
| SWATH-MS | Sequential Window Acquisition of all Theoretical Mass Spectra |
| SMAT | Società Metropolitana Acque Torino |
| SPE | Solid Phase Extraction |
| SDG | Sustainable Development Goal |
| TOF | Time-of-flight |
| TN | Total Nitrogen |
| TOC | Total Organic Carbon |
| VIP | Variables Importance in Projection |
| WWTP | Waste Water Treatment Plant |
| WEI+ | Water Exploitation Index plus |

| | |
|-----|---------------------------|
| WFD | Water Framework Directive |
| WM | Wide Mouth |
| WHO | World Health Organization |

Section I

Introduction

Chapter 1 Water at the heart of circular economy

Transition towards a Circular Economy (CE) has become a popular and important issue for environmental management in the last years [1]. CE is a sustainable development strategy that aims to boost resource efficiency and minimize waste production, while increasing economic and social benefits [2]. Actually CE is a rethinking of the current linear economic systems of "take, use, dispose", by promoting the employment of reuse, recycle, redesign, remanufacture, reduce and recover (the Rs approach) in a way to create a closed-looped system. Valuable materials from one product, at the end of its life, could be recovered and reused as source materials for the production of another product, minimizing resource inputs and waste [3]. The three main principles of CE include designing out waste and pollution, regenerating natural systems and keeping materials in use [4]. There is no clear evidence for the origin of the Circular Economy concept, with a variety of researchers contributing to it since the 1970s [3]. However, it was not until 2010 that its practices started being applied in different sectors, like waste management, sustainable design, food production, etc [5]. In the European level, the CE concept was adopted after two communications with titles "Towards a circular economy: a zero waste programme for Europe" [6], published in 2014, and "Closing the loop - An EU action plan for the Circular Economy" [7] published in 2015. Both communications highlighted the importance of adopting more rational use of resources and sustainable waste management in every branch of industry, and every group of materials and waste, while moving towards the CE [1]. As water is essential for human and ecosystems survival, and plays a significant role for industries as a carrier of both materials and energy, implementation of the CE principles in the water sector is necessary in order to improve the management of the already under pressure water bodies.

1.1 Water Scarcity and Sustainable Development Goals *SDGs*

Water is a finite resource - as its amount on Earth remains always the same and circulates through its natural cycle - of vital importance for living organisms, and indispensable for economic development, food production, and energy supplies [1,8]. Due to its high value for the society it should be used in a very sustainable way. However, the water crisis is already a reality in regions around the world, with studies reporting that 1 billion people lack access to clean water, and around one-third of the world population has already experienced severe water stress conditions [9,10]. It is estimated that by 2030 close to 3.9 billion people (almost half of the world's population) will be

living under conditions of severe water scarcity [11]. In the European level, the Water Exploitation Index plus (WEI+) showed that 33% of the population and 20% of the area has already faced severe water stress conditions (Figure 1), a percentage that is estimated to increase to 50% by 2030 [12].

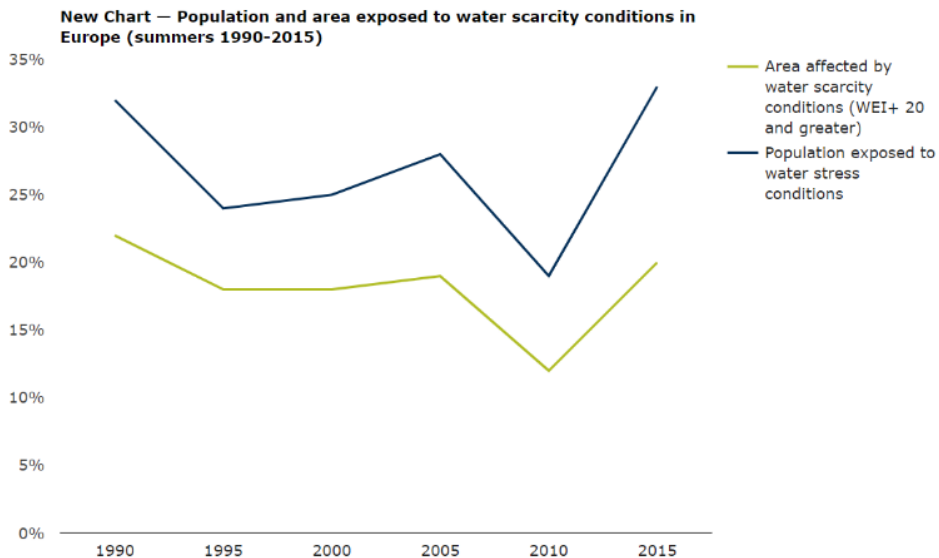


Figure 1: Water Scarcity Conditions in Europe (WEI+).

The continuously increasing world's population, that provokes increased human needs for water and requests for food, is responsible for the lack of sufficient available water resources on the demand side. Agricultural practices are increasingly growing in order to provide higher food yields, achieving a high water consumption, estimated as the 70% of the total one [13]. Their intensified land use with the overexploitation of water sources for irrigation purposes are adding in the risk of water scarcity, posing also challenges to the sustainable management of water. On the supply side, water scarcity is considered as a consequence of climate change. Nowadays, extreme floods or drought periods, alongside with warming temperatures affected significantly the hydrological cycle adding stress on water bodies [5]. Droughts can also have a negative impact on the biodiversity of an area. Different plant and animal species may go entirely extinct in extreme climate change conditions [14]. Furthermore, humanity's increasing water consumption is adding to the decrease of a region's biodiversity, which subsequently can affect humanity. The imbalance between demand and supply can be tremendous for the world population, since lack of clean water and sanitation can result in hunger,

poverty, diseases, and migration [15]. Therefore, protection of water resources for present and future generations is one of the most crucial environmental issues nowadays. A variety of solutions within the context of CE - such as integrated water resources management and wastewater reuse – have been proposed for addressing it. However, apart from quantity, maintaining the high quality of water bodies, is also vital for the population's water demands satisfaction. Extensive use of chemicals in agriculture alongside with uncontrolled discharge of big volumes of untreated industrial and domestic wastewater, undermine water quality posing threats to human health and aquatic ecosystems [16].

Humanity is called to take actions and make collective efforts in order to address these problems. In this context, the Sustainable Development Goals (SDGs) were adopted in 2015 by all the United Nations Member States, as a universal call for action to manage the social, economic, and environmental aspects of sustainable development by 2030 [17]. After the completion of the Millennium Development Goals (MDGs), the 17 SDGs and their 169 global targets were adopted in order to improve life's quality and achieve a better and more sustainable future for everyone (Figure 2) [18]. Water is directly and indirectly linked to all of the 17 SDGs, with SDG 6 being entirely devoted to clean water and sanitation. More specifically, SDG 6 aims to achieve access to safe drinking water and sanitation for all by 2030, improve water quality by reducing pollution and release of hazardous chemicals in water bodies for safe reuse, protection and restoration of water-related ecosystems. Finally, SDG 6 aims to address water scarcity by sustainable withdrawals and better water resources management, highlighting the dependance of social and economic development on the sustainable water management [19,20].



Figure 2: Water centric 17 SDGs for each sector.

1.2 Sustainable water management

In order to face water scarcity and meet the SDGs, different strategies have been published. In December 2019, EU published the European Green Deal [21], which aims to provide better solutions and conditions for achieving a more sustainable climate neutrality and circular economy by 2050. The strategy highlights the need to take actions for protecting and restoring ecosystems, use resources in a more sustainable way, reassure a toxic-free environment and improve human health [21,22]. In order to achieve these goals, EU published in March 2020, the new Circular Economy Action Plan [7], which points the importance of reducing waste externalities, maintain sources in use and regenerate the natural capital. For achieving a more sustainable management, which means to meet the current water supply needs without endangering the ability of future generations to do the same [23], these can be translated as [4,24]:

- optimization of the amount of water, energy and chemicals used in water systems' operation,
- maintain water, minerals, energy and chemicals in use and maximize their reuse,
- preserve the natural capital by river restoration, pollution prevention, and quality of water reassurance,

- ensure minimum disruption of natural water cycle from human actions.

In practice, different methods have translated these principles in the so-called Rs approach, including the concepts of:

- **reduce** water use and pollution at source,
- **reclamation** which is associated with the efficient removal of pollutants from wastewater using highly effective techniques,
- **reuse** of treated water for non-potable needs, as irrigation,
- **recycle** which is associated to reclaimed water from treated wastewater for direct potable use,
- **recovery** of materials, minerals and chemicals as potential resources,
- **rethink** of how to use natural resources in a more sustainable way.

Adoption of these concepts will eventually achieve reduction of the gap between availability and water demand by eliminating the adverse effects of pollution, identification of potential pollution sources, reduction of excessive water abstraction and waste production, better management of wastewater, and finally assurance of sufficient amount of good quality water to meet the needs of human and ecosystem [1,25].

1.3 Green Chemistry to enable circularity and sustainability

As already discussed, adopting a circular economy approach within a sustainable water management system is essential for increasing water resilience and preventing a global water crisis [23]. A key concept in order to enable sustainable development and circular economy is the implementation of Green Chemistry Principles (GCPs).

In general, green chemistry focuses on designing innovative products, with less consumption of natural sources and minimized waste disposal [26]. The twelve GCPs (Table 1) were created by Paul Anastas and John Warner, who tried to explain a more environmental-friendly process or product [27,28]. According to them, the pollution's prevention can be achieved before occurring by using new techniques and products that reduce or eliminate the use and generation of hazardous substances to human health and the environment [29].

Table 1. The twelve principles of Green Chemistry.

| The twelve principles of Green Chemistry (GCPs) | |
|--|--|
| 1. Prevent waste | 7. Use renewable feedstocks/materials |
| 2. Maximize atom economy in syntheses | 8. Avoid chemical derivatives |
| 3. Design less hazardous chemical syntheses | 9. Minimize waste by using catalytic reactions and not stoichiometric reagents |
| 4. Design safer chemicals and products | 10. Design chemicals and products to degrade after use |
| 5. Use safer solvents and reaction conditions | 11. Analyze in real time to prevent pollution |
| 6. Increase energy efficiency | 12. Minimize the potential for accidents |

Green chemistry focuses on a safe and sustainable design and development of materials and processes, by eliminating waste and improving energy efficiency, which are main components towards a transition to sustainable energy, resources and processes and circular economy [30]. Moreover, since its implementation plays an important role in environment, through pollution prevention and human health through elimination of hazardous compounds - while providing an economic sustainability - it can be a centric piece on addressing the SDGs [28]. Finally, green chemistry could have a significant contribution on the increased demand for development of innovative chemicals and technologies, that are cost-efficient and don't produce waste, in order to address the presence of CECs in the aquatic environment.

Chapter 2 Towards a toxic-free environment

Creating a toxic-free environment is one of the goals set by the European Green Deal towards a more sustainable and circular economy. In order to achieve it more actions for pollution monitoring, preventing its generation and finding better remediation solutions are necessary, alongside with revisions of policies and regulations.

Concerning water management and protection, till now EU has published different directives for achieving good environmental standards. In 2000, the Water Framework Directive (WFD) 2000/60/EC was adopted, aiming to prevent and reduce pollution, promote sustainable water use and protect the aquatic environment [31]. The EU WFD, established a watch list of priority substances (PSs) with high risks to the aquatic environment in order to set their monitoring assessment and establish their Environmental Quality Standards (EQSs). These values represent the maximum concentrations that a pollutant or a group of contaminants can be present in different environment compartments without posing threats to human health or ecosystem. The first list of priority compounds was established in 2008, by the Directive 2008/105/EC [32], including 33 PSs and 8 other pollutants. The choice of these compounds was based on data of their acute and chronic effects to ecosystem and human health. The EU Member States were required to set monitoring campaigns of the PSs and a good status of water bodies was determined if the detected concentrations were not exceeding the established Environmental Quality Standards. In 2013, the list was updated to 45 PSs and 8 other pollutants by the Directive 2013/39/EU [33], grouped as single or classes of substances and containing pesticides, industrial additives, pharmaceuticals, hormones, personal care products, food additives, surfactants, flame retardants and others. Moreover, the 2013/39/EU Directive proposed a first Watch List of 17 unregulated organic pollutants, not regularly included in monitoring programs but able to cause adverse effects on human health and ecosystems, and recommended their occurrence assessments in order to prioritize them and develop innovative treatment technologies for their abatement. This Watch List was later established by the Decision 2015/495/EU [34].

Although, there aren't any Environmental Quality Standards available for these pollutants, an evaluation system based on the frequency and the extent of exceedance of Predicted No-Effect Concentration (PNEC) in order to prioritize the compounds for monitoring and treatment has been proposed from the NORMAN Association [35]. In December 2020 the revised Drinking Water Directive 2020/2184 [36] has been

published, aiming to ensure the quality standards and increase the transparency for the consumer. More specifically, it highlights the need of better understanding, monitoring and evaluating the risks posed from newly detected chemicals in water bodies and their combined effects when present in mixtures. Moreover, it reports that risk assessment and management of the supply systems (Water Safety Plan approach), improvement of treatment techniques and increase of collaboration between stakeholders will allow better reduction of contaminants' release in the aquatic environment and subsequently their impact on drinking water sources.

2.1 Challenges

However, the biggest challenge of these strategies and regulative frameworks, that aim to ensure good quality of water bodies is the pollution originating from the so-called Contaminants of Emerging Concern (CECs).

Currently, water pollution is a serious problem that undermines the already scarce water resources. The majority of European countries rely on surface and groundwater for their drinking water needs, whose quality is affected from natural and anthropogenic factors. Metals, single organic ions and more complex organic molecules, as well as biological components can derive from various sources, such as natural disasters, agricultural run-off, industrial and domestic discharges, increasing population and economic growth, and can affect the quality of water bodies [8,37]. The existence of these factors in the aquatic environment represents a serious threat for human health and ecosystems. In recent years, the availability of robust and sensitive analytical methods and techniques has allowed the identification and detection of a wide variety of pollutants, with those deriving from anthropogenic sources commonly being referred to as micropollutants since their presence in water bodies is usually at trace levels (between few ng/L to some µg/L) [8]. These contaminants may be classified as legacy - whose toxic effects are already known and control measures have been established - or as Contaminants of Emerging Concern (CECs) [37]. This last class concerns compounds that are not currently regulated or included in routine monitoring programs, but are thought to have potential adverse effects to ecosystems and human health, and may serve as candidates for future legislations. CECs are not necessarily chemicals that have been recently introduced in the environment, they might as well include contaminants that have been present for a longer time, but their significance has not been evaluated since now or their occurrence was not known due to lack of adequate analytical techniques [38].

More than 1000 substances are considered as CECs or Emerging Pollutants (EPs) with a variety of sources being responsible for their occurrence in the aquatic environment, such as hospital effluents, landfill leachates, runoff from agriculture, and mainly industrial and domestic wastewater due to insufficient treatment techniques (Figure 3) [39,40]. Different studies have reported the inability of conventional treatment methods used in Waste Water Treatment Plants (WWTPs) to efficiently remove CECs, posing threats to the receiving environment [41,42].

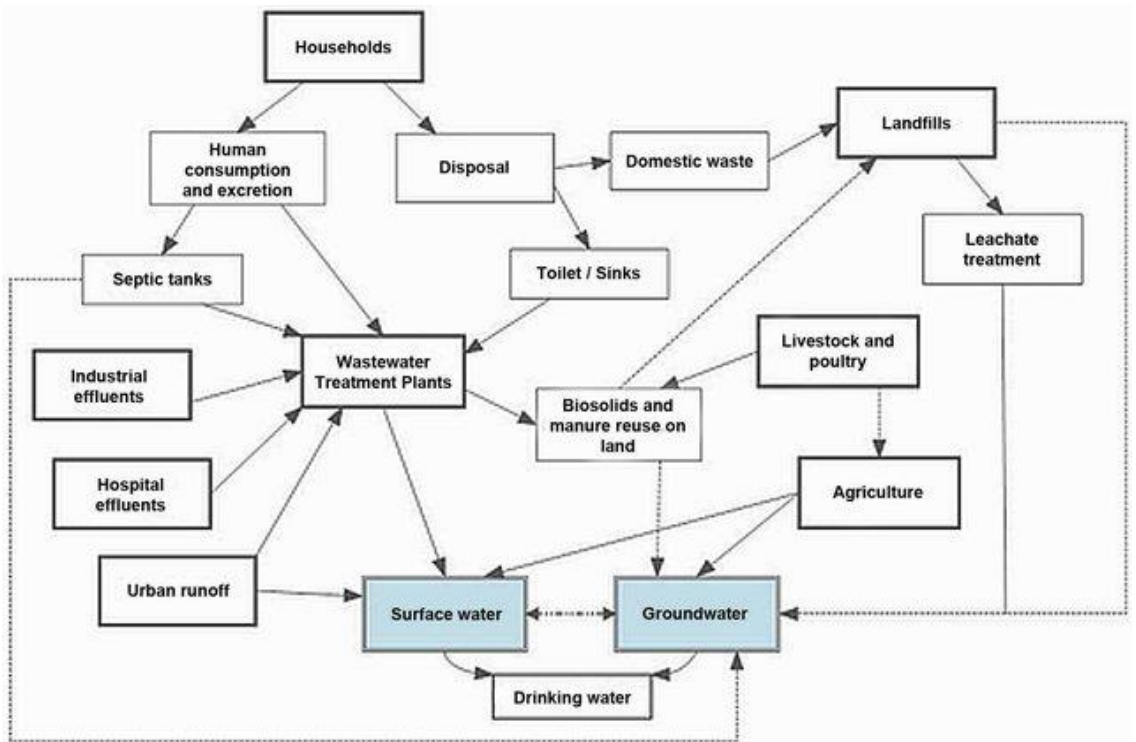


Figure 3: Sources of CECs in the aquatic environment [40].

As CECs are considered compounds included in the following 16 categories [43]: Pharmaceuticals, Personal Care Products, Sunscreens and UV/filters, Veterinary medicines, Pesticides and Herbicides, Persistent Organic Pollutants (POPs), like Perfluoroalkyl Substances (PFASs) and Polybrominated Diphenyl Ethers (PDBEs), Endocrine Disrupting Compounds (EDCs), including estrogens – both naturally occurring and synthetic-, Nanomaterials, Microplastics, Drinking water by-products, Antibiotic Resistance Genes (ARGs) and their transformation products.

Currently, in the European level, there are no existing regulatory limits for the majority of these substances. However, the European Environmental Agency considers that CECs or EPs should be closely monitored by taking into account the risk management approach and the combined exposure assessment, as they are increasingly being detected in the aquatic environment.

2.1.1 Per and polyfluoroalkyl substances (PFASs)

Per and polyfluoroalkyl substances (PFASs) represent a broad class of organic compounds, widely used over the past decades. They belong in the category of fluorosurfactants, since they host a substitution of hydrogen atoms by fluorine, in their carbon chain, building in that way the hydrophobic part of the surfactant [44,45]. PFAS are man-made organic compounds, with two major processes being followed for their production, the electrochemical fluorination and the telomerisation. However, since 2002 mainly the telomerisation method is applied. In general, perfluorinated compounds (PFCs) like perfluoroalkyl sulfonamides and fluorotelomer alcohols can be degraded naturally under aerobic conditions and transformed to perfluoroalkyl carboxylates (PFCAs) and perfluoroalkyl sulfonates (PFSAs) [46]. Based on the length of their perfluorocarbon chain, PFASs (PFCAs and PFSAs) can be classified as short and long chain compounds. In those with long chain are included PFCAs with eight or more atoms of carbon and PFSAs with at least six atoms of carbon [47,48] (Table 2).

Table 2. Categorization of PFASs according to their chain length [46].

| Perfluoroalkane Sulfonates (PFSA) | | Perfluoroalkyl Carboxylates (PFCA) | |
|--|---|--|---|
| Short Chain $n \leq 5$ e.g. PFBS | Long Chain $n \geq 6$ e.g. PFHxS, PFOS and PFDS | Short Chain $n \leq 7$ e.g. PFBA, PFPeA, PFHxA and PFHpA | Long Chain $n \geq 8$ e.g. PFOA, PFNA, PFDA, PFUdA, PFDoA, PFTrDA, PFTeDA, PFHxDA and PFODA |

PFASs represent a vast variety of molecules, with unique physicochemical properties, like extreme hydrophilic and lipophilic character or thermal and chemical stability. Most importantly, these compounds host one of the strongest chemical bonds (C-F) in their

carbon chain, which makes them very stable towards natural degradation or under conditions of heat, acid and oxidation [47,49]. Due to these characteristics, they are essential elements for a wide range of industrial and commercial operations. Some examples include stain- and water- resistant fabrics for clothes and carpeting, cleaning products, food packaging and cookware, paints and fire-fighting foams.

The widespread use of PFASs and their abilities to remain intact in the environment result in continuously increasing contamination levels in the water bodies. As their main sources are considered the industrial and municipal wastewater treatment plants, landfill leachate, dry and wet atmospheric deposition, soil and street surface runoff [48]. Different studies have reported their occurrence in various raw water sources, in concentrations between ng/L and few µg/L, and other environmental matrices, including food [50-55]. Accumulation of certain PFASs in humans and animals has also been reported, with serious concerns being raised about the potential health effects. Due to their stability properties, and the incapability of conventional treatment technologies to efficiently remove them from water, PFASs have also been detected in drinking water, raising significant threats of adverse human health effects [56,57]. Therefore, as mentioned in the EU Chemicals Strategy for Sustainability published in October 2020 [58], it is crucial to find and establish new and more effective degradation techniques, alongside with the prevention of their contamination.

2.1.2 Pharmaceuticals and hormones

Pharmaceuticals and hormones (PhACs) represent another big category of anthropogenic contaminants present in the aquatic environment [43], as a result of increasing population's growth and economic activities. These compounds can have adverse effects both to human health and aquatic ecosystems, like morphological anomalies, endocrine disruption and increasing antimicrobial resistance. In Europe, their use is continuously increasing, with 3000 compounds currently being active in the market [59]. Due to their large consumption, pharmaceuticals and hormones can reach the aquatic environment through different routes mainly including animal and human excretion, improper domestic or industrial discharge and landfill leaching [43], depending upon the substance and its properties. The last decades, a wide variety of studies has reported their presence in water systems of all types, in concentrations ranging from ng/L to few µg/L [60-66], raising concerns about their potential effects on human health, especially after a long term exposure to low level concentrations. However, none of these compounds is regulated in European Union, even though some

of them are included in the Watch List of the WFD, mainly due to the fact that they have insignificant biological effects at their low occurrence concentrations in the environment. Nevertheless, till now none of their assessments considered their effects when they bioaccumulate, or the synergistic interactions of simultaneous contamination with multiple compounds [67].

In order to face these problems, and fill knowledge gaps related to PhACs occurrence concentrations, risk assessments and monitoring of “hotspot” locations, the European Commission published in 2019 the European Union Strategic Approach to Pharmaceuticals in the Environment [6]. This approach proposes that Member States should promote the more careful use of PhACs, develop compounds that are not harmful for the environment, improve their environmental risk assessment and management of waste, broaden their environmental monitoring assessments and finally identify further knowledge gaps to be resolved, such as finding cost-effective remediation methods, towards the transition to a circular economy based environmental management.

2.1.3 Pesticides

Active ingredients from pesticides' formulation are among the most frequently detected organic micropollutants in the aquatic environment, due to their widespread use in agriculture and forestry [68]. They may contaminate the different water bodies types after application through spray drift, surface runoff, and soil leaching [69,70]. Depending on their physicochemical properties and more importantly on their chemical stability, pesticides may undergo natural degradation processes resulting to metabolites equally or even more toxic, considered as contaminants as well. The rapid population growth has increased the needs for food production, enhancing the need for pesticides to protect the yields from undesirable organisms. Pesticides are classified into different categories, based on their toxicity level, target application and chemical properties. Moreover, they can be categorized as organochlorines, organophosphorus, carbamates, pyrethrin and pyrethroids [71]. Advances in analytical techniques have allowed, since 1970s, the multi residual detection of pesticides in different water types and especially in drinking water, raising particular concerns about the greater impacts that can have on human health [72,73]. For this reason, pesticides concern few of the contaminants of emerging concern that -since 1998- are regulated in Europe. The published directives have set a maximum concentration of 0,1 µg/L for individual compounds, their metabolites and transformation products, and 0,5 µg/L for their total amount. Moreover, in a work programme launched in 1992, EU started a review process for all

active ingredients used in pesticides, based on human health, environmental, and ecotoxicity risk assessments and on food residues monitoring assessments and taking into account the requirements of the 91/414/EEC directive [74]. The outcomes are categorizing the active substances as banned, for “essential use” and authorized. Currently, 84 compounds are banned in EU including Atrazine, Alachlor, Dichlorvos, and Diazinon, which however are still detected in water bodies.

Another widely detected class of pesticides, whose approval in agricultural uses is a debated topic, is the glyphosate-based ones. Their active ingredient - N-(phosphonomethyl)glycine (Glyphosate) - affects a broad-spectrum of plants, making it a useful tool for agricultural, public and domestic uses [75-78]. At present, glyphosate has a massive global usage of about 700,000 tons per year, which has caused its ubiquitous occurrence in the aquatic environment, posing threats to humans and ecosystems [75,77-80]. A variety of degradation techniques has been proposed for removing glyphosate from the aquatic environment. However, there is still the need of finding green and cost-efficient techniques, with the potentiality to be used in large scale applications, such as treatment plants, within the circular economy context.

2.2 Improving Monitoring Assessments

In the European level, even if EU Member States achieved significant improvement of water bodies quality after adopting the requirements of the different directives, the current status of many European water bodies is still uncertain, regardless the advanced techniques and the numerous developed methods. According to the latest European Environmental Agency (EEA) Water Europe report [81] the majority of European water bodies faced difficulties in achieving a minimum of good ecological status as set by the different EU directives [1]. This fact highlighted the need of new measurements' and regulations' adoption, focusing mainly on reducing the negative effects of pollution that stress water bodies, and better managing of waste, towards the implementation of CE principles in the water sector [82].

Robust monitoring of water quality is a must for conserving the existing unpolluted resources and restoring the polluted ones within a sustainable water management, and a prerequisite for a safe regulation of chemicals [83]. Monitoring data are fundamental for identifying problems, taking decisions for reducing emissions and evaluating their effectiveness, as well as policy development and registration of chemical substances (ex. under the Regulation for Registration, Evaluation, Authorization and Restriction of Chemicals in Europe –REACH) [84]. Initiatives for new monitoring programmes, like the Watch List strategy published under the WFD, aim to fill the gaps concerning the occurrence and impacts of micropollutants in the European water bodies. However, these approaches, based on target analyses of a specific number of pollutants fail to evaluate the holistic quality status of the different water bodies. Therefore, improving monitoring assessments by combining them with risks evaluation on ecosystem and human health, taking into account also the simultaneous exposure to combined effects of chemical mixtures, is necessary [15].

More specifically, analyses targeting to specific analytes are not able to distinct between the non-existence and the transformation of a compound due to degradation processes. Instead, suspect and non-target screening approaches based on High-Resolution Mass Spectrometry (HRMS), are able to provide a more comprehensive picture of chemicals' presence in water bodies by addressing complex chemical mixtures analytically. Even the valuable information that can be obtained from non-target screening, it can never replace the target analysis but only trigger it after being the first step of the assessment [85]. Finally, bioassays are another available method able to address unknown mixture risks present in the water bodies [86].

2.2.1 Approaches for comprehensive monitoring

A more comprehensive monitoring of chemicals in the aquatic environment will be achieved after the combination of the three previously approaches. In this way, the detection of newly emerging compounds and their transformation products, as well as the identification of their toxicity effects, necessary information for future abatement plans, will be achieved [87]. Target, suspect and non-target screening analyses, based on modern Liquid or Gas Chromatography instruments coupled to HRMS, allow the detection of organic micropollutants in water at trace level concentrations (ng/L). Practically, this means that the combination of these three approaches based on HRMS is able to detect simultaneously complex mixtures of a huge variety of substances at a high level of sensitivity, while identifying unknowns as well [83]. Complementary to chemical analyses, bioassays are increasingly used as bioanalytical tools for water quality assessments in order to measure the combined effects of trace-level mixtures of chemicals [88]. Different biological tests, including cell models, receptors, tissues or small organisms can be used for measuring the effects of chemicals on various biological endpoints. Although bioassay data cannot be used for thorough risk assessments, can identify the presence of one or more compounds that cause effects on biological battery tests relevant to human health and the environment. As each chemical causes different effects, water quality monitoring requires a suitable and adequate set of bioassays, based on chemicals' human health effects [89] or environmental pressures [90]. Consequently, combining chemical and effect-based methods has the potential to improve water quality monitoring, by revealing the cause of an effect and the effect itself.

2.2.2 Sustainability in monitoring assessment

New monitoring approaches need to prove that they are cost-efficient and not time consuming, for being able to be applied in large-scale programmes. For this reason, a massive request for simpler, faster and lower-cost methods, in order to enable more extensive and efficient monitoring of a wide-range CECs in water bodies while minimizing waste, is growing [9]. Currently, the analytical techniques used in monitoring assessments include Liquid and Gas Chromatography coupled to Mass Spectrometry. Usually, a samples' pretreatment step is necessary prior to the analysis, in order to remove potential interferences and preconcentrate the sample. Solid Phase Extraction (SPE) is currently the most used technique, despite its drawbacks of high organic solvents' consumption or high costs [8]. Since improvements in monitoring assessments

are required, the replacement of traditional methods, with more sustainable and energy-efficient alternatives following the principles of Green Chemistry, is necessary and can result to great socioeconomic and environmental impacts. However, many of the opportunities for sustainability and greenness of an analytical method are directly related to the sample preparation process. During this step, the most efficient way to reduce waste is to introduce the sample to the analytical apparatus with no or little pretreatment. However, since in the majority of times, the no pretreatment step is impossible and the analytes must first be extracted from their matrices prior to analyses, sample preparation techniques should be optimized to reduce energy input, solvent use, waste production, and operator exposure as much as possible [91]. This can be accomplished by implementation of the Green Analytical Chemistry Principles, which promote the use of alternative eco-friendlier solvents and avoidance of toxic organic solvents, greener (less polluting) extraction procedures or complete elimination of the pretreatment step, restriction of the size and number of samples and the use of reusable samples extraction devices [91]. Moreover, the importance of methods' development which target to mixtures of multiple analytes and are based on simplified and automated analytical protocols, alongside with reduction of energy consumption and use of mathematical modelling of the data based on chemometrics is encouraged. Consequently, the greenness of the analytical methods promoted for large-scale monitoring assessments applications, is a fundamental step for increasing sustainability in the water management, alongside with the use of modeling tools for identifying contamination hotspots [92]. The combination of all these approaches, can provide the water sector with "smart" tools for more realistic monitoring and risk assessment of mixtures of CECs in water bodies, and help take decisions concerning treatment methods.

2.3 Remediation Methods

Except of the importance of pollution monitoring and preventing its generation at source towards creating a toxic-free environment within a more sustainable and circular economy, it is crucial to find more sufficient remediation solutions [37]. According to EU regulations and laws, treatment of wastewater is obligatory [1] and WWTPs represent the primary barrier of preventing water bodies' contamination. However, it is well-known that WWTPs are one of the most significant CECs pollution sources, since the conventional physical and biological treatment technologies that they rely on are unable to efficiently remove recalcitrant organic compounds, discharging them directly into the environment [93-95]. Drinking water treatment plants (DWTPs) impose another barrier

concerning the prevention of unintended human exposure to CECs. However, studies have shown that utilized conventional treatment techniques such as coagulation, flocculation, filtration and chlorination have low removal efficiencies as well [96,97], while enhanced treatment technologies like ozonation and activated carbon adsorption are able to efficiently remove them [96,98-100]. However, even though these techniques are effective, may require expensive chemicals and equipment. Moreover, as water sector is focusing on alternative water sources for meeting the current potable or non-potable needs while preserving the natural resources, is promoting water reuse. However, a not appropriate water treatment system for the specific reuse needs could lead to undermined and unreliable water quality (e.g. generation of disinfection byproducts in reclaimed water for potable needs) posing risks to human health and ecosystems [101]. For this reason, more efficient, cost-effective treatment techniques dealing with a vast variety of chemicals, while minimizing waste and energy consumption need to be developed. Among the alternatives, Advanced Oxidation Processes (AOPs) represent one viable option [39].

AOPs are eco-friendly chemical methods that rely on chemical and physical processes “in situ” generating highly reactive radical species (such as HO^\bullet and $\text{SO}_4^{\bullet-}$) for the oxidation of organic compounds [39], without producing waste. The generated radicals are optimal and powerful oxidants, since they do not generate additional waste, they are not corrosive for equipment, they are not toxic and they have a very short lifetime [102]. Their efficiency relies on the fact that the generated radicals are unselective and able to degrade a vast variety of compounds through non-selective reactions into smaller or inorganic molecules (Figure 4) [103]. The degradation of the organic contaminants occurs through hydrogen abstraction, electrophilic attack or electron transfer by the hydroxyl radicals.

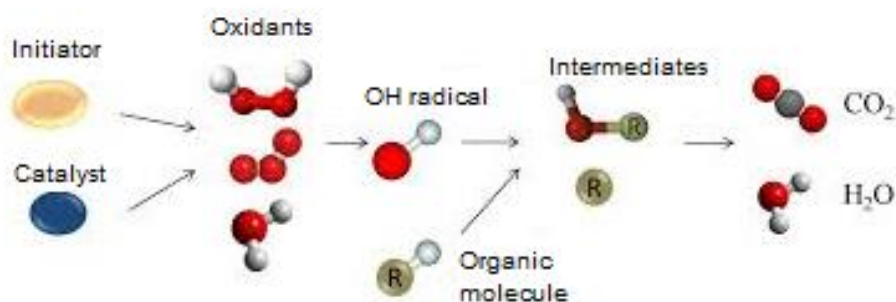


Figure 4: AOPs degradation mechanism when employing a catalyst.

AOPs include a vast variety of processes like Fenton, photo- and electro-Fenton processes, ozonation, UV photolysis, H₂O₂ and heterogeneous photocatalysis (Table 3) [39]. Processes involving treatment with O₃ and UV irradiation are the simplest processes proven to be able to efficiently degrade CECs and are already established in large scales applications, like DWTPs and water reuse facilities. Nonetheless, several AOPs such as electrochemical treatment, microwave, plasma and ultrasound related processes are continuously being studied by the scientific community [104].

Table 3. Different Advanced Oxidation Processes (AOPs) and their sources of radicals.

| Advanced Oxidation Processes (AOPs) | Source of Radicals |
|--|--|
| Photolysis | UV irradiation |
| O ₃ -based processes | O ₃ |
| | O ₃ /UV |
| | O ₃ /H ₂ O ₂ |
| | O ₃ /H ₂ O ₂ /UV |
| H ₂ O ₂ -based processes | H ₂ O ₂ /UV |
| | H ₂ O ₂ /Fe ²⁺ (Fenton) |
| | H ₂ O ₂ /Fe ³⁺ (Fenton-like) |
| | H ₂ O ₂ /Fe ²⁺ /UV (Photo-Fenton) |
| Heterogeneous photocatalysis | TiO ₂ /UV |
| | TiO ₂ /UV/H ₂ O ₂ |
| Sonochemical oxidation | Ultrasounds (water sonolysis) |
| Electrochemical oxidation | Electricity (water electrolysis) |

AOPs represent a group of sustainable remediation technologies that mainly follows the principles of Green Chemistry, with rapid reaction rates, mineralization of organics and a great potential to reduce toxicity of organic compounds. Moreover, they do not generate waste that need further treatment (such as the membranes, and activated carbon absorption). However, they have also some drawbacks, such as not being able to treat large volumes of low concentration pollutants (realistic conditions), and their efficiency being affected from the presence of organic matter or inorganic ions, which slow the rates of radicals' reactions. Finally, AOPs can generate several toxic

intermediates which sometimes are even more harmful than the parent compound, proposing the need of additional treatment for their removal [103,105]. For this reason, risk and toxicity of byproducts assessments should be included when planning and designing these processes as well, in order to identify and deal with their potential advantages and disadvantages, and set them in use [101].

2.3.1 UV photolysis

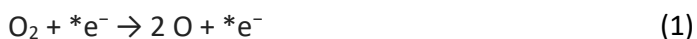
Photolysis of micropollutants present in water can occur directly via photon absorption in order to reach excited states, or indirectly through chemical reactions mediated by active oxygen species [37,102,106,107]. These can be translated to several potential mechanisms by which UV irradiation could mediate the transformation of CECs in water [106]. Firstly, UV irradiation can cause the direct photochemical dissociation of a compound, through photon absorption, followed by different chemical reactions such as bond cleavage and oxidation-reduction. Moreover, UV irradiation can interact with H₂O and O₂ and form reactive oxygen species including hydroxyl radicals ($\cdot\text{OH}$) that interact with the micropollutant and cause degradation. In any process that uses UV irradiation, the above mentioned reactions can take place. The performance of such processes depends on several factors, such as the intensity and wavelength of the UV irradiation, as well as the water matrix. As UV irradiation is considered a group of electromagnetic radiations with different wavelengths (e.g., 10-400 nm) and is classified into different categories such as UV-A (365nm), UV-B (302nm), UV-C (254nm), and Vacuum UV.

One of the main advantages of this AOP is that photolysis represents one sustainable alternative degradation method, since it is a non-invasive and free of added chemicals technique that can destroy the targeted CECs without generating waste or transferring them from one matrix to another [37]. Hence, UV-based AOPs are one of the most widely studied degradation methods for organic micropollutants. Pharmaceuticals, hormones and pesticides are some of the studied compounds classes reported in literature. These processes can promote the degradation of micropollutants via direct or indirect photolysis. However, UV doses used in conventional drinking water treatment lines (e.g., disinfection) are relatively low and not able to ease their removal. For this reason, different additives such as oxidants or catalysts may be added to improve the degradation processes mainly through the formation of radicals. Examples of such water treatment processes are the advanced oxidation process involving the

addition of hydrogen peroxide (e.g., UV-C/H₂O₂) and photocatalysis involving the addition of different catalysts (e.g., UV-C/TiO₂) [108-111].

3.2 Electrochemical Oxidation

Water plasma is another treatment technique utilizing non-selective and reactive radicals in order to degrade organic micropollutants. Even if water plasma is a technology firstly explored in the late 1980s, currently is attracting lots of attention for water treatment applications. Water-plasma technologies include both thermal and non-thermal plasmas used to remove biological or chemical contamination [112,113]. The main advantages of this process, is its ability to activate different reactions in a short time and at lower temperature (room temperature) using a relatively simple equipment [114]. Such reactions include the formation of highly oxidating species such as H[•], O and [•]OH radicals, free electrons and ozone, as well as oxidants like hydrogen peroxide (Equations 1-7) and ultraviolet light due to the plasma discharge light emission. Therefore, the degradation mechanism of pollutants is complex.



Different electrodes configuration has been explored for water plasma treatment methods, with the most used type being the corona discharge, which is created by applying a high-intensity electric field to sharp pointed electrode tips. Based on the polarity of the electrode, the corona discharge exists in positive and negative forms [114]. Water plasma has proven to be an efficient and cost-effective remediation technique – in short processing times [115]- of many recalcitrant compounds such as pharmaceuticals, pesticides, synthetic dyes and phenols without regenerating waste [116]. However, since its use is still in a lab-scale, further research about its sustainability in industrial-scale applications is needed.

Section II

Aims and structure of this
thesis

In the context of the AQUALity-ETN project, this PhD thesis aimed at the development of analytical methods for CECs' detection and assessment in different water matrices used for drinking water production including a toxicological assessment. The main research activities of this PhD project were carried out at the Research Centre of Società Metropolitana delle Acque Torino S.p.A (SMAT), the company in charge of the water cycle management in the Metropolitan Area of Turin (Piedmont, Italy). Further collaborations and secondment periods in other institutions within the consortium of AQUALity contributed in the conclusion of this thesis.

The main goal of this work was to evaluate a more sustainable way of water management systems within the context of Circular Economy (CE), by improving water quality monitoring assessments, after combining screening and identification techniques with higher detection sensitivity (even at trace levels) for a broader and more diverse group of Contaminants of Emerging Concern (CECs) with risk assessments, taking as well into account the cocktail effect of their mix. Based on the results of these assessments, the evaluation of conventional and modern degradation techniques to remove them was carried out including toxicological assessments and identification of by-products.

Within this thesis, three monitoring assessments were carried out, including both targeted and non-targeted screening approaches. Two different analytical methods - using High Performance Liquid Chromatography coupled to a Triple Quadrupole Mass Spectrometer (HPLC-MS/MS) - for evaluating the pollution rates, at trace level concentrations, were developed, following the principles of Green Analytical Chemistry, which required them to be fast, cost-effective and more green. Both methods were validated according to the requirements of ISO 17025 and those established by Accredia, the Italian National System for laboratories' accreditation.

More specifically, during the first year of this PhD project an innovative method for the determination of seventeen Perfluoroalkyl Substances (PFASs) in different water matrices (including surface, ground and drinking water) at really low quantification levels was developed. The target compounds included a mix of seventeen linear perfluoroalkyl substances, with a chain length ranging from four to eighteen atoms of carbon, in order to investigate the variability among their physicochemical properties. The key characteristic of this method is the achievement of the compounds' detection in a maximum analysis time of 10 minutes and in really low concentration levels without using any preconcentration step. In the literature, only three methods using a direct injection analysis [51,117,118] are reported. However, all of them target at less than 17

compounds, and have a longer analysis time than the one reported here. Other reported methods achieved as well low quantitation levels (in the ppt range), but for a smaller number of target compounds and after the use of an extraction/pre-treatment step [46] [119-123]. Furthermore, until now very few standard methods (Table 4) – EU aims to develop standard methods for PFAS until 2024 - exist for the determination of PFASs in water samples, with only one, the American Society for Testing and Material (ASTM) method, not including any pre-treatment step. However, this method is not referring to determination of PFASs in drinking water samples. Similarly, even if both the ISO 25101:2009(E) [124] and the United States Environmental Protection Agency (EPA) 537 [125] methods, concern drinking water matrices, they use a solid phase extraction (SPE) step and focus on a smaller number of compounds than the one reported in the developed method.

The developed method was used in the first monitoring assessment study reported in this thesis, which aimed at estimating the PFASs occurrence levels in the Metropolitan Area of Turin. A correlation study between the obtained results and the potential pollution sources in the area (such as WWTPs, industries and civilian airports) was carried out using spatial multivariate statistical analysis tools, in order to develop a statistical framework that investigates if the presence of PFASs in water bodies is associated with the number of pollution sources within a watershed. In this way, a geographical model that supports “smart” water quality monitoring programs, was developed in order to not only take into account the number of inhabitants or the volume of supplied water when planning quality monitoring programs, as done till now. The results of this study (**P1**) are summarized in **Section III, Chapter 3**.

Table 4. A comparison of standard methods for PFASs analysis [46].

| Standard Method | EPA 537 | ISO 25101:2009(E) | ASTM D7979-16 | ASTM D7868-14 |
|------------------------------|---|--|---|---|
| Sample volume | 250 mL | 500 mL | 5 mL | 2 g |
| Sample matrix | Drinking water | Drinking water, groundwater, surface water and seawater | Water; wastewater sludge, influent and effluent | Solid and biosolid |
| Analytes | PFAS and FOSAAs 14 PFAS | PFOS and PFOA | PFAS, FOSAAs, FTSS, n:2 FTUCAs and FTCAs | PFAS, FOSAAs, FTSS, n:2 FTUCAs and FTCAs |
| Preservation | Trizma for buffering and removal of free chlorine | Sodium thiosulfate pentahydrate for removal of free chlorine | None | None |
| Holding time | Before extraction: 14 days refrigerated at ≤ 6 °C Postextraction: 28 days at room temperature | 14 days at 4 ± 2 °C | 28 days at $0-6$ °C | 28 days at $0-6$ °C |
| Extraction Method | SPE-WAX (SPE Weak anion exchange) | SPE | Direct injection | Solvent extraction followed by filtration using polypropylene filters |
| Analytical instrument | LC-MS/MS (liquid chromatography tandem with mass spectrometry) | LC-MS/MS and LC/MS | LC-MS/MS | LC-MS/MS |
| Reporting limits | 2.9–14 ng/L | 2–10,000 ng/L | 10–400 ng/L | 25–1000 ng/L |

The second developed method focused on the quantification of sixteen different pharmaceutical compounds and hormones (PhACs) at trace levels in water. It was developed following the principles of Green Analytical Chemistry and validated according to the criteria of ISO/IEC 17025 [126] as well. However, taking into account the variety among the compounds and their properties, in this method a preconcentration step was necessary for achieving low Quantification Levels for all of them. The second monitoring assessment reported in this thesis used this method with the aim to evaluate the concentrations of the sixteen target PhACs in raw drinking water sources (surface and groundwater) of the Metropolitan Area of Turin. However, in order to avoid a blind monitoring, like the first assessment campaign, the geographical tool developed in Chapter 3 was used for choosing the sampling points in higher risk of pollution, according to their geographical position close to potential pollution sources (WWTPS, hospitals and care houses). Finally, a risk assessment was carried in order to evaluate the potential adverse effects that PhACs' occurrence in drinking water can have on human health after a long-term exposure to low doses. Human health risks were evaluated considering average detected concentrations for individual compounds and their mixtures, and provisional guideline values for those that a drinking water regulatory value didn't exist. Since risk assessments are required for establishing priority substances for monitoring and if necessary managing their removal – as reported in the just issued Drinking Water Directive 2020/2184/UE [36]- the overall aim of this study was to fill some knowledge gaps existing in literature for PhACs' risk assessments due to limited datasets and synergistic effects of contaminants mixtures [127,128]. The results of this study (**P4**) are collected in **Section III, Chapter 4**.

However, one fundamental obstacle in evaluating water quality with current monitoring approaches is the fact that they focus on a very small number of chemicals. In order to more comprehensively assess the presence of contaminants in the aquatic bodies and prioritize pollutants within regulatory applications, target, suspect and non-target screening approaches should be combined. Hence, in this thesis a non-target monitoring assessment based on High Resolution Mass Spectrometry (HRMS) was included. Surface water samples from two different countries – Greece and Italy – were analyzed after a conventional SPE step by Liquid Chromatography coupled to a hybrid Quadrupole-Time-of-flight Mass spectrometer (LC-QTOF-MS) using the Sequential Window Acquisition of all Theoretical Mass Spectra mode (SWATH-MS). A suspect list of 100 compounds, with available standards in the laboratory, was developed and a stock solution containing them, was prepared and analyzed as well. After searching the chromatograms for exact

masses of suspects -as an additional help for identifying compounds in the samples- a non-target screening was carried out. However, due to the complexity of environmental samples, thousands of HRMS features were generated by their nontarget screening analyses (in this thesis detected fragments/ions will be referred to as features prior to their identification). For this reason, data reduction efforts for collecting reliable information were pursued, followed by the application of Multivariate chemometric methods - such as Principal Component Analysis (PCA) and Partial Least Squared Discriminant Analysis (PLS-DA) - in order to process the big obtained datasets, reveal pollution patterns and prioritize the features responsible for the discrimination among the samples. The use of a database with more than 4000 entries contributed to high confidence identification of the compounds, which was based on mass accuracy, retention time, isotopic ratio pattern, and MS/MS fragmentation pattern searching. For those that a correspondence to the database was not found, an empirical formula calculation was done by the instrument's software, followed by an *In silico* fragmentation using an online library to recognize the MS/MS fragment ions and successfully identify unknown compounds. The results of this study are reported in **Section III, Chapter 5.**

The importance of monitoring assessments except the fact that they provide valuable information about water bodies' quality status, they can be a helpful tool for treatment decisions. In general, conventional treatment methods used in WWTPs and DWTPs have been found to face difficulties in efficiently removing CECs from water, especially when they occur in trace level concentrations [66,97,129,130]. For this reason, the need of finding new more efficient degradation processes has emerged with great attention being raised towards Advanced Oxidation Processes (AOPs), a green alternative. AOPs rely on the generation of highly reactive species, able to oxidize the emerging pollutants. These techniques have shown great potential in removing efficiently CECs from water bodies, especially when they are combined with the conventional treatment processes. For this reason, after taking into account the results obtained from the monitoring assessments, as well as the general trends of aquatic contamination, the evaluation of the pollutants' removal by conventional and advanced water treatment technologies was carried out, during the second half and third year of this PhD over secondment periods in other Universities. The obtained results are summarized in Section IV. More specifically, during a secondment period at the University of Aalborg, an attempt of finding environmental sustainable treatments was done by evaluating the degradation efficiency of UV photolysis.

UV photolysis has been reported as a degradation technique of low environmental impact and minimized environmental footprint since it doesn't have a lot of prerequisites and doesn't generate waste [131]. The contaminants' degradation after exposure to UV irradiation can be a result of direct photochemical transformation by photons' absorption, or of interactions with generated reactive oxygen species. In the study (P2) reported in **Section IV, Chapter 6**, the effects of irradiation on glyphosate - the most detected pesticide in the aquatic environment - were investigated. Samples of the compound were prepared in both MilliQ water and different real water matrices (including drinking and groundwater) from the Municipality of Aalborg, Denmark and exposed to UV-A (365nm), UV-B (302nm) and UV-C (254nm) irradiation under different UV doses. A test battery including aquatic organisms from different trophic levels was used in order to evaluate glyphosate's biological effects and bioactive transformation products before and after treatment. Studies measuring the effectiveness of UV irradiation alone as a remediation treatment for pesticides and more specifically glyphosate, should include an assessment of its potential to alleviate the parent compound's toxicity to non-target organisms as well as the one from the transformation products. For this reason, in this study we combined LC-MS analysis in order to follow glyphosate's degradation and identification of byproducts with bioassays to assess the compound's removal and toxicity to non-target aquatic test organisms before and after treatment was done. Samples taken before and after exposure to different wavelengths of UV irradiation, were analyzed with LC-Orbitrap-MS, and based on a suspect screening, identification of by-products was done.

Finally, in **Section IV, Chapter 7** is presented the High Voltage Pulsed Electric Field process, a promising technology for degrading persistent emerging contaminants that are recurrent in water bodies and can pose a threat to both human health and aquatic environment. One major category of such compounds includes PFASs, which host one of the strongest chemical bonds in their carbon chain (C-F) that makes them highly persistent against conventional treatment methods. Moreover, the fact that these contaminants coexist in mixtures and the variety of physicochemical properties among them, harshens the opportunity to find one efficient degradation technique. Hence, in this study, the performance of non-thermal plasma technique was evaluated. The reactor used for the treatment is a patent of IRIS S.r.l., and the technique basically applies on one or several very high voltage (HV) pulses of very short duration to a reactor containing the contaminated water samples, generating in this way pressure waves, UV light and formation of chemically active species such as $\bullet\text{OH}$, $\bullet\text{H}$, $\bullet\text{O}$, $\bullet\text{O}_2^-$, $\bullet\text{HO}_2$, $\bullet\text{H}_2\text{O}_2$,

•O₃ [132], that can break the organic molecules. Based on the results from the study reported in Chapter 3, the most abundant compounds in the area were prioritized for treatment. The selected compounds had different chain lengths, included both perfluoroalkane sulfonates and perfluoroalkyl carboxylates, and all of them were linear. Samples of individual compounds as well as their mix were prepared in MilliQ water and real water matrix, and analyzed with LC-QTOF-MS in both HRMS MRM and SWATH acquisition modes. The MRM mode contained the parameters of the method reported in Chapter 3 and was used for the quantification of the compounds before and after treatment, in order to follow the degradation profiles (**P3**). The SWATH mode, based on a full scan acquisition, was used for identifying by-products with both suspect and non-target screening (**P5**).

Section III

Freshwater Quality Monitoring and Assessment

Chapter 3 Per and polyfluoroalkyl Substances (PFASs)

Firstly, an advanced quantitative analytical method for detecting PFASs in different water type samples was developed and validated. The overall aim was to obtain a fast, cost-effective and green method – following the Green Analytical Chemistry principles– in order to be applied in a large scale monitoring assessment of PFASs levels in the Metropolitan Area of Turin. Finally, an estimation study of the potential pollution sources was carried out by correlating the PFASs occurrence results with industrial sites, civilian airports and WWTPs that exist in the area, using spatial and multivariate statistical analysis tools.

1. Materials and Methods

1.1 Reagents and Chemicals

A mix (PFAC-MXB) of seventeen PFASs was examined in this study containing compounds with various carbon chain lengths (between four to eighteen atoms of carbon): thirteen linear perfluoroalkylcarboxylic acids and four perfluoroalkylsulfonates (Table 5). The standard mix solution PFAC-MXB was purchased from Wellington Laboratories (Guelph, ON, Canada) with chemical purities of >98% and a concentration of 2000 ng/mL in Methanol/Water <1% for every individual perfluoroalkylcarboxylic acid and perfluoroalkylsulfonate. Another mix (MPFAC-MXA) containing seven mass-labelled (^{13}C) perfluoroalkylcarboxylic acids and two mass-labelled (^{18}O and ^{13}C) perfluoroalkylsulfonates was used as internal standards (Table 5). The mix solution MPFAC-MXA was purchased from Wellington Laboratories (Guelph, Ontario, Canada) with chemical purities >98% and a concentration of 2000 ng/mL in Methanol/Water <1% for every individual mass-labelled perfluoroalkylcarboxylic acid and mass-labelled perfluoroalkylsulfonate and with isotopic purities of 99% per ^{13}C and >94% per ^{18}O . UHPLC-grade Methanol was purchased from Sigma-Aldrich, Co (St. Louis, MO, USA), MilliQ was obtained from MilliPore (MA, USA) and Ammonium acetate for LC-MS LiChropur[®] was purchased from Merck KGaA (Darmstadt, Germany).

Table 5. Target compounds and their related internal standards.

| Target Compounds (PFAC-MXB) | | Internal Standard Compounds (MPFAC-MXA) | |
|---------------------------------------|--------------|---|--------------|
| Full Name | Abbreviation | Full Name | Abbreviation |
| Perfluoro-n-butanoic acid | PFBA | Perfluoro-n-[¹³ C ₄]butanoic acid | MPFBA |
| Perfluoro-n-pentanoic acid | PFPeA | Perfluoro-n-[1,2- ¹³ C ₂]hexanoic acid | MPFHxA |
| Perfluoro-n-hexanoic acid | PFHxA | Perfluoro-n-[1,2- ¹³ C ₂]hexanoic acid | MPFHxA |
| Perfluoro-n-heptanoic acid | PFHpA | Perfluoro-n-[1,2- ¹³ C ₂]hexanoic acid | MPFHxA |
| Perfluoro-n-octanoic acid | PFOA | Perfluoro-n-[1,2,3,4- ¹³ C ₄]octanoic acid | MPFOA |
| Perfluoro-n-nonanoic acid | PFNA | Perfluoro-n-[1,2,3,4,5- ¹³ C ₅]nonanoic acid | MPFNA |
| Perfluoro-n-decanoic acid | PFDA | Perfluoro-n-[1,2- ¹³ C ₂]decanoic acid | MPFDA |
| Perfluoro-n-undecanoic acid | PFuDA | Perfluoro-n-[1,2- ¹³ C ₂]undecanoic acid | MPFuDA |
| Perfluoro-n-dodecanoic acid | PFDoA | Perfluoro-n-[1,2- ¹³ C ₂]dodecanoic acid | MPFDoA |
| Perfluoro-n-tridecanoic acid | PFTrDA | Perfluoro-n-[1,2- ¹³ C ₂]dodecanoic acid | MPFDoA |
| Perfluoro-n-tetradecanoic acid | PFTeDA | Perfluoro-n-[1,2- ¹³ C ₂]dodecanoic acid | MPFDoA |
| Perfluoro-n-dexadecanoic acid | PFHxDA | Perfluoro-n-[1,2- ¹³ C ₂]decanoic acid | MPFDA |
| Perfluoro-n-octadecanoic acid | PFODA | Perfluoro-n-[1,2- ¹³ C ₂]decanoic acid | MPFDA |
| Potassium perfluoro-1-butanefulfonate | L-PFBS | Sodium perfluoro-1-hexane [¹⁸ O ₂] sulfonate | MPFHxS |
| Sodium perfluoro-1-hexanesulfonate | L-PFHxS | Sodium perfluoro-1-hexane [¹⁸ O ₂]sulfonate | MPFHxS |
| Sodium perfluoro-1-octanesulfonate | L-PFOS | Sodium perfluoro-1-[1,2,3,4- ¹³ C ₄] octanesulfonate | MPFOS |
| Sodium-1-decanesulfonate | L-PFDS | Sodium perfluoro-1-hexane [¹⁸ O ₂]sulfonate | MPFHxS |

1.2 Study area and sampling

As already mentioned SMAT is the company in charge of managing and supplying water to 293 municipalities in Piedmont, Italy (Figure 5, Table A1) and a population of about 2,3 million inhabitants. The sampling campaign was organized by SMAT and carried out between March 2018 and October 2018. In total 930 samples were collected from the whole territory, according to the specifications and requirements of ISO 5667 [133], including 5% of surface, 19% of groundwater and 76% of drinking water. As surface water were considered samples taken from rivers, streams, and at the intake of SMAT DWTP (including river and lagoon water), as groundwater those taken from pumps at each wellhead, and as drinking samples those collected at the end of the DWTP treatment line, fountains, and tanks. The samples were collected in polypropylene bottles WM (wide mouth) with caps (volume 125 mL), purchased from SciLabware Limited (Stoke-on-Trent, Staffordshire, ST4 4RJ, United Kingdom), stored at 4°C prior to their analysis, and analyzed within 15 days from their collection, in order to prevent biodegradation of the matrix and reassure the recovery of the analytes.

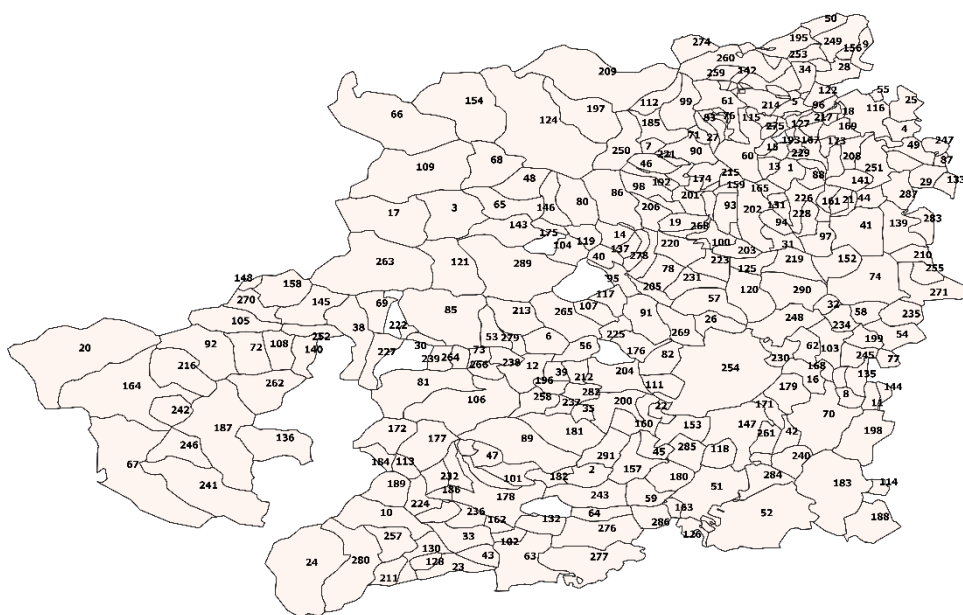


Figure 5: Map of the municipalities in the study area (Metropolitan Area of Turin, Piedmont, Italy), more information available on Table A1.

1.3 Sample preparation

The samples were injected directly into the analytical system without any pretreatment step. A filtration step was not necessary as the samples—mostly drinking water—were not contaminated with soils or suspended organic matter. Two working standard solutions—in 50% MeOH/50% H₂O for the first and in 100% H₂O for the second—were prepared with a dilution from each of the two stock solutions, and used for the calibration. The purchased solutions were stored at 4°C, while the other four were stored at room temperature. A volume of 700 µL of each sample was transferred into 0,7 mL Polypropylene Short Thread Micro-Vials (purchased from CPS Analytica for Chemistry, Milan, Italy), and 1 µL of the Internal Standard mix (50 ng/L) was added.

1.4 Instrumental Analysis

Analyses were carried out using the SCIEX QTRAP[®] 6500 system (SCIEX, Framingham, MA, USA) with a Thermo Scientific Dionex UltiMate 3000 UHPLC system and a RS-3000 autosampler (Dionex Softron GmbH, Germering, Germany). The UHPLC instrument was equipped with a Luna[®] C18 (2) HPLC Column (5 µm particle size, 30 mm × 2.0 mm; Phenomenex Inc., Torrance, CA, USA) installed between the eluent mixer and the autosampler, in order to delay the potential contamination originating from the UHPLC system. The chromatographic separation was achieved using a Luna[®] Omega PS C18 HPLC Column (1,6 µm particle size, 50 mm × 2,1 mm; Phenomenex Inc., Torrance, CA, USA)—heated to 40 °C—by injecting a 50 µL sample volume at a mobile phase consisted of a mixture of 20 mM Ammonium Acetate in Water (A) and Methanol (B), lasting a total time of 12 minutes. The gradient profile, with a flow rate of 0,550 mL/min, started with 98% A and 2% B, increasing to 100% B in 6 minutes, and, after keeping this ratio for 1,5 minutes, reversed into the initial conditions (Table A2). The parameters of the mass spectrometer are summarized in Tables A3, A4.

1.5 Method Validation

In order to reassure the validity of the results, a validation process was necessary and carried out following the ISO/IEC 17025 [126] guidelines and those set by Accredia, the Italian National Accreditation System. Six-point calibration curves of final concentrations 5, 10, 25, 50, 90 and 120 ng/L were built for each target compound, and fifteen replicates of each point were analyzed in order to estimate the uncertainty, trueness, linearity, recovery and limits of Detection (LOD) and Quantification (LOQ). In order to reassure the quality of the method and the best performance of the instrument during the analyses,

blank and control (QC) samples were analyzed after every ten samples. QC samples were prepared by diluting the standard solution in MilliQ water with a final concentration of 50 ng/L and adding 50 ng/L of the internal standard mix. The quantitation was performed using the software MultiQuant™ 3.0.3 software (SCIEX, Framingham, MA, USA).

1.6 Spatial and Statistical Analysis

Spatial analysis was performed in order to develop a framework that investigates the correlation between contamination levels and the number of potential pollution sources within a watershed. For this reason, information about 176 industrial sites (Figure A1) and 800 WWTPs present in the study area were taken from Arpa Piemonte [134,135], and in particular, the geographical data used (coordinates in WGS 84 system and maps of the area) were obtained from the free Diva-Gis platform [136]. The QGIS 3.4 software was used for estimating the correlation between the sampling points with detected concentrations above the LOQs and the industrial sites and WWTPs that exist in the area, within a radius of 5km from the sampling points, and can potentially be characterized as potential pollution sources. However, the lack of available information concerning emerging pollutants employed by the industrial sites, led us to choose them according to their sector activities and manufacturing products known to potentially employ PFASs. QGIS 3.4 was also used for building the thematic maps, which provide a distinct visualization of the spatial distribution of the detected concentrations in the area. The GeoDa 1.12 software was used in order to build the Ordinary Least Squares (OLS) and the Spatial Regression models, following the equation (8):

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \lambda \quad (8)$$

where Y is the dependent variable, X_1 , X_2 are the explanatory or independent variables, β_0 , β_1 , β_2 are the coefficients and λ is the random spatial error term. Spatial autocorrelation between the area units was evaluated utilizing the Moran's I statistic, while the Akaike info criterion was used to check the strongest model for the correlation prediction between the potential pollution sources and the contaminated areas [137].

2. Results

2.1. Method development and optimization

During the development of an analytical method aiming to detect trace level contamination, like in nanograms per liter (or parts per trillion; ppt), cross contamination effects can have a large impact on the accuracy and validity of the analytical results. More specifically, since PFASs are compounds with a vast variety of applications,

background contamination can originate even from the laboratory equipment [117]. In order to minimize these effects, online SPE-LC-MS/MS methods for PFASs detection have been proposed and reported in the literature [51]. However, an elimination of the pretreatment step may minimize better these risks, alongside with meticulous and methodical manipulation of the samples. In this study, teflon and glass materials were avoided, as well as a filtration step - even if it was not necessary for drinking water samples - in order to minimize any contamination of the samples. Furthermore, a smaller HPLC column was added between the pump and the injector in order to delay possible contamination originating from the solvents [51,118-120].

However, direct analysis of spiked water was proven unsuccessful, as the results didn't show sufficient recovery levels for the compounds with longer chains. For this reason, taking into account the diversity of chemical properties among the target compounds as well, we chose to add Methanol in the samples following the reports of the EPA 537 method [125]. Optimization of the solvents' percentage ratio in both working standard mixes was done, and satisfying results were achieved after preparing them in 50% MeOH/50% H₂O. Moreover, in order to ensure the best detection results, a mix of isotopically labelled internal standards was added at a concentration of 50 ng/L. For quantifying the molecules, the ratio between the peak areas of the target compounds and that of their related internal standard was used.

Concerning the chromatographic conditions, the best compounds' separation was achieved after using a shorter column as an isolator, placed before the main one, in order to separate the target compounds occurring in the analytical samples, from those that were potentially present in the solvents. Furthermore, the parameters of the gradient elution of the mobile phase were optimized as well before achieving the best for analyte's separation and shape of peaks. Firstly, an elution starting from 40% Methanol and 60% Ammonium Acetate in water (5 mM) increasing to 100% Methanol and returning to the initial conditions within 6 minutes was tried. These conditions provided a satisfactory separation of the longer chain compounds, but some of the shorter chain ones were co-eluting at the beginning of the chromatogram. After different efforts, the best conditions concluded to those finally used (Figure 6, Tables A2, A3).

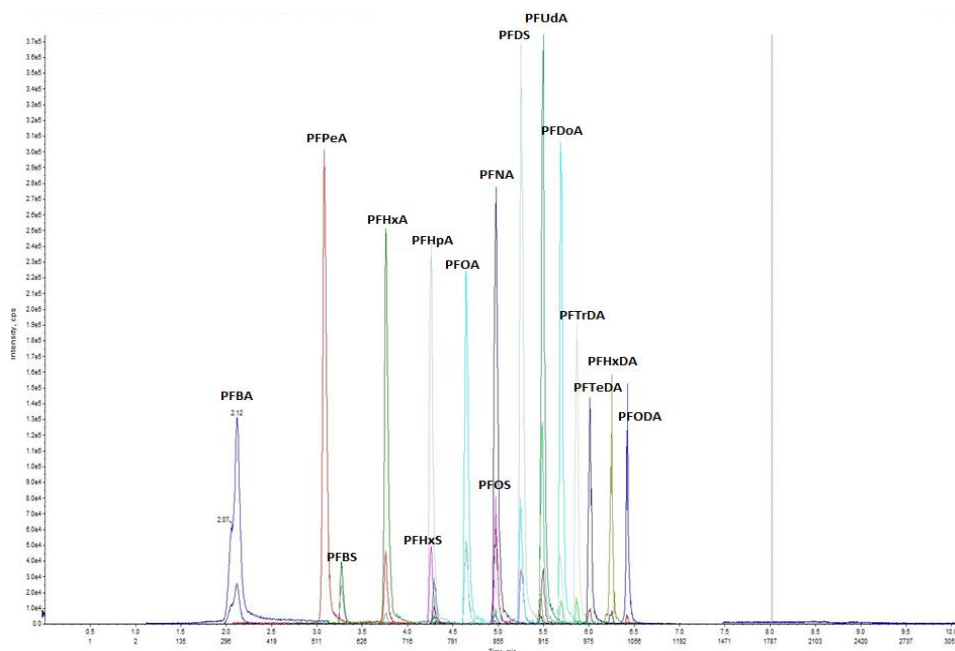


Figure 6: A typical chromatogram.

2.2 Validation and Quality Assurance

Concerning the validation parameters, linearity is based on the linear regression analysis of the obtained quantitative data. The regression coefficients (R^2) of the six-point calibration curves were calculated by the ratio between the peak area of the target compound, the peak area of the relative Internal Standard (IS) and the different concentrations of the calibration curve. A $1/x$ concentration-weighting factor was used in order to give more emphasis to the lower concentrations and to ensure the best assay performance. Good coefficient results between a range of 0,980 and 0,999 (Table 6) were obtained for all the target PFASs.

Furthermore, the measurement of the systematic and random errors is a crucial step during the validation of a method. According to ISO/IEC 17025 [126], the required values for uncertainty accepted values should be $RSD \leq 20\%$ and for a sufficient recovery within the range of 70%–120%. For this method, the variability and reproducibility of the results were calculated for every point of the calibration curves, offering satisfactory results within the acceptance ranges (Table 6). The recovery of the compounds was also tested in 4 different real water matrices, spiked with the mix of the target compounds at the same concentrations (50 ng/L) offering satisfactory results (Table 7). Similarly, the

accuracy of the method, based on the trueness of the results, was calculated for every analyte and found within the range of $\leq 30\%$ (Table 6).

Table 6. Validation results.

| Compounds | Concentration ng/L | Trueness % | Recovery % | Uncertainty % | Linearity | LOQ |
|-----------|-----------------------|---------------|---------------|------------------|-----------|-----|
| PFBA | 50 | -8,6 | 91,38 | 3,13 | 0,997 | 5 |
| PFBS | 50 | -4,2 | 95,78 | 2,89 | 0,997 | 5 |
| PFPeA | 50 | 1,3 | 101,28 | 1,67 | 0,999 | 5 |
| PFHxA | 50 | -5,2 | 94,84 | 2,82 | 0,998 | 5 |
| PFHxS | 50 | -3,9 | 96,12 | 6,07 | 0,997 | 5 |
| PFHpA | 50 | -4,7 | 95,28 | 3,63 | 0,999 | 5 |
| PFOA | 50 | -3,7 | 96,28 | 4,36 | 0,998 | 5 |
| PFOS | 50 | -8,0 | 92,04 | 1,96 | 0,992 | 5 |
| PFNA | 50 | -9,6 | 90,38 | 4,82 | 0,999 | 5 |
| PFDA | 50 | -14,4 | 85,58 | 7,59 | 0,995 | 5 |
| PFUdA | 50 | -5,3 | 94,72 | 9,64 | 0,998 | 5 |
| PFDoA | 50 | -7,2 | 92,78 | 8,72 | 0,999 | 5 |
| PFTrDA | 50 | -26,8 | 73,22 | 17,53 | 0,989 | 5 |
| PFTeDA | 50 | -30,0 | 69,98 | 13,76 | 0,987 | 5 |
| PFHxDA | 50 | -15,1 | 84,88 | 15,97 | 0,995 | 5 |
| PFODA | 50 | -13,7 | 86,34 | 5,48 | 0,980 | 5 |
| PFDS | 50 | -51,3 | 48,68 | 15,78 | 0,997 | 5 |

Finally, in order to calculate the Limits of Detection and Quantification for each analyte the requirements of the International Conference on Harmonisation (ICH) Method [139] were followed. More specifically, the standard deviation of the y -intercepts and the slope of the calibration curves were obtained from the 15 replicate analyses of the calibration curves. The LOD was calculated following the equation (9) and LOQ following the equation (10)

$$\text{LOD} = 3,3 \sigma / S \quad (9)$$

$$\text{LOQ} = 10 \sigma / S \quad (10)$$

where σ is the standard deviation of y -residuals and S is the slope of the calibration curve. The results for the LOQs of the target PFASs varied between 3 (for shorter chain compounds) to 8 ng/L (for longer chain compounds). However, for practical reasons and

data processing uniformity as a Limit of Quantification for every compound (as stated also in the ASTM D7979-17 method [140]) was considered the lowest point of the calibration curve satisfying the trueness and uncertainty criteria of less than 30% (considering the 15 replicates). In this way, the LOQ for every compound resulted at 5 ng/L (Table 6). However, the validation results for PFDS were not satisfactory (Table 6). Hence, this compound was removed from our method.

Table 7. Recovery results after spiking real water samples.

| | | Real Sample 1 | Real Sample 2 | Real Sample 3 | Real Sample 4 |
|-----------|--------------------|---------------|---------------|---------------|---------------|
| Compounds | Concentration ng/L | Recovery % | Recovery % | Recovery % | Recovery % |
| PFBA | 50 | 107,52 | 106,06 | 90,84 | 107,05 |
| PFBS | 50 | 106,92 | 107,22 | 90,76 | 103,90 |
| PFPeA | 50 | 108,22 | 105,85 | 90,53 | 109,99 |
| PFHxA | 50 | 104,96 | 106,89 | 91,69 | 107,11 |
| PFHxS | 50 | 104,96 | 106,73 | 90,88 | 105,98 |
| PFHpA | 50 | 102,19 | 102,93 | 92,57 | 105,56 |
| PFOA | 50 | 103,63 | 114,74 | 92,09 | 107,14 |
| PFOS | 50 | 95,92 | 97,31 | 86,33 | 96,11 |
| PFNA | 50 | 106,18 | 102,78 | 97,41 | 104,91 |
| PFDA | 50 | 95,29 | 90,22 | 81,93 | 90,39 |
| PFUdA | 50 | 93,49 | 83,01 | 81,57 | 80,78 |
| PFDoA | 50 | 93,19 | 100,73 | 96,98 | 97,45 |
| PFTTrDA | 50 | 97,89 | 87,62 | 82,73 | 97,13 |
| PFTeDA | 50 | 101,03 | 84,89 | 94,62 | 88,16 |
| PFHxDA | 50 | 100,61 | 101,19 | 84,50 | 101,98 |
| PFODA | 50 | 86,63 | 90,16 | 87,91 | 93,71 |
| PFDS | 50 | 72,26 | 75,31 | 52,28 | 74,51 |

2.4 Monitoring Assessment results

The developed method was used in an estimation study of the PFASs' pollution levels in the Metropolitan Area of Turin. In total, 930 samples were collected from all the steps of a water supply system (from the catchment till the tap), through a sampling campaign organized by SMAT, during March and October 2018. Between the samples, 5% included surface water, 19% groundwater and 76% drinking water.

All the detected concentrations -both for individual compounds as well as their mix- were significantly lower than the drinking water performance values set by the Italian Ministry of Health (30 ng/L for PFOS and 500 ng/L as sum of PFAS), and the parametric limit values reported in the revised Drinking Water European Directive 2020/2184 [36] (100 ng/L as single and 500 ng/L as sum of PFAS). In this study, the highest detected concentration for the sum of the target compounds was 57 ng/L. Only four out of the sixteen compounds were detected in the area in concentrations above their Limits of Quantification (5 ng/L) and they were the perfluoro-n-butanoic acid (PFBA), perfluoro-1-hexane sulfonate (PFHxS), perfluoro-n-octanoic acid (PFOA) and sodium perfluoro-1-octanesulfonate (PFOS). For PFBA the highest detected concentration was 19 ng/L (Figure 7a), while for PFHxS and PFOA, they were 15 ng/L and 9 ng/L, respectively (Figure 7b,c). Concerning PFOS, the highest detected concentration was 23 ng/L (Figure 7d), which was the highest among the four detected compounds. The highest concentrations for these four compounds were detected in raw sources. These results were in contrast with those reported in other studies, where as a general finding the detected concentrations of carboxylates in the aquatic environment are higher than those of sulfonates [137].

In general, PFBA, PFHxS, PFOA and PFOS were present alone or in mixtures of two or three compounds in the analyzed samples. However, only in a groundwater sample of one municipality, which was the one that hosts the civilian airport on the area, were present all together. Even if their sum concentration was low, this was an expected result since it is known that PFASs constitute some of the main components of the aqueous film-forming foams (AFFF) used in that areas [46]. Detected PFASs concentrations were observed in all the different types of water sources included in this study, with those concerning treated water being significantly lower, but still above the individual LOQ levels. These results alongside with the fact that the highest concentrations were detected close to industrial sites and WWTPs, confirm the studies in literature claiming that conventional treatment methods (both in DWTPs and WWTPs) are not sufficient in efficiently removing PFASs from water. Understanding the correlation between pollution sources and detected concentrations in drinking water, and whether increased concentrations are associated with their number within a watershed is important for water utilities in order to identify the exposure risks. For this reason, in this study multivariate spatial regression models [141] were developed in order to identify correlations between contaminated sampling points and the selected potential pollution sources.

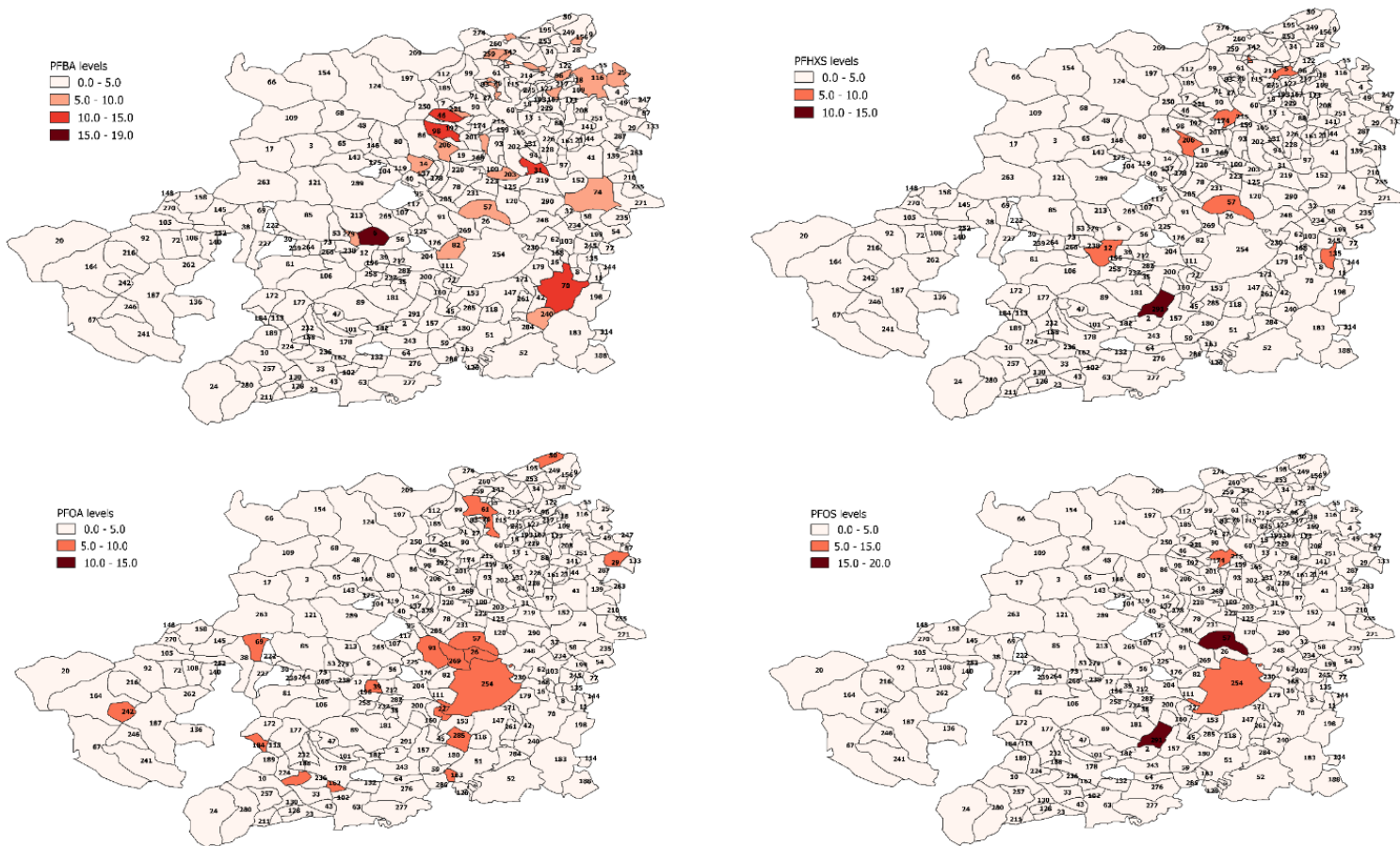


Figure 7: Concentration levels(ng/L) among the different municipalities of the study area for the four detected compounds: (a) PFBA; (b) PFHxS; (c) PFOA; (d) PFOS.

2.5 Spatial and Statistical Analysis

In this study, spatial analysis was performed in order to study the geographic correlation between the “positive” sampling points and the potential contamination sources that are surrounding them. The main principle of spatial statistical analysis is that space is an influence on the observations of a study, meaning that it is very likely for values obtained from nearby points to be more similar to each other than with values obtained from more distant points. Regression analysis is a tool that allows to examine and model these location-oriented relationships, and helps to explain spatial patterns. In this study, the results obtained from the screening assessment were used in order to build three different regression models, the OLS, Spatial Lag and Spatial Error Regression. The model that best explained the relationship between the concentration levels of the compounds (dependent variables) and the number of WWTPs and industrial activities in their surrounding area (independent variables) was chosen, based on the obtained Akaike criterion and R^2 values (Table 8). In order to build the regression models spatial weights -as measures of the influence- were taken into account.

Table 8. Summary of the different regression models.

| Coefficients | | OLS | Spatial Error | Spatial Lag |
|--------------|-----------------------|---------|---------------|-------------|
| PFBA | Industrial Sites | 1,1012 | 1,1371 | 1,1233 |
| | WWTPs | 0,2790 | 0,2795 | 0,2792 |
| | R^2 | 0,660 | 0,662 | 0,662 |
| | Akaike info criterion | 1020,04 | 1021,45 | 1019,78 |
| PFHxS | Industrial Sites | 0,1667 | 0,1688 | 0,1678 |
| | WWTPs | -0,0110 | -0,0118 | -0,0113 |
| | R^2 | 0,270 | 0,303 | 0,296 |
| | Akaike info criterion | 848,65 | 850,48 | 848,58 |
| PFOS | Industrial Sites | 0,2719 | 0,2859 | 0,2785 |
| | WWTPs | -0,0119 | -0,0123 | -0,0121 |
| | R^2 | 0,078 | 0,081 | 0,076 |
| | Akaike info criterion | 909,98 | 911,79 | 909,97 |
| PFOA | Industrial Sites | 0,6166 | 0,576574 | 0,59239 |
| | WWTPs | -0,0311 | -0,0251 | -0,0261 |
| | R^2 | 0,242 | 0,225 | 0,233 |
| | Akaike info criterion | 968,21 | 963,78 | 965,32 |

Based on the results obtained from the different regression models, the Spatial Error model was the one that explained better the spatial correlation between the compounds PFBA, PFHxS and PFOS and the potential pollution sources in the area. Concerning PFOA, the spatial regression model that best described the relationship between the contaminated areas and the industrial activities and WWTPs was the OLS. The results of the chosen models are summarized in Table 9. These results explained 8%–66% of the variance in the water samples for the four PFAS compounds that were detected. Increasing PFBA concentrations were positively associated with the number of industrial sites and the WWTPs present in the surrounding area of the sampling point, meaning that each potential pollution source is associated with a 66% increase in PFBA levels. This relationship resulted as statistically significant ($p < 0,001$), and as the strongest statistical association across the compounds and contamination sources.

Table 9. The chosen spatial regression models for PFASs concentrations in drinking water.

| Compounds | Industrial Sites | WWTPs | λ^* | R^2 |
|---------------------------------|------------------|------------------|---------------|-------|
| PFBA coefficient p-value | 1,1371 0,001 | 0,2795 0,001 | 0,092 0,05 | 0,66 |
| PFHxS coefficient p-value | 0,1688 0,002 | -0,0118 0,523 | 0,033 0,07 | 0,30 |
| PFOS coefficient p-value | 0,2859 0,001 | -0,0123 0,527 | 0,082 0,07 | 0,08 |
| PFOA coefficient p-value | 0,6166 0,001 | -0,0311 0,151 | 0,159 0,07 | 0,24 |

* Spatial error term coefficient showing spatial influence.

The other three detected compounds showed positive correlations with the number of the industrial sites, with statistically significant relationships ($p < 0,05$), indicating that each additional industrial site is associated with a 30% increase in PFHxS, 8% increase in PFOS and 24% in PFOA levels. However, the regression models showed negative association between the number of WWTPs and the increasing levels of PFHxS, PFOA and PFOS, with relationships lacking statistical significance ($p > 0,05$) (Table 9). This

indicates that PFAS releases from WWTPs are important but less significant than those from industries, following the findings obtained from Hu et al. [137]. However, the low number of sampling points with concentrations above the LOQ for these three compounds (for PFHxS, only 7 out of the 930 samples were positive, whereas for PFOS, only 6 out of the 930) can explain these results. The Moran's I statistic was used for evaluating the spatial autocorrelation between the area units. Basically this test is used for examining if the attribute values of features (compounds) cluster or not, taking into account locations of other features. A result of -1 would mean a checkered pattern, a result of 0 would mean a random pattern and a result of 1 would mean a clustered pattern. In this study the results of Moran's I statistic for the four regression models built were between 0,131 (for PFOS) and 0,541 (for PFBA), showing a random pattern of spatial autocorrelation among the area units.

The spatial analysis performed in this study was challenging due to the low detection results and the lack of available information, as reported by the λ coefficients that represent the spatial error. Geospatial data for many potentially important PFAS point sources were not present, as well as information about the companies' production processes. Moreover, no data about the employment of this class of substances (as PFAS are not regulated yet) or the airborne emissions were available in order to evaluate the importance of the atmospheric releases. Information about where and if the intake of the water supply was upstream from the point source of pollution was not accessible as well.

3. Conclusions

In this chapter, a new green, fast and validated method with high sensitivity in detecting a mix of sixteen different PFASs in drinking water samples at trace level concentrations is presented. The key characteristic of this method is the absence of an extraction step and a direct injection into the analytical system. Even the numerous difficulties faced in order to achieve its best performance, good recovery results, and really low Quantification Limits (5 ng/L) were achieved for all the compounds. The developed method was applied in the first assessment of PFASs occurrence levels in the Piedmont region of Italy. Despite the low detected pollution rates, a correlation between the "positive" sampling points and the potential pollution sources in the territory was done in order to understand their influence on the pollution levels and take decisions for reduction of contamination at source. The results showed that the number of point sources within a watershed significantly affects PFASs occurrence levels, providing us

with significant predictors for guiding future choice of sampling points at higher risk. However, the lack of information through the correlation study didn't allow for better assessment, highlighting the need for stronger cooperation and active participation between Regional Health and Environmental Protection Agencies, Water Companies and Stakeholders within policy making.

In conclusion, the results of this study highlighted the fact that chemical analysis alone is not able to evaluate the potential pollution of water bodies sufficiently. In order to take control measures for a safe and sustainable water supply is important to identify the hazardous components, their occurrence areas, and also the points at higher contamination risk. For this reason, and considering the costs, efforts and environmental impact of wide screening assessments, a "smart" monitoring program is better performing thanks to the prioritization of sites at major risks.

Chapter 4 Pharmaceuticals and Hormones

As described in Chapter 3, improvement of monitoring assessments for better evaluating the water bodies' quality is fundamental for better water supply and regulations. Hence, in this chapter is presented the second monitoring assessment done within this PhD thesis, followed a risk-based approach. More specifically, the geographical model developed in Chapter 3 was used in order to identify the points at higher risk of pollution and include them in the evaluation of pharmaceuticals and hormones levels in the Metropolitan Area of Turin. For the selection of the target compounds different analytical protocols' requirements were taken into account and a human health risk assessment for the detected concentrations – both for individual compounds and their mixed effects – was carried out in order to prioritize contaminants for treatment, with the higher aim of managing better a safe drinking water supply.

1. Materials and Methods

1.1 Selection of compounds

For this study, a target list containing different pharmaceutical compounds and hormones, was prepared based on the just revised European Drinking Water Directive (2020/2184/UE) [36], the requirements of the Regional Environmental Protection Agency (ARPA Piemonte) analytical protocol [142] and the NORMAN prioritization framework of emerging substances [35]. In this way, we concluded to sixteen different compounds: Ketoprofen, Atenolol, Trimethoprim, Ofloxacin, Azithromycin, Ciprofloxacin, Cyclophosphamide, Sulfamethoxazole, Erythromycin, Clarithromycin, and Caffeine (Table 10). Caffeine was included in this study as a tracer of anthropogenic pollution.

1.2 Reagents and Chemicals

Stock solutions of the target compounds were prepared in UHPLC-grade MeOH, all purchased from Sigma-Aldrich, Co (St. Louis, MO, USA). MilliQ was obtained from Millipore (MA, USA), LiChropur Formic Acid 98%-100% and LiChropur Ammonia (NH₃) solution 25% for LC-MS were purchased from Merck KGaA (Darmstadt, Germany), Ethylenediaminetetraacetic acid trisodium salt dihydrate (Na₄EDTA) and Hydrochloric Acid were obtained from Fluka Analytical (Sigma-Aldrich, MO, USA). Ammonium acetate for LC-MS was purchased from Fisher Chemical Scientific (Geel, Belgium).

Table 10. Selected compounds and their current regulation status in EU.

| Compounds | Chemical Group | CAS number | Regulation Status |
|-------------------|------------------------------------|------------|---------------------------------|
| Atenolol | β -Blockers | 29122-68-7 | NORMAN framework prioritization |
| Azithromycin | Macrolide Antibiotic | 83905-01-5 | EU Watch List/ARPA protocol |
| Clarithromycin | | 81103-11-9 | EU Watch List/ARPA protocol |
| Erythromycin | | 114-07-8 | EU Watch List/ARPA protocol |
| Caffeine | Stimulant | 58-05-2 | NORMAN framework prioritization |
| Carbamazepine | Anticonvulsant | 298-46-4 | NORMAN framework prioritization |
| Ciprofloxacin | Fluoroquinolones antibiotics | 85721-33-1 | NORMAN framework prioritization |
| Ofloxacin | | 82419-36-1 | NORMAN framework prioritization |
| Cyclophosphamide | Alkylating agent | 50-18-0 | NORMAN framework prioritization |
| Diclofenac | Analgesics anti-inflammatory drugs | 15307-79-6 | EU Watch List/ARPA protocol |
| Ketoprofen | | 22071-15-4 | NORMAN framework prioritization |
| Ibuprofen | | 15687-27-1 | NORMAN framework prioritization |
| Sulfamethoxazole | Antibacterial sulfonamides | 723-46-6 | EU Watch List/ARPA protocol |
| Trimethoprim | | 738-70-5 | EU Watch List/ARPA protocol |
| 17-beta Estradiol | Estrogens | 50-28-2 | EU Watch List/ARPA protocol |
| Estrone | | 53-16-7 | NORMAN framework prioritization |

1.3 Study Area and Sampling

The focus area of this study was the same with Chapter 3 (Figure 5, Table A1). However, in order to avoid a large-scale blind monitoring and all the costs and effort that has, a risk-based approach was followed. More specifically, in order to more sufficiently evaluate the pollution levels of PhACs in the area, a prioritization of the sampling points at higher risk was done based on the geographical model presented in Chapter 3. In this case, as potential pollution sources were considered 44 hospitals and care houses, and 24 major WWTPs, that occur within a radius of 5km from the 683 already existing sampling points in the SMAT network, usually included in regular monitoring campaigns (Figure 8). As a result, 270 sampling points were characterized as “hotspots” and were included in this study.

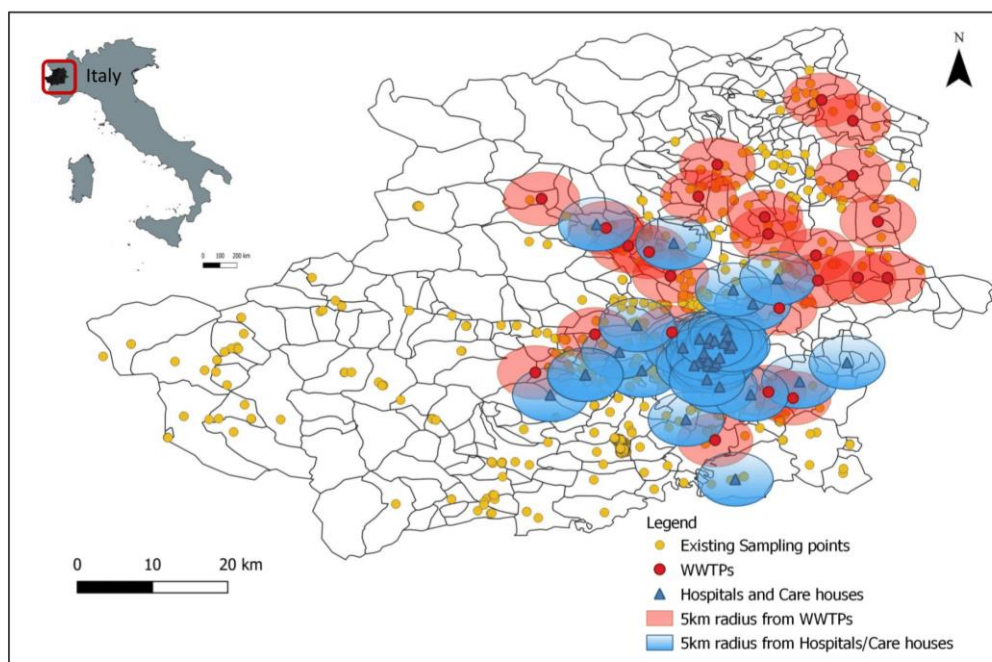


Figure 8: Map of the study area, including all the SMAT existing sampling points in the catchment areas, WWTPS and hospitals/care houses taken into account as potential pollution sources.

The sampling campaign was organized by SMAT, and carried out between October 2019 and October 2020. In total 328 samples including both raw and treated drinking water samples, following the specifications and requirements described in Chapter 3. As raw drinking water sources were considered surface (rivers, streams, and those taken at the

intake of SMAT's DWTP) and groundwater, while as treated were considered samples taken from fountains, tanks and the end of the SMAT's DWTP line. The samples were collected into amber glass bottles (1L) – previously decontaminated and rinsed with MeOH, according to the EPA 1694 method – stored at 4°C prior to their analysis and analyzed within 7 days of their sampling.

1.3.1 SMAT Drinking Water Treatment Plant

In figure 9 -which shows a photo of SMAT's DWTP- and figure 10 –which shows its schematic diagram- are shown the different pre-treatments steps of the plants. They include the catchment (D), where a wire mesh filter removes the majority of coarse contaminants and sediments, and a static horizontal flow pre-settling basin (E) consisting of a circular pool, equipped with a rotary dredge for the mechanical removal of sludge. In this stage additives can also be used in order to promote the sedimentation process. At the outlet of the pre-settling basin, the treatment plant is divided into three different lines. Po1 and Po2 (output 1100L/s), and Po3 (output 1500L/s). For Po3, an ozonation process (F) takes place, then the water is transferred to three CYCLOFLOC clarifiers (capacity of 1,5 m³/sec) with addition of aluminum polychloride. The basic principle of this system is the precipitation of sludge formed by the aluminum polychloride clumps and added microsand. During this clarification-flocculation stage (H), in one CYCLOFLOC basin, a further oxidation treatment is carried out with addition of sodium hypochlorite, for the elimination of ammonia and the nitrogenated compounds (G). Lastly, the water is filtered in two filtration batteries, each of which contains twelve overlapping units. The uppermost and lowermost filters, consist of 0,80 m thick filtering surface of granular activated carbon, trapping any particles left in the water after the settling process. After the filtration, the water is transferred into a tank where a final disinfection treatment with chlorine dioxide takes place in order to avoid the regrowth of bacterial colonies along with the distribution system during the delivery of the water to the users (I). For the two identical treatment lines Po1 and Po2, after the first sedimentation process, a pre-chlorination step takes place by means of chlorine dioxide and sodium hypochlorite, then the water is headed to an "Accelerator" type sludge recirculation tank for the clarification step (L). Subsequently, the treated water is filtered on granular activated carbon and accumulated in a tank (M) where the final disinfection step with chlorine dioxide occurs (Figures 9,10).

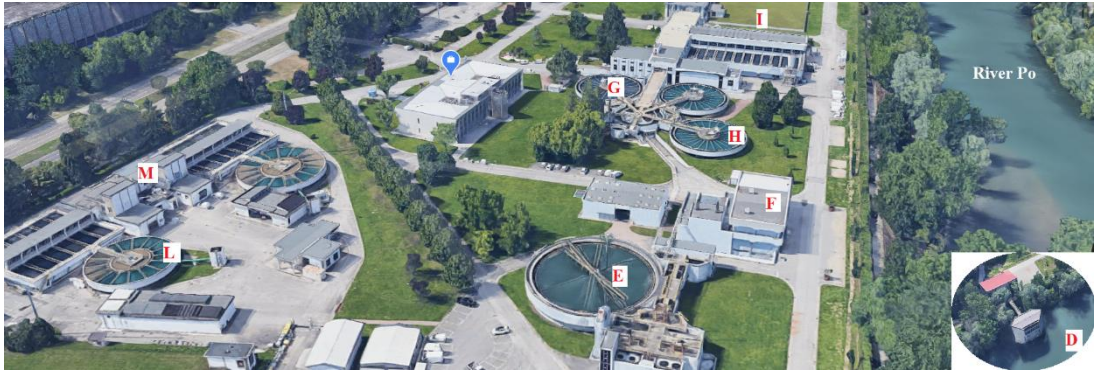


Figure 9: SMAT Drinking Water Treatment Plant

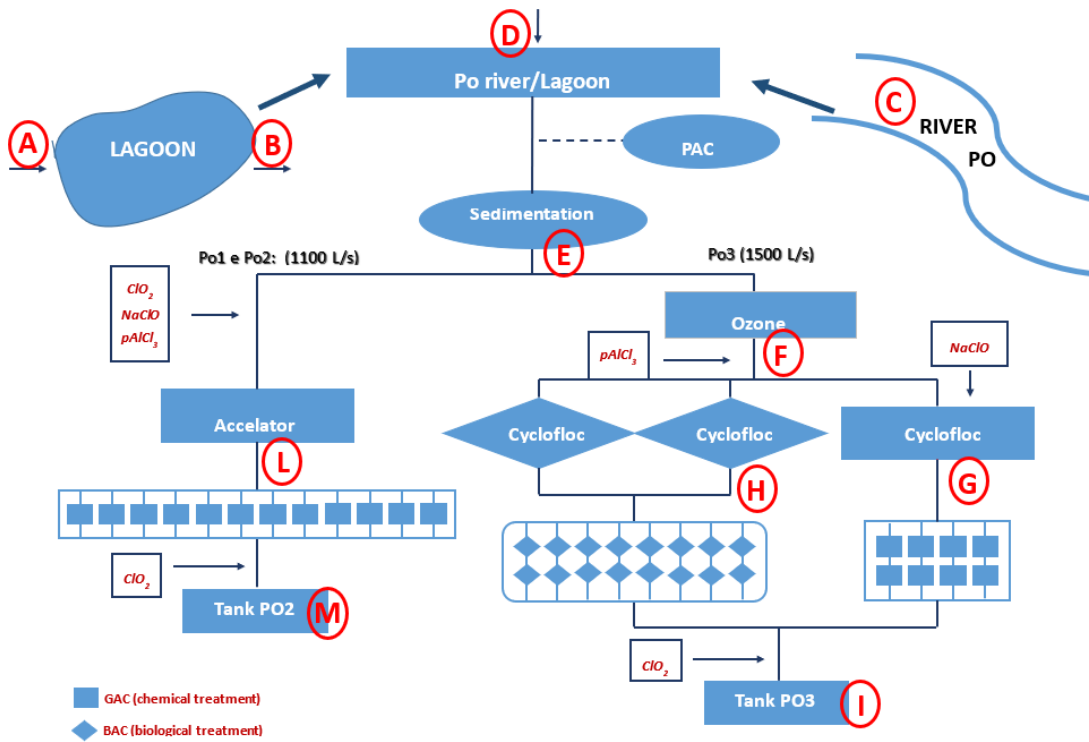


Figure 10: Schematic diagram of treatment lines in SMAT DWTP.

1.4 Sample preparation and instrumental analysis

In order to achieve adequate extraction of the mainly acidic target analytes, the samples' pH was adjusted to 2,0 using HCl, and after the addition of 500mg of Na₄EDTA to each of them. The 1L samples were loaded on to Oasis-HLB (200mg) solid phase extraction (SPE) cartridges (Waters, Milford, MA, USA) - which were preconditioned with 12 mL MeOH, followed by 6 mL MilliQ and 6 mL MilliQ with pH 2,0 - with a flow rate of 10mL/min. Elution of the analytes was done with 12 ml MeOH and after the solvent's evaporation with a rotary evaporator (BUCHI Rotavapor R-114), they were reconstituted in 1 ml MilliQ water. The chromatographic separation was achieved with a HPLC-triple quadrupole MS system equipped with a C18 HPLC column. The MS system operated in both Positive and Negative ESI using different MRM scan modes targeted to every analyte, while three subsequent HPLC methods were developed, due to the heterogeneity among the compounds. For the substances ionized in the Positive ESI mode (Table A5) a sample volume of 5µL was injected into a mobile phase consisted of a mixture of 0,1% Formic Acid in MilliQ Water (A) and Methanol (B), in a total run time of 10 minutes. The gradient profile, with a flow rate of 0,250 mL/min, started with 98% A and 2% B, increased to 100% B after 6 minutes, and after keeping this ration for 2 minutes, returned to the initial conditions. For the Negative ESI compounds (Table A5) a volume of 10µL of sample was injected into a mobile phase consisted of a mixture of 0,02% Ammonia in MilliQ Water (C) and Methanol (D) with a total run time of 10 minutes, while Ibuprofen was analyzed alone. In this case a sample volume of 10µL was injected into a mobile phase of a mixture of 0,1% Ammonium Acetate and 0,1% Formic Acid in MilliQ Water (C) and Methanol (D) in a total run time of 10 minutes. For both analysis cycles the gradient profile, with a flow rate of 0,400 mL/min, started with 98% C and 2% D, increased to 100% D after 6 minutes, and after keeping this ratio for 2 minutes, returned to the initial conditions.

1.5 Method Validation

For reassuring the applicability of this method a validation study was carried out according to the requirements of ISO/IEC 17025 [126] and the ICH method [139], described in Chapter 3. Six-point calibration curves with final concentrations of 1000, 2000, 4000, 6000, 8000 and 10000 ng/L (preconcentration factor of 1000 after the SPE step) were built for each target compound and used for quantification and calculation of linearity, trueness, uncertainty and recovery as well as LOD and LOQ for each target compound. Blank and quality control samples were analyzed in order to ensure the best performance of the instrument and the repeatability of the results during the whole analysis time. Quality control samples were prepared at a final concentration of 4000 ng/L (in the middle of the calibration curve range) and analyzed after every ten samples. The quantitation was performed using the MultiQuant™ 3.0.3 software (SCIEX, Framingham, MA, USA) and the 1/x concentration-weighting factor was applied.

1.6 Human Health Risk Assessment

A human health risk assessment study was carried out by comparing PhACs' concentrations to guideline values. The risk assessment was carried out for both individual compounds and their mixture, and as PhACs' concentrations were considered their average detected concentrations. In order to avoid wrong estimation, non-detects were considered at a value of ¼ of the individual LOD for each target molecule as proposed by Houtman et al. [62], since removing them or setting their values as zero would have under or overestimated their average concentrations. Moreover, compounds with n-octanol-water partition coefficient ($\log K_{ow}$) higher than 3 were not included in the risk assessment study as in general, there is a smaller possibility that they will pass all the drinking water treatment line steps and end up in the treated water [143]. For this reason, $\log K_{ow}$ values for each compound were obtained with the KOWWIIN algorithm of the EPI Suite 4.11 software [144], and only those with higher values than 3 were included in the study.

The Risk Quotient (RQ_i) for individual compounds (equation 11) was obtained by the ratio between the average Detected Concentration (MEC_i) and the corresponding guideline value or, where it didn't exist, the calculated provisional guideline value ((p)GLV) [143]. The pGLVs were calculated using the equation 12,

$$RQ_i = MEC_i / pGLV_i \quad (11)$$

$$pGLV_i (\mu\text{g/L}) = [\text{ADI} \times \text{BW} \times 10\% \text{ drinking water allocation}] / \text{DWI} \quad (12)$$

where ADI is the Acceptable Daily Intake ($\mu\text{g}/\text{kg bw}/\text{day}$); BW is the body weight set at a default value of 70kg, as it is the closest to the average European bodyweight value of 70,8 kg [146]; DWI is the drinking water intake (L/day) set at a default value of 2 L/day as reported by WHO; a 10% of drinking water allocation factor was taken into account as drinking water is not the only exposure way for humans [62,143,147]. ADI values for each target compound were obtained from the literature, and in the case of absence they were calculated after dividing N(L)OAEI values with an uncertainty factor of 100 [148]. For RQ values ≥ 1 there is the possibility of risk, if a lifelong exposure to the compound occurs only after drinking water consumption, while for RQ values $\leq 0,2$ the risk for adverse human health effects is considered negligibly low [62,143]. For the calculation of the mixed health Risk Quotient (RQ_{mix}) the Concentration Addition (CA) concept [149] was followed by comparing the sum of individual RQs.

2. Results

2.1 Validation and Quality Assurance

Good coefficient results with R^2 within the range of 0,995-0,999 were obtained for all the molecules, indicating good linear correlation. Concerning the systematic and random errors, satisfying results within the required ranges (as reported in Chapter 3) were obtained for each point of the calibration curve, with those obtained for 4000 ng/L being reported in Table 11. The recovery of the compounds after the SPE treatment was checked in 4 different real water samples at concentrations of 4 ng/L and 10 ng/L, and resulted in a range of 85,5-128% for all the compounds. For the calculation of the limits of Detection and Quantification, equations (9) and (10) were followed and resulted in a range of 0,010-3,492 ng/L for LOD and 0,034-11,369 ng/L for LOQ (Table 11).

Table 11. Validation results for every target compound

| Compounds | C (ng/L) | Trueness % | Uncertainty % | Linearity | LOD (ng/L) | LOQ (ng/L) |
|-------------------|-------------|---------------|------------------|-----------|---------------|---------------|
| Atenolol | 4000 | -3,977 | 3,047 | 0,9996 | 0,196 | 0,655 |
| Azithromycin | 4000 | -10,090 | 7,290 | 0,9951 | 0,736 | 2,454 |
| Caffeine | 4000 | -1,300 | 1,912 | 0,9991 | 0,322 | 1,073 |
| Carbamazepine | 4000 | -14,831 | 9,761 | 0,9999 | 0,066 | 0,219 |
| Clarithromycin | 4000 | -3,519 | 2,786 | 0,9996 | 0,031 | 0,074 |
| Ciprofloxacin | 4000 | -1,603 | 0,892 | 0,9996 | 0,788 | 2,625 |
| Cyclophosphamide | 4000 | -0,161 | 2,563 | 0,9996 | 0,010 | 0,034 |
| Diclofenac | 4000 | -7,627 | 2,531 | 0,9998 | 0,376 | 1,254 |
| Erythromycin | 4000 | -4,793 | 3,114 | 0,9998 | 0,244 | 0,814 |
| Ketoprofen | 4000 | -10,221 | 4,476 | 0,9999 | 0,115 | 0,385 |
| Ofloxacin | 4000 | -1,1769 | 2,735 | 0,9978 | 0,493 | 1,644 |
| Sulfamethoxazole | 4000 | -4,823 | 2,202 | 0,9983 | 0,110 | 0,366 |
| Trimethoprim | 4000 | -7,457 | 5,497 | 0,9998 | 3,492 | 11,369 |
| 17-beta Estradiol | 4000 | -7,129 | 6,546 | 0,9972 | 0,303 | 1,010 |
| Estrone | 4000 | -23,144 | 3,655 | 0,9971 | 0,400 | 1,333 |
| Ibuprofen | 4000 | -1,599 | 1,770 | 0,9969 | 0,412 | 1,375 |

2.2 Monitoring Assessment results

In total 325 samples were analyzed, of which 287 were groundwater and 24 surface water. For the samples with the highest detected PhACs concentrations, treated samples from the same areas were analyzed in order to reassure the quality of treated water, and take the appropriate countermeasures if necessary.

Concerning the raw drinking water sources, in 40 samples none of the target compounds was detected above their individual LOQs, while in 52 samples only one compound was detected. This result, highlights the fact that in the majority of the samples a mix of PhACs usually is present. The maximum number of coexisting compounds in one sample

in this study was eleven, and it was detected only in one groundwater sample. The point from which the sample was taken, is close to two WWTPs and one care house, confirming the higher risk of pollution when more point sources are closer to one sampling point, as indicated from the geographical model. The average detected concentration as sum of all the target compounds was found to be 28,32 ng/L, with a range between 2,02 and 523,36 ng/L in groundwater samples and 18,54 ng/L (2,02-82,05 ng/L) in surface water.

The range of individual detected concentrations was ranging between 0,08 ng/L and 483,94 ng/L, as it is concluded from the results reported in Table 12 and Figure 11. Only two out of the sixteen target compounds – ofloxacin and erythromycin- were not detected in the study area, in concentrations higher than their individual LOQs (1,64ng/L and 0,81ng/L for ofloxacin and erythromycin subsequently). Human consumption trends in the area, the physicochemical characteristics of the compounds enabling them to be adsorbed on different particles of the soil or biodegradation processes could explain the absence of these compounds. However, these results are in accordance with a study from Verlicchi et.al [64], that doesn't report higher than their individual method detection limits concentrations of Ofloxacin and Erythromycin in surface water from the Po Valley in Italy. On the contrary, the most detected compounds in the study area were Caffeine and Ketoprofen. Caffeine is considered as one of the most abundant compounds in the aquatic environment worldwide, and was present in 176 groundwater samples with an average detected concentration of 4,61 ng/L (1,15-65,92 ng/L), while in 23 surface water samples was detected with an average concentration of 5,34 ng/L within a range of 1,31-61,28 ng/L. However, its occurrence concentrations in the study area are significantly lower than those reported in other studies, whose ranges are in the scale of µg/L. The second most abundant compound in the area, was Ketoprofen as it was present in 143 groundwater samples, with an average concentration of 6,51 ng/L within a range of 0,16-152,98 ng/L, and in 21 surface water samples with an average concentration of 5,84 ng/L (0,43-71,84 ng/L). As shown by the results, Ketoprofen's concentrations varied significantly across the territory, with the areas closer to WWTPs showing higher levels, following trends from other studies [65,150], and could be correlated with socioeconomic aspects and consumption trends.

The next two more abundant compounds in the area were the two target estrogens, raising fears about their negative endocrine disrupting effects on humans and animals. Estrone's average concentration in the 117 groundwater samples that was detected was 4,03 ng/L, with a range between 1,09 and 125,97 ng/L, and in the 12 surface water

samples was 1,003 ng/L (1,30-8,33 ng/L). Concerning, 17-beta estradiol, which is the only compound from those included in this study that is subjected to a guideline (its concentration should not exceed 1 ng/L in drinking water), it's concentration in the 114 raw surface and groundwater that was present, was higher than the guideline level with an average of 1,50 ng/L, with the highest being detected in an area close to a hospital, highlighting the risks originating from discharges of untreated wastewater effluents. A mix containing only the two hormones was present in 24 samples, while in 148 none of the two was detected.

Table 12. Detection results of the target PhACs in the study area.

| Compounds | QF* n = 325 | C_{min} (ng/L) | C_{max} (ng/L) | C_{average} (ng/L) | C_{median} (ng/L) | Q1 (ng/L) | Q3 (ng/L) |
|-------------------|------------------------------|---|---|---|--|----------------------------|----------------------------|
| Atenolol | 12,54 % | 1,07 | 483,94 | 18,73 | 3,96 | 1,64 | 7,93 |
| Azithromycin | 4,18 % | 2,55 | 82,46 | 14,84 | 3,28 | 2,64 | 14,63 |
| Caffeine | 61,67 % | 1,15 | 65,92 | 5,69 | 3,53 | 2,21 | 5,51 |
| Carbamazepine | 39,37 % | 0,23 | 183,49 | 6,93 | 2,44 | 1,07 | 5,24 |
| Clarithromycin | 21,95 % | 0,10 | 101,30 | 7,57 | 1,48 | 0,40 | 4,60 |
| Ciprofloxacin | 3,83 % | 2,86 | 7,00 | 4,16 | 3,25 | 2,88 | 5,33 |
| Cyclophosphamide | 11,15 % | 0,08 | 1,10 | 0,31 | 0,26 | 0,19 | 0,34 |
| Diclofenac | 11,15 % | 1,26 | 121,46 | 12,41 | 3,62 | 2,22 | 11,89 |
| Erythromycin | 0,00 % | <LOD | <LOQ | <LOD | <LOD | <LOD | <LOD |
| Ketoprofen | 49,83 % | 0,4 | 152,88 | 8,28 | 2,58 | 1,40 | 7,31 |
| Ofloxacin | 0,00 % | <LOD | <LOQ | <LOD | <LOD | <LOD | <LOD |
| Sulfamethoxazole | 29,27 % | 0,41 | 99,47 | 4,94 | 1,91 | 0,92 | 3,68 |
| Trimethoprim | 2,79 % | 12,87 | 87,16 | 37,80 | 31,02 | 22,07 | 41,71 |
| 17-beta Estradiol | 36,59 % | 1,08 | 9,00 | 1,28 | 1,18 | 1,45 | 2,04 |
| Estrone | 40,77 % | 1,35 | 125,97 | 7,69 | 3,20 | 2,12 | 5,71 |
| Ibuprofen | 2,79 % | 1,46 | 10,54 | 3,77 | 3,15 | 1,78 | 3,73 |

*QF = quantification frequency

The highest concentrations of PhACs in raw drinking water sources of the study area were detected for Atenolol 483,94 ng/L, Carbamazepine 183,49 ng/L, Ketoprofen 152,88 ng/L, Estrone 125,97 ng/L, and Diclofenac 121,46 ng/L. All of them were detected in groundwater samples taken from points around WWTPs, indicating that conventional treatment techniques are not able to efficiently remove them from wastewater and highlighting the need to find new ones. As a conclusion, the findings of this study are in accordance with occurrence patterns in Italy and other countries, reported in literature [62,63,65,96,97,129,150-154].

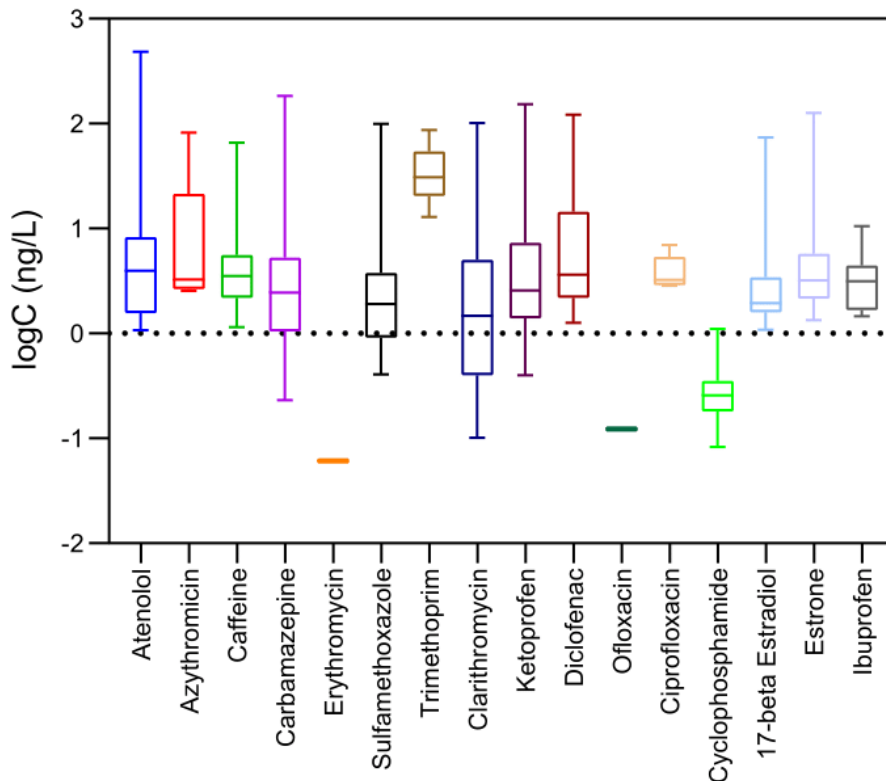


Figure 11: Boxplots showing the differences among the PhACs' detected concentrations in drinking water sources of the study area.

In literature the occurrence of PhACs in tap water around the world has been reported, as well, mainly due to insufficient treatment in DWTPs [97,148,151,155-157]. They claim that conventional treatment techniques like flocculation and sedimentation are not sufficient in removing PhACs and hormones from water completely, especially when

their concentrations are in trace levels (in the order of ng/L) [18]. However, addition of steps like ozonation and activated carbon filters, have be proved as a sufficient improvement of the treatment lines. Hence, the best solution for water companies in order to reassure a good and safe water quality supply is to combine these techniques. This concept has been incorporated in the study DWTP -as it is presented in Figures 9 and 10.

In order to evaluate the efficiency of the treatment lines in removing PhACs from water, treated samples were included in this study. The results showed that Atenolol, Azithromycin, Clarithromycin, Ciprofloxacin, Cyclophosphamide, Diclofenac, Erythromycin, Estrone, Ofloxacin, Sulfamethoxazole, Trimethoprim, and 17-beta-estradiol were not present in concentrations higher than their individual LOQs, in contrast with Carbamazepine, Caffeine, Ibuprofen and Ketoprofen. The reasons of these occurrence trends could vary among different phenomena, such as biodegradation, sufficient removal through adsorption on carbon filters and chlorination for those absent, and consumption trends and hydrophilic behavior ($\log K_{ow} < 3$) for those present. Nevertheless, the detected concentrations of PhACs after the treatment line were significantly lower than those found in raw water sources, as shown in Figure 12.

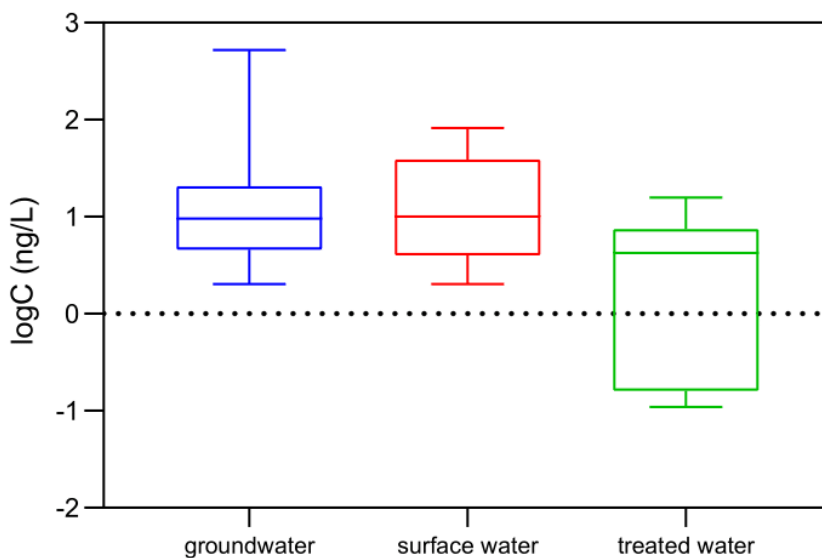


Figure 12: Average of sum detected concentrations \pm standard deviation for groundwater, surface and drinking water.

2.3 Human Health Risk Assessment

Based on the log K_{ow} values obtained, only ten out of the sixteen target compounds – Atenolol, Caffeine, Carbamazepine, Ciprofloxacin, Cyclophosphamide, Diclofenac, Erythromycin, Ofloxacin, Sulfamethoxazole, Trimethoprim - were resulted as potentially able to be present in drinking water. Even if they were not detected in the treated water samples, they were included in the risk assessment study. On the other hand, log K_{ow} values for Ibuprofen and Ketoprofen resulted higher than 3, which means that their hydrophobic character probably would not allow them to pass from all the treatment steps. However, since they were detected in treated samples they were included in the assessment as well.

Risk Quotients were calculated for both individual compounds and their mixtures, as the ratio between the average detected concentrations and the pGLV values. However, pGLV values could not be derived from toxicological data in literature – confirming the existence of knowledge gaps in PhACs risk assessment estimation. Hence, ADI values - and where not available N(L)OAEL values- were used for their calculation. All the ADI and N(L)OAEL values were obtained from literature, with the most restrictive value found, being selected (Table 13). The obtained pGLVs ranged from 0,07 $\mu\text{g/L}$ for Ofloxacin to 5285 $\mu\text{g/L}$ for Caffeine. The derived $RQ_{i\text{average}}$ were much lower than 0,2 (Table 13) ranging between $9,80 \times 10^{-7}$ for Cyclophosphamide and $2,21 \times 10^{-3}$ for Carbamazepine, indicating that none of the target compounds could potentially pose a risk of adverse health effects to humans, even after a lifelong exposure. In accordance with other studies, these outcomes show that even most of the compounds were detected in raw drinking water sources, they do not pose threats to human health individually [148, 158-160], mainly because of their low quantification frequency. However, in order to have a more holistic view of the risks and examine the need of strict measurements to guard pollution on specific points, the RQs were calculated also using the highest detected concentrations for each compound. The results obtained were again much lower than 0,2 ranging between $9,52 \times 10^{-6}$ for Cyclophosphamide and $1,54 \times 10^{-1}$ for Carbamazepine indicating low human health risks (Table 13).

In this context, since a calculation of the risks posed only from individual compounds would result in a total risk underestimation [148,161], the mix Risk Quotient (RQ_{mix}) was calculated as a sum of individual RQs. Since, toxicological data for mixtures of compounds are rare in literature, the Concentration Addition (CA) [161] concept was followed. This concept assumes that no interactions among the different compounds of

the mixture will occur, because of their same action mechanism and toxicity targets [149,162]. Moreover, it is expected that all the different components of a mixture will contribute to the total toxicity depending however on their concentration, resulting to the expectation that even if the individual compounds do not pose a risk, their mixture could potentially do, due to the addition effect [162]. In fact, the obtained results confirmed this assumption with the combined risk of exposure being higher than the individual one, but still it was negligibly low (lower than 0,2 for the sum of $RQ_{i\text{average}}$, and 0,2 for the sum of $RQ_{i\text{max}}$).

Table 13. Human health risk assessment parameters for target PhACs.

| Compounds | Log K_{ow} | ADI ($\mu\text{g}/\text{kg}$ bw/day) | Source | pGLV ($\mu\text{g}/\text{L}$) | MEC (ng/L) | $RQ_{i\text{average}}$ | $RQ_{i\text{max}}$ |
|-------------------|--------------|--|--------|------------------------------------|---------------------------------|------------------------|-----------------------|
| Atenolol | -0,03 | 2 | [163] | 7 | 2,29 | $3,27 \times 10^{-4}$ | $6,91 \times 10^{-2}$ |
| Azithromycin | 3,24 | N/A | N/A | N/A | N/A | N/A | N/A |
| Caffeine | 0,16 | 1510 | [164] | 5285 | 3,52 | $6,65 \times 10^{-7}$ | $1,25 \times 10^{-5}$ |
| Carbamazepine | 2,25 | 0,34 | [163] | 1,19 | 2,63 | $2,21 \times 10^{-3}$ | $1,54 \times 10^{-1}$ |
| Clarithromycin | 3,18 | N/A | N/A | N/A | N/A | N/A | N/A |
| Ciprofloxacin | -0,001 | 12 | [163] | 42 | 0,37 | $8,75 \times 10^{-6}$ | $1,67 \times 10^{-4}$ |
| Cyclophosphamide | 0,97 | 33 | [166] | 115,5 | 0,11 | $9,80 \times 10^{-7}$ | $9,52 \times 10^{-6}$ |
| Diclofenac | 0,57 | 200 | [166] | 700 | 1,5 | $2,14 \times 10^{-6}$ | $1,74 \times 10^{-4}$ |
| Erythromycin | 2,48 | 0,7 | [163] | 2,45 | 0,06 | 0 | 0 |
| Ketoprofen | 3,00 | 20 | [163] | 70 | 4,07 | $5,81 \times 10^{-5}$ | $2,18 \times 10^{-3}$ |
| Ofloxacin | -0,20 | 0,02 | [166] | 0,07 | 0,12 | 0 | 0 |
| Sulfamethoxazole | 0,48 | 510 | [166] | 1785 | 1,39 | $7,78 \times 10^{-7}$ | $5,57 \times 10^{-5}$ |
| Trimethoprim | 0,73 | 190 | [163] | 665 | 1,78 | $2,68 \times 10^{-6}$ | $1,31 \times 10^{-4}$ |
| 17-beta Estradiol | 3,94 | N/A | N/A | N/A | N/A | N/A | N/A |
| Estrone | 3,43 | N/A | N/A | N/A | N/A | N/A | N/A |
| Ibuprofen | 3,79 | 400 | [166] | 1400 | 0,2 | $1,52 \times 10^{-7}$ | $7,53 \times 10^{-6}$ |

3. Conclusions

Given the production, consumption and disposal trends of CECs in the aquatic environment, a successful implementation of the just issued Drinking Water Directive requirements is challenging. Finding more adequate strategies in order to protect the quality status of water bodies is necessary. Within this context and following the conclusions derived from the study presented in Chapter 3, a prediction of pollution risk approach prior to the monitoring assessment was followed. Spatial and statistical tools identified the areas at higher risk, based on the number of point sources present within the watershed. Raw water samples from these areas were included in a more comprehensive assessment aiming to quantify the occurrence levels of PhACs in drinking water sources, which was the first in the Piedmont region in Italy. A fast and green method using advanced analytical technique was developed and used for this study. The results confirmed the presence of the target compounds in the area, in concentrations of the ng/L scale, with those geographically closer to the considered pollution sources showing higher detection rates. A risk assessment study was then followed in order to evaluate the potential effects on human health, considering the mixture occurrence as well. Concluding, this study provided us with important information that will contribute to decisions making for a safer and more sustainable water management and supply system, by identifying point sources and potential health risks.

Chapter 5 Non-target screening monitoring assessment

The increasing number of chemicals released in the aquatic environment makes it practically impossible to evaluate the quality of water bodies, after following only target analyses dependent on individual standards. Therefore, suspect and non-target screening methods can be used, as they are able to reveal the full spectrum of occurring compounds, and provide valuable information. High-resolution mass spectrometry is a fundamental tool in order to facilitate these analyses.

In this study, a non-target screening of surface water samples was carried out in order to evaluate the current status of different European bodies. SWATH-HRMS acquisition mode – a technique relatively new and not widely studied in the field of environmental monitoring assessments - was used in order to detect as many analytes as possible, without losing those of lower intensity. Finally, multivariate statistical tools were used in order to identify pollution patterns among the samples, and find the compounds that are responsible for their discrimination.

1. Materials and Methods

1.1 Sample collection

In this study 17 surface water samples from different points in two different European countries were included. The samples concerned river water – taken from the Po river, in Turin, Italy – and lake water – collected from 2 different lakes in Italy, Orta and Comabbio and the lake Pamvotis in Ioannina, Greece. All sampling points were selected to be close to known pollution sources. Information about the sampling points are summarized in table A6, while their maps in Figure A2. The sampling campaign was executed between July and September 2020. For each sampling location, 1L of water was collected into amber glass bottles with stoppers of Teflon lined screw caps. The samples were filtered with 0,7 μ m GF-F fiberglass filters (Whatman, UK) and stored in dark at 4°C prior to their extraction and analysis.

1.2 Reagents and Chemicals

MilliQ was obtained from Millipore (MA, USA), UHPLC-grade Methanol was purchased from Sigma-Aldrich (Saint Louis, MO, USA), LC-MS grade Water LiChrosolv® and Ammonium acetate for LC-MS LiChropur® were purchased from Merck KGaA (Darmstadt, Germany). Analytical standards of 100 compounds were included in this

study, and their information are summarized in Table A7. As an internal standard (IS) the Flunixin-d3 was used.

1.3 Sample preparation

For the extraction of the analytes two protocols using different SPE materials were followed. The Oasis HLB (200mg/6 mL) SPE cartridges (Waters, Milford, MA, USA) were used for the extraction of non-polar and slightly polar compounds, while the ENVI-Carb Plus (0,4 g/1 mL) reversible tubes (Sigma-Aldrich, Co, St. Louis, MO, USA) were used for high polarity compounds. Each sample was divided into two 500 mL portions and each portion was spiked with 1 µg/L Internal Standard (IS) solution. The first 500 mL portion of each sample was loaded on to the Oasis-HLB cartridges, previously preconditioned with 5 mL MeOH, followed by 5 mL HPLC-grade water at a flow-rate of 5mL/min. Elution of the analytes was done with 2x5 mL MeOH. Extracts were evaporated until dry under a gentle nitrogen stream and reconstituted in 500µL of 10:90 (v:v) MeOH:H₂O. The second 500 mL portion of each sample was loaded on to the ENVI-Carb cartridges, after conditioning them with 1 mL Methylene Chloride, followed by 2x3 mL MeOH and 3 mL Deionized water at a flow rate of 5mL/min. The elution was conducted with 2x5 mL of 50:50 MeOH: Methylene Chloride (v/v) after arranging them in the forward direction. The obtained aliquot was evaporated until dry, under a gentle nitrogen stream and reconstituted to 500µL of 10:90 (v:v) MeOH:H₂O. Three replicates of control samples for each extraction protocol were prepared from MilliQ water.

1.4 Instrumental analysis

Analyses were carried out using a Quadrupole-time-of-flight mass spectrometer (X500R QTOF, SCIEX, Framingham, MA, USA) coupled to an ultra-performance liquid chromatography (UHPLC) system (ExionLC, Shimadzu, Japan). The chromatographic separation was achieved using a Luna® Omega Polar C18 100 LC Column (3µm particle size, 100 x 2,1mm) heated at 40°C, by injecting a 50µL sample volume into the mobile phase at a flow of 0,350 mL/min. The mobile phase used for the positive ionization mode consisted of a mixture of 5mM Ammonium Formate in H₂O (A) and 5mM Ammonium Formate in MeOH (B), and the elution followed a gradient profile starting from 95% A and 5% B, keeping the ratio for 1 minute and then gradually changing to 100% B within 14 minutes. This profile was kept for 2 minutes and then gradually reversed into the initial conditions until 20 minutes of elution, accordingly also for the negative ionization. For the negative ionization mode, the mobile phase consisted of a mixture of 5mM Ammonium Acetate in H₂O (C) and 5mM Ammonium Acetate in MeOH (D). Also in this

case the elution followed a gradient profile, starting from 95% C and 5% D, holding this ratio for 1 minute and then gradually changed to 100% B within 14 minutes, which was kept for 2 minutes and then gradually reversed into the initial conditions until 20 minutes of elution. During the whole analysis period, the samples were cooled at 4°C inside the autosampler.

The X500R QTOF source parameters for the positive polarity were as follows: ion source gas 1: 45 psi, ion source gas 2: 55 psi, curtain gas: 30 psi, collision gas (CAD): 7 psi, temperature: 350°C; spray voltage: 5500V. The parameters used in full-scan MS mode were as follows: accumulation time: 0,05 sec; declustering potential: 50 V; TOF start mass: 100 Da; TOF stop mass: 1000 Da. A generic collision energy spread of 35 ± 15 was used. For the Q1 isolation strategy (MS/MS) the parameters were TOF start mass: 50 Da; TOF stop mass: 1000 Da; total number of windows: 24; window accumulation time: 0,035 s. An external calibration was performed daily, using a mixture of 10 compounds with a mass range between m/z 132,90 and m/z 2034,63. This mixture was also automatically injected every 5 samples in order to maintain the mass accuracy below 2 ppm. Similarly, the source parameters for the negative polarity were as follows: ion source gas 1: 45 psi, ion source gas 2: 55 psi, curtain gas: 30 psi, collision gas (CAD): 7 psi, temperature: 350°C; spray voltage: -4500V. The parameters used in full-scan MS mode were as follows: accumulation time: 0,05 s; declustering potential: -80 V; TOF start mass: 100 Da; TOF stop mass: 1000 Da. A generic collision energy spread of $(-35) \pm 15$ was used. For the Q1 isolation strategy (MS/MS) the parameters were: TOF start mass: 50 Da; TOF stop mass: 1000 Da; total number of windows: 24; window accumulation time: 0,035 sec. An external calibration was performed daily, using a mixture of 10 compounds with a mass range between m/z 68,99 and m/z 2233,91. This mixture was also automatically injected every 5 samples in order to maintain the mass accuracy below 2 ppm. SCIEX OS 1.7 software (SCIEX, Massachusetts, USA) was used for data acquisition and elaboration.

1.5 Data elaboration

In general, the SCIEX OS software accepted as features the detected fragments that had at least 10 data points across each peak, a minimum intensity of 2000 counts/s, were within $\pm 0,01$ Da and had a peak area 10 times greater than the blank sample. Moreover, it performed peak alignment across samples if features were detected within $\pm 0,2$ minutes from each other, and assigned adducts. Due to the large volume of the obtained data, some reduction steps - after visual inspection of the detected features - were

followed, based on low peak intensity, bad integration quality, isotopes and adducts from the analytical apparatus removal. The last step in the reduction process was to remove features that were present in the blank, due to the fact that even if the software subtracted the blank prior to the peak list formation, it gave the possibility to use only one sample.

Moreover, elaboration of the data was done also with the software MZmine 2.52 [167] in order to prepare the tables for statistical analysis. Firstly the raw data obtained from the instrument were converted into .mzXML format (compatible with MZmine) using the software ProteoWizard [168]. Then they were loaded in MZmine and were elaborated following the steps of peak picking, peak deconvolution, peak alignment, and isotope removal. The parameters used are summarized in Table A8. The result was a table which as rows had the observations (samples) and as columns the variables (detected features-ions).

1.5.1 Suspect Screening workflow

For the suspect screening, 100 analytes (Table A7) were taken into account. Their selection was done according to the in house availability of chemical standards. A mix of them was prepared (100 ng/L) in 10:90 (v:v) MeOH:H₂O and analyzed at the end of the sequence in order to obtain MS spectra and retention time data and compare them with those found in the samples for facilitating and enhancing reliability in the identification.

1.5.2 Non-target screening workflow

After data reduction, the instrument's software prepared peak lists including information about the exact m/z value, retention time RT (min), and peak area for each detected feature in each sample. All the peaks were visually inspected for shapes and intensities, and those not showing satisfactory results not being included in the further identification process. Moreover, peaks were inspected for isotopic patterns, with those that in their MS1 (precursor ions) spectra had peaks with differences of ¹H, ³⁷Cl, or ⁸¹Br, from the major ion peak, being further processed. Moreover, the SCIEX OS software provided molecular formulas for each peak, based on the fragmentation in MS1 (precursor ion) and MS2 or MS/MS (product ion) spectra, following specific parameters. More specifically, atoms up to C₄₉H₇₅Br₂Cl₅F₃I₃N₁₀O₁₆P₁S₃ were considered, the mass error of the parent ion had to be ± 5 ppm and the MS/MS fragments had to support the proposed formula within an error of ± 10 ppm.

For the identification of the compounds, libraries provided from the SCIEX OS software were used. They included entries for 4656 components, including pesticides, pharmaceuticals, hormones, personal care products, perfluoroalkyl substances and different toxins with high resolution data. SCIEX OS evaluated the matching grade between the measured MS spectra and those registered in the libraries, and scored each match on a scale of 0 to 100. Only those with a compatibility score of more than 65 were considered, and were manually inspected in order to discard false matches, like those for whom parent ion and molecular formulas didn't match. For those features that a match in the instrument's libraries was not found, searches on the online library ChemSpider, to which SCIEX OS is directly connected, were done. ChemSpider also offers the possibility of *In silico* fragmentation, for spectral predictions that could enhance the identification process.

1.5.3 Statistical analysis

The reduced peak lists, containing information about accurate mass, retention time, and peak area were exported from MZmine and further processed using Microsoft Excel in order to transform it into a table compatible with the SIMCA 14.1 software, which was used for the multivariate analysis of the results. Principal Components Analysis (PCA) and Partial Least Square Discriminant Analysis (PLS-DA) were done in order to identify patterns among the samples, while the Variables Importance in Projection (VIP) technique was used in order to categorize the features that are responsible for the discrimination among the samples. Unit Variance (UV) and Pareto scaling were used as weighting methods, in order for every variable to give equal relevance to the model.

2. Results

2.1 SWATH-MS acquisition

When performing a non-target LC-MS analysis of water samples, many analytes are eluting from the column and are entering the mass spectrometer. Using a Data Dependent Acquisition (DDA) mode, a full scan MS - as Q1 isolation- on a broad range of m/z values is acquired and the peaks of the analytes are categorized based on their intensity. However, MS/MS fragmentation is captured only for the most abundant analytes of the sample, risking to lose information about compounds at a lower concentration level. For this reason, in this study we used a Data Independent Acquisition (DIA) technique – the so-called SWATH Acquisition. In that mode, the mass spectrometer uses a wider Q1 isolation size -with the possibility to create smaller Q1 windows in the m/z regions where the most compounds are expected to occur- and

collects MS/MS spectra for every detected analyte, by reducing the risk to lose low abundance analytes in complex samples.

However, this type of acquisition in non-target screening analyses generates massive data, that are difficult to handle and require a proper preprocessing workflow in order to obtain reliable information. In this study, 38631 features were detected in total, from which 20796 in the positive and 17835 in the negative ESI mode. Therefore, elaboration of the data prior to identification was done, including the following steps: (a) peak picking, including the removal of chromatographic peaks, based on peak intensities and shapes, (b) blank subtraction, including removal of features present in the matrix, (c) peak alignment, where detected features' peaks are linked among different samples, (d) adducts and isotopes grouping and removal. In this way, we resulted to peak lists containing information for 2741 features detected in Positive and 2526 detected in Negative ESI mode. The high percentage of decrease on the data volume is reported and confirmed also by other studies, that follow this procedure [169,170].

2.2 Multivariate statistical analysis

Due to the complexity and the big volume of data obtained from HRMS non-target screening assessments, statistical analysis is a fundamental processing step that allows better results' visualization. Thus, the reduced peak lists need to undergo further elaboration in order to prioritize features for identification. Multivariate Data Analysis (MVDA) is a useful approach for handling big data and achieving maximum recovery of information [171]. Explanatory analysis of these data can be done based on unsupervised methods such as PCA and Hierarchical Clustering Analysis (HCA), or supervised methods like PLS-DA.

PCA is a multivariate technique that aims to convert a set of observations, possibly related to each other, into a new set of small variables, which have the property of being linear combinations of their initial observations, without being related to each other. Basically, PCA is used for data visualization by reducing their dimensionality into orthogonal principal components (PC) that explain the largest amount of variance among the observations. The first PC (PC_1) defines the highest variance among the samples, the second PC (PC_2) is a vector of 90° to PC_1 that defines the highest variance that can't be explained from PC_1 , and finally PC_n is a vector of 90° to PC_{n-1} that defines the highest variance that can't be explained from PC_{n-1} . PLS-DA is another multivariate method, which aims to investigate the features that are more statistically important for the discrimination among the observations. More specifically, in the PLS-DA model the X-matrix -which contains information about the variables and observations- is transformed by adding a Y parameter, which contains the classification of the

observations. Basically, PLS-DA is a linear classification model between the X-matrix and the Y-observations that enables the selection of the most predictive or discriminant features in the data which help to classify the observations [171].

In this study, PCA was performed on the data for examining the relationships between different samples and study the possibility of different locations or sample types being responsible for their variance. The PCA models were constructed after transformation of the data with two types of scaling, the UV and the Pareto, in order for every variable to give equal relevance to the model. In UV scaling every variable is multiplied with the ratio $1/sd_j$, where sd_j is the variable's standard deviation, while in Pareto is multiplied with the ratio $1/\sqrt{sd_j}$. The coefficients R^2 and Q^2 were used for evaluating the goodness of fit and prediction of every model to the dataset used. R^2 explains how well the model fits the data, with high values (close to 1) indicating the best conditions, while Q^2 explains how well the model predicts the data, with values $> 0,5$ indicating good predictivity. These two coefficients tend to have a completely different behavior as the number of PCs increases. More PCs result to an increasing R^2 value (going closer to 1), showing a better fitting of the data. On the contrary, the Q^2 value reaches a maximum, after of which the addition of PCs is decreasing the predictivity of the model. Concerning the PLS-DA models, were constructed taking into account the two different scaling types, and were validated using the Permutation test. This test uses random tests in order to assess the risk that a difference between the observations is random and not statistically significant. More specifically, its idea is to compare the goodness of fit (R^2 and Q^2) of the original PLS-DA model with the goodness of fit of several random tests, based on the alternation of the Y-observations while the X-matrix are kept intact. A model is valid when the R^2 and Q^2 values of random tests are lower than those of the initial model, and when the regression line between the Q^2 values of the initial test and of the random tests intersects the vertical axis at, or below zero (Figure A3). Finally, the VIP plot was used in order to rank the variables (detected features). The VIP values are calculated as the sum of PLS squares, weighted by the sum of squares explained in each model. VIP values > 1 are considered as important and responsible for the discrimination among the samples variables, while VIP values $< 0,5$ are considered as not important [169-171].

Figure 13 shows the obtained PCA models for the features detected in ESI (+). In the UV scaling-PCA the variability explained by PC_1 was 22%, and 11% by PC_2 , with R^2 equal to 0,221 and Q^2 equal to 0,108. In the Pareto scaling-PCA model, PC_1 explained 25% of the variability and PC_2 15%, with R^2 equal to 0,250 and Q^2 equal to 0,131. These information show that the Pareto scaling-PCA model fits and predicts better the data of this study. Nevertheless, in both models samples S2 and S10 cluster closely together in the score plot and far from the rest of the samples, indication similarities in the chemical composition. Both samples are collected from the two different lakes in Italy (Table A6).

Moreover, in the score plot the sample S9 is plotted far away from the rest of the samples, indicating the occurrence of different compounds. All these three samples, which were collected from the two Italian lakes, were the most chemical dissimilar, highlighting the influence of different locations on the variance. Samples S12, S5, S7 were clustered closely to the other samples, with S12 being alone and the S5, S7 being closer. This result is reasonable, since S5 and S7 are samples from the same lake while S12 concerns a river sample. Finally, the rest of the samples, even if originating from different lakes in different countries are plotted together, without highlighting variance among them.

Figure 14 shows the obtained PLS-DA models (in both scaling types) for the features detected in ESI (+). In both models is obvious the classification of the samples between countries (green-Greece, blue-Italy), while S2 and S10 are clustered together and far from the rest of the samples. In the UV scaling-PLS-DA the variability explained by PC₁ was 83%, and 55% by PC₂, with R² equal to 0,831 and Q² equal to 0,398. In the Pareto scaling-PCA model, PC₁ explained 81% of the variability and PC₂ 49%, with R² equal to 0,810 and Q² equal to 0,331. These information show that the model obtained after the transformation of the data with the UV scaling can predict better the features responsible for the discrimination among the observations. However, the validation of the two models showed that it was not valid (Figure A3), since it was not fitting satisfactorily the data (R² value was very close to 1, while the intersection of the Q² regression line with the vertical axis was higher than 0, indicating the overestimation of the model). On the other hand the Pareto-scaling PLS-DA model provided satisfactory results. Hence, it was used in order to build the VIP plot, and select the significant features for the discrimination among the samples, that would be identified (Table 16).

The models for the data detected in ESI (-) are not reported, since they resulted in overestimation of the variance among the samples, due to the high number of zero values.

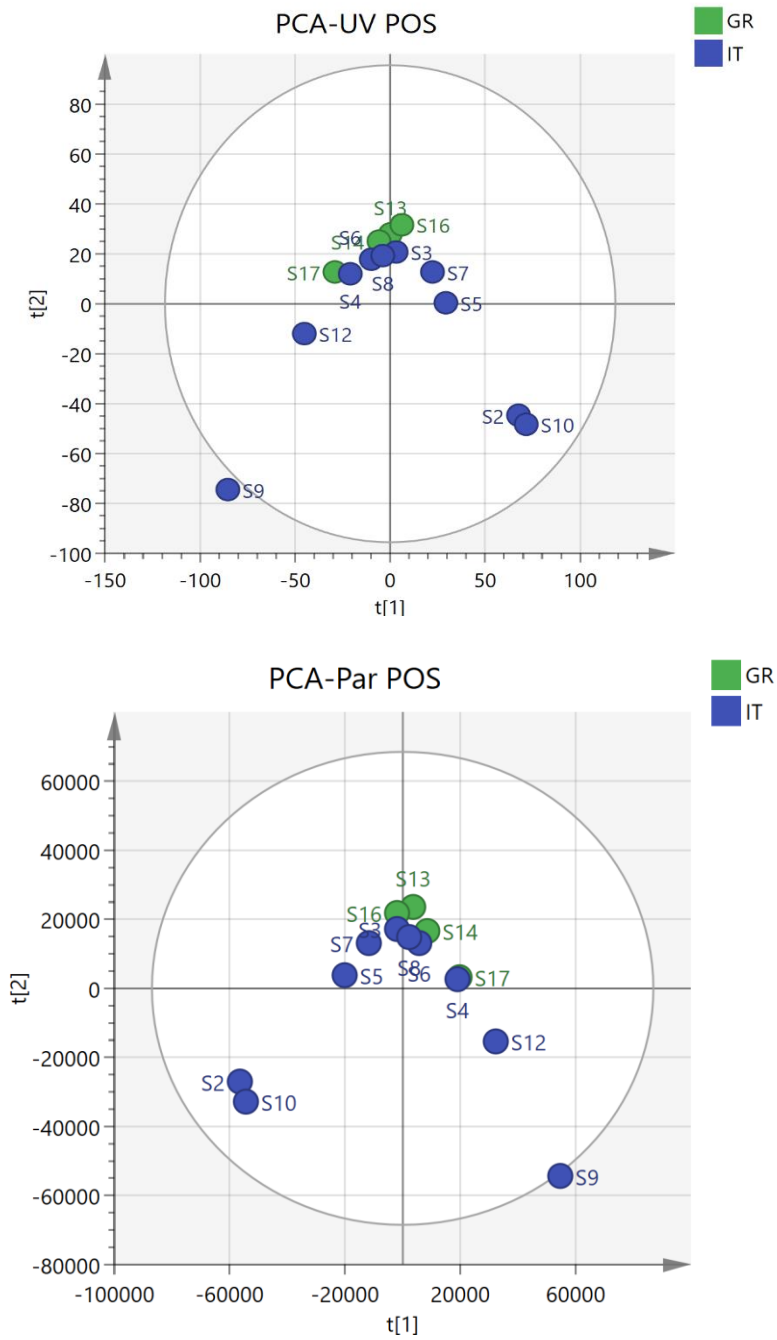


Figure 13: PCA models with UV and Pareto Scaling, for the features detected in ESI (+).

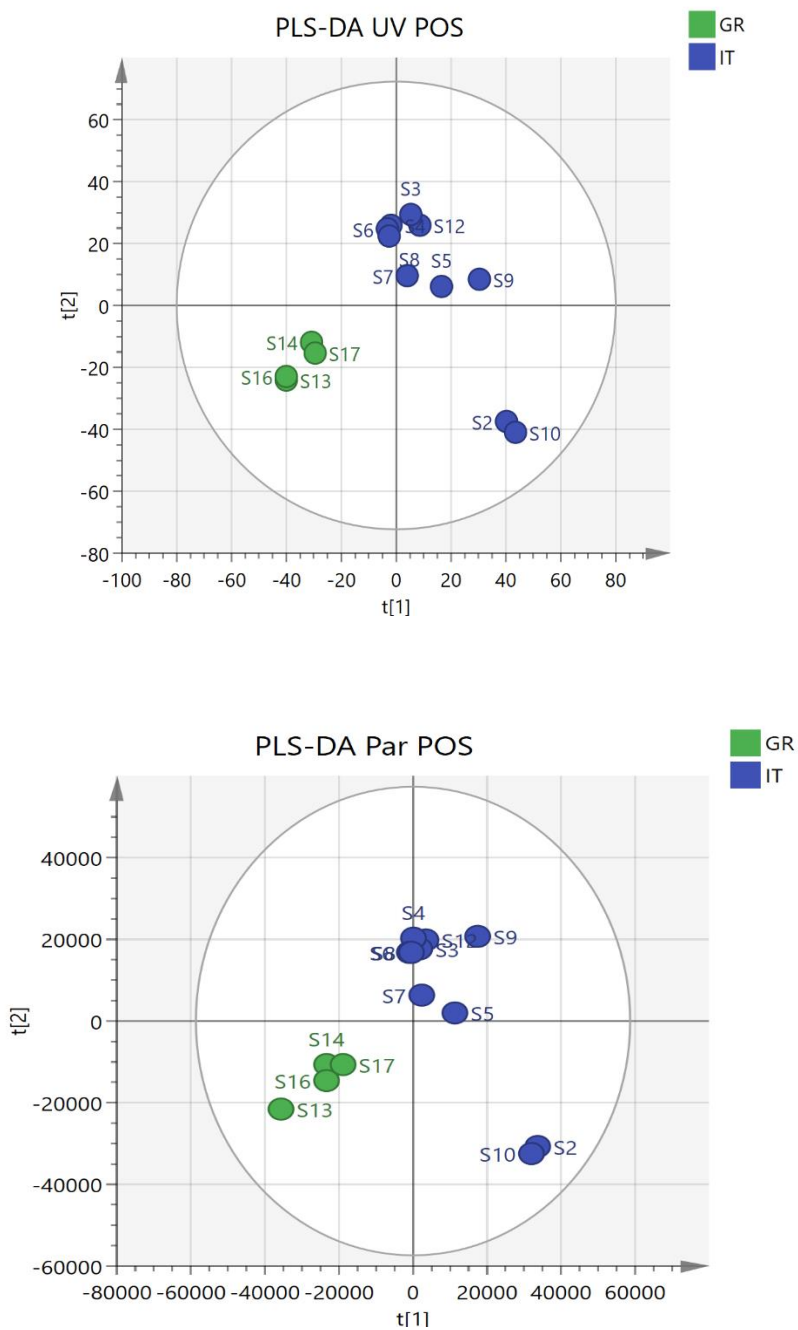


Figure 14: PLS-DA models with UV and Pareto Scaling, for the features detected in ESI (+).

2.3 Suspect and non-target screening

The suspect screening compounds detected in the samples and their responses are summarized in Table 14. All the suspect screening detections are confirmed identifications; as analytical standards were used for these analyses. Pesticides, like Terbutylazine, Atrazine, Metolachlor and Primicarb were the most detected compounds in the samples, followed by the pharmaceuticals Carbamazepine, Ibuprofen, Clarithromycin and Sulfamethoxazole. PFAS was the next most frequently detected class of compounds, with PFOA and PFOS being detected in 7 out of the 17 samples.

Table 14. Suspect screening analytes' detection information.

| Compound | Observed <i>m/z</i> | RT (min) | Compound | Observed <i>m/z</i> | RT (min) |
|-------------------|------------------------|-------------|-------------------|------------------------|-------------|
| 17-beta estradiol | 271,3814 | 5,16 | Epoxiconazole | 330,0897 | 6,27 |
| Acetamidrid | 223,0743 | 7,89 | Molinate | 188,1109 | 10,86 |
| Acetochlor | 270,1262 | 11,12 | Ofloxacin | 362,1582 | 5,19 |
| Alachlor | 270,1143 | 12,54 | Oxadiazon | 362,0819 | 3,52 |
| Ametryn | 228,1271 | 10,40 | Penconazole | 284,0719 | 11,35 |
| Atenolol | 267,1707 | 2,90 | Pendimethalin | 282,1156 | 15,23 |
| Atrazine | 216,1007 | 9,95 | Pethoxamid | 296,1414 | 11,13 |
| Atrazine-desethyl | 188,0701 | 8,00 | PFHxA | 312,9730 | 9,05 |
| Azythromycin | 749,5164 | 8,95 | PFOA | 412,9662 | 10,31 |
| Boscalid | 343,0411 | 10,6 | PFOS | 498,9321 | 12,54 |
| Caffeine | 195,0877 | 6,89 | Phosalone | 368,0218 | 7,26 |
| Carbamazepine | 237,1019 | 9,64 | Pirimicarb | 239,1498 | 9,67 |
| Chlorfenvinphos | 358,9777 | 11,5 | Prochloraz | 376,0388 | 11,66 |
| Chloridazon | 222,0429 | 7,76 | Procymidone | 301,0241 | 8,91 |
| Chlorotoluron | 213,0791 | 9,75 | Prometryn | 242,1419 | 5,37 |
| Ciprofloxacin | 332,1457 | 4,09 | Pronamide | 257,1238 | 11,03 |
| Clarithromycin | 748,4841 | 10,70 | Propachlor | 212,0837 | 10,08 |
| Cyanazine | 241,0966 | 9,07 | Propazine | 230,1166 | 10,52 |
| Cyclophosphamide | 262,0792 | 8,19 | Propiconazole | 342,0874 | 9,34 |
| Dichlorvos | 220,9541 | 13,23 | Pyraclostrobin | 388,1058 | 11,56 |
| Diazinon | 305,1078 | 11,50 | Pyrimethanil | 200,1182 | 10,48 |
| Diclofenac | 296,1163 | 2,45 | Parathion (ethyl) | 292,1762 | 3,29 |
| Dimethenamide | 276,0822 | 10,70 | Simazine | 202,0855 | 9,27 |
| Ethofumesate | 287,2694 | 1,48 | Spirotetramat | 374,197 | 11,10 |
| Erythromycin | 734,4693 | 10,20 | Spiroxamine | 298,2745 | 10,69 |
| Estrone | 269,1529 | 5,27 | Sulfamethoxazole | 254,0598 | 7,13 |

| Compound | Observed <i>m/z</i> | RT (min) | Compound | Observed <i>m/z</i> | RT (min) |
|------------------|------------------------|-------------|-----------------------------|------------------------|-------------|
| Fenamidone | 312,1176 | 10,53 | Tebufenozide | 353,1672 | 14,92 |
| Fenbuconazole | 337,1211 | 11,15 | Terbuthylazine | 230,1165 | 16,69 |
| Fenhexamid | 302,071 | 10,91 | Terbuthylazine- desethyl | 202,0855 | 15,06 |
| Hexazinone | 253,1659 | 9,38 | Tetraconazole | 371,9982 | 7,93 |
| Ibuprofen | 204,9879 | 1,56 | Thiacloprid | 253,0310 | 8,39 |
| Indoxacarb | 528,0856 | 14,20 | Thiamethoxam | 292,0012 | 4,29 |
| Kresoxim-methyl | 314,1616 | 10,82 | Thiobencarb | 258,0719 | 11,6 |
| Ketoprofen | 255,1026 | 10,36 | Triticonazole | 318,1342 | 5,18 |
| Lenacil | 235,1448 | 10,13 | Tolyfluanid | 364,0021 | 9,35 |
| Linuron | 249,0193 | 10,46 | Trimethoprim | 291,1446 | 6,73 |
| Metazachlor | 278,106 | 10,09 | Vinclozolin | 287,1192 | 4,49 |
| Methyl parathion | 264,1987 | 3,54 | Zoxamide | 336,0391 | 6,38 |
| Metolachlor | 284,1412 | 11,26 | | | |

Concerning the non-target screening, the identification of the compounds was done based on libraries using the accurate mass, retention time, isotopic pattern and MS/MS spectra of the detected ions. In this way, 28 compounds were identified in the negative ionization mode, and 327 in the positive (with VIP values >1). This big difference between the numbers of detected compounds in ESI (+) and ESI (-) can be explained by the lack of available information on the libraries, as well as the ionization type of the majority of the compounds. Among the results, a variety of chemicals was detected including pharmaceuticals - 39% in ESI (+) and 21% in ESI (-), pesticides - 34% in ESI (+), PFAS - 17% in ESI (-), toxins, food additives, drugs and other. These results are summarized in Tables 15 and 16.

Table 15. Compounds detected in ESI (-).

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples |
|---|---------------------------|------------------------|-------------|---|
| 15-Acetyldeoxynivalenol | 337,166 | 337,1636 | 1,39 | 1,2,3,5,6,7,8,10,11,12, 13,14,15,16,17 |
| Agistatin E | 227,093 | 227,0925 | 1,39 | 1,3,5,6,7,10 |
| Altersolanol A | 334,961 | 334,9592 | 2,15 | 11,3 |
| Asperlactone | 183,067 | 183,0668 | 1,39 | 14 |
| Aspinonene | 187,098 | 187,0981 | 1,37 | 14,15,16,3,6,8 |
| Aspyrone | 183,067 | 183,0661 | 1,41 | 13,15,2,4,5,6,7,9,10 |
| Chanoclavine | 255,102 | 255,1022 | 14,75 | 17,12 |
| Closantel- ¹³ C ₆ | 667,060 | 667,0618 | 18,40 | 11 |
| Deoxynivalenol | 295,228 | 295,2284 | 1,44 | 13,14,15,16,17,11,12, 1,2,3,5,6,7,8,10 |
| Fatty acid C20:4 Garbage | 303,234 | 303,2335 | 18,95 | 2,3,4,5,7,8,9,10 |
| Heptelidic acid | 279,124 | 279,1239 | 1,42 | 8 |
| Ibuprofen | 204,991 | 204,9879 | 1,56 | 11 |
| Isofusidienol A | 299,260 | 299,2592 | 17,75 | 11 |
| Mycophenolic acid | 319,142 | 319,1398 | 1,45 | 12 |
| Nivalenol | 311,114 | 311,1134 | 1,42 | 1,2,3,4,5,6,7,8,9,10,11 ,12,13,14,15,16,17 |
| Norsolorinic acid | 369,244 | 369,2434 | 18,54 | 17 |
| Palitantin | 253,144 | 253,1436 | 1,64 | 14 |
| Patulin | 153,093 | 153,0916 | 1,49 | 3 |
| Penicillic acid | 169,088 | 169,0861 | 1,38 | 13 |
| Pentobarbital Negative | 225,074 | 225,0736 | 13,69 | 17 |
| PFBA | 213,056 | 213,0563 | 15,74 | 11,12 |
| PFHpA | 362,970 | 362,9688 | 15,21 | 1,16,12,11,3,4 |
| PFHxA | 312,973 | 312,9728 | 12,82 | 1,2 |
| PFOA | 412,966 | 412,9659 | 16,02 | 12,1,16,3,11, |
| PFPeA | 263,165 | 263,1642 | 11,24 | 2 |
| Radicicol | 363,182 | 363,1806 | 1,63 | 11,13,16,17,14,12,2,1 5 |
| Secobarbital | 236,917 | 236,9155 | 1,05 | 15 |
| Vedaprofen | 281,177 | 281,1766 | 15,15 | 16 |

Table 16. Compounds detected in ESI (+).

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|---|---------------------------|------------------------|-------------|---------------------------|---------|
| 25I-NBOMe | 428,255 | 428,2550 | 14,43 | 12 | 1,97464 |
| 2-Benzyltetronic Acid | 190,912 | 190,9090 | 1,21 | 11,1,6,7,10 | 1,71775 |
| 2C-E | 210,112 | 210,1115 | 1,51 | 12,6,8,7,10 | 1,50137 |
| 2C-P | 241,154 | 241,1559 | 1,48 | 11,16,17,15 | 1,34957 |
| 3,4-Dimethoxyphenethylamine | 182,082 | 182,0819 | 1,46 | 16,15,13 | 1,77064 |
| 4-Beta-hydroxystanozolol | 182,082 | 182,0817 | 1,38 | 15,16,17,11,1,8,3,6,7,10, | 1,07988 |
| 4-Beta-hydroxystanozolol | 345,231 | 345,2308 | 1,40 | 14 | 1,93558 |
| 4-EEC Ethylethcathinone | 206,103 | 206,1024 | 1,60 | 14 | 1,93883 |
| 4-Fluoroamphetamine | 154,142 | 154,1424 | 12,72 | 2 | 1,92382 |
| 4-Fluoromethamphetamine | 168,066 | 168,0628 | 1,39 | 14,16,13,6 | 1,90835 |
| 4-methylnitrosamino-1-3-pyridyl-1-butanol | 210,135 | 210,1357 | 1,41 | 6 | 1,99311 |
| 4-MTA | 182,080 | 182,0818 | 1,52 | 3,6 | 1,82334 |
| 5-Hydroxythiabendazole | 218,211 | 218,2113 | 14,23 | 7 | 1,90379 |
| 5-Methyl-mellein | 207,138 | 207,1382 | 1,43 | 6 | 1,91168 |
| 7-Aminodesmethylflunitrazepam | 270,280 | 270,2797 | 18,96 | 12,4,9 | 1,61323 |
| 8-Hydroxyquinoline | 146,060 | 146,0598 | 10,34 | 1,10 | 1,80248 |
| 9-Hydroxyrisperidone | 427,340 | 427,3401 | 19,31 | 15,16,13,11,3,10,8 | 1,23423 |
| Acecarbromal | 279,233 | 279,2328 | 1,80 | 14 | 1,90548 |
| Aceclidine | 170,154 | 170,1539 | 13,32 | 4,9 | 1,86929 |
| Acetamiprid | 223,064 | 223,0742 | 7,90 | 4 | 1,96346 |

| Library Detection | Theoretical m/z | Detected m/z | RT (min) | Samples | VIP |
|--------------------------------|--------------------|-----------------|-------------|----------------------------------|---------|
| Acetochlor | 270,126 | 270,1262 | 11,1 | 12,4,9 | 1,62789 |
| Actinoquinol | 254,154 | 254,1542 | 18,03 | 15 | 1,93761 |
| Aflatoxin B2 | 315,196 | 315,1986 | 17,52 | 16,9 | 1,97832 |
| Ajmaline | 327,079 | 327,0794 | 17,82 | 13,15,16 | 1,80012 |
| Albendazole-D3 | 269,248 | 269,2494 | 1,83 | 2,3,5,7,8,10 | 1,45079 |
| Albendazolsulfonamin | 240,139 | 240,1397 | 1,89 | 15 | 1,92363 |
| Aldicarb-sulfoxide | 207,159 | 207,1591 | 14,24 | 10 | 1,95348 |
| Alpha-PPP | 204,087 | 204,0812 | 1,89 | 8 | 1,93077 |
| Alpha-Pyrrolidinopentiophenone | 232,170 | 232,1743 | 1,56 | 12,4,9,10 | 1,53898 |
| Alverine | 282,093 | 282,0956 | 16,13 | 12,3 | 1,81898 |
| Aminoflubendazol | 256,263 | 256,2632 | 1,79 | 16 | 1,96888 |
| Amisulpride | 370,217 | 370,2169 | 18,70 | 5 | 1,95816 |
| AMT | 174,992 | 174,9924 | 1,42 | 13,16,14,17,11,12,1,6,8,10,2,3,9 | 1,10455 |
| a-Nortestosteron | 275,149 | 275,1436 | 9,13 | 8 | 1,92023 |
| Apophedrin | 170,081 | 170,0821 | 1,51 | 15,16 | 1,80837 |
| Aramite | 352,307 | 352,3063 | 1,78 | 17,8 | 1,86342 |
| Aspinolide B | 285,222 | 285,2217 | 19,87 | 12 | 1,99593 |
| Atenolol | 267,171 | 267,1707 | 2,90 | 17 | 1,91923 |
| Atratone | 212,201 | 212,2009 | 17,11 | 12 | 1,91676 |
| Atrazine | 216,086 | 216,1007 | 9,95 | 4,9,12 | 1,65839 |
| Atrazine-2-hydroxy | 198,185 | 198,1845 | 16,60 | 12 | 1,99523 |
| Atrazine-desethyl | 188,070 | 188,0698 | 13,02 | 12 | 1,97218 |
| Atrazine-desisopropyl | 174,055 | 174,0550 | 6,75 | 8,6,5,7 | 1,62605 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|----------------------------|---------------------------|------------------------|-------------|--------------------|---------|
| Austdiol | 237,112 | 237,1122 | 1,52 | 13,15 | 1,81243 |
| Axeen | 227,128 | 227,1281 | 1,50 | 11 | 1,99334 |
| Azaperol | 330,210 | 330,2105 | 1,69 | 9,12,4 | 1,76984 |
| Beflubutamid | 356,196 | 356,1954 | 13,29 | 9,12,9,4 | 1,68854 |
| Benazolin | 244,090 | 244,0907 | 14,37 | 15,1 | 1,84029 |
| Benzocetamine | 250,087 | 250,0874 | 1,87 | 12 | 1,95955 |
| Benzoximate | 364,325 | 364,3136 | 18,40 | 12 | 1,92986 |
| Benzylpiperazine BZP | 177,054 | 177,0541 | 15,99 | 3,6,7 | 1,6353 |
| Betamethasone-21-phosphate | 473,320 | 473,3265 | 1,72 | 4 | 1,91477 |
| Bifenazate | 323,146 | 323,1347 | 1,42 | 7,10 | 1,86396 |
| Bioallethrin | 303,231 | 303,2312 | 1,80 | 17,8 | 1,82974 |
| Biotin | 245,227 | 245,2267 | 1,80 | 17,3 | 1,9933 |
| Buprenorphine | 468,325 | 468,3256 | 13,54 | 10,8,1,10,3,5,4,2 | 1,00012 |
| Butylate | 218,211 | 218,2113 | 14,27 | 14,17,4 | 1,76627 |
| Butylone | 222,113 | 222,1186 | 13,39 | 17,13,14,11,12,4,9 | 1,20488 |
| Caffeine | 195,087 | 195,0870 | 11,39 | 12 | 1,97385 |
| Cannabinol | 311,164 | 311,1641 | 17,82 | 2 | 1,9719 |
| Capsaicin | 306,291 | 306,299 | 17,84 | 12,4,9 | 1,63193 |
| Carbamazepine | 237,103 | 237,1025 | 15,22 | 13,12 | 1,86959 |
| Carboxin | 236,107 | 236,1070 | 16,35 | 13,14,16 | 1,75915 |
| Cathine | 169,097 | 169,0986 | 1,49 | 14,3,5,9 | 1,74871 |
| Cathinone | 150,027 | 150,0262 | 19,77 | 6 | 1,94434 |
| CBD | 315,195 | 315,1954 | 1,45 | 1,14,17,11,4,9 | 1,56184 |
| Cerulenin | 224,118 | 224,1271 | 10,92 | 16,4 | 1,84696 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|----------------------|---------------------------|------------------------|-------------|-------------------------|---------|
| Chanoclavine | 279,249 | 279,2491 | 19,73 | 1 | 1,96412 |
| Chloroquine | 320,288 | 320,2886 | 17,67 | 1 | 1,91292 |
| Chlorphenethiazine | 305,175 | 305,1788 | 1,45 | 1 | 1,93923 |
| Cimetidine | 253,216 | 253,2163 | 1,57 | 1 | 1,99649 |
| Citreoviridin A | 403,234 | 403,2343 | 1,68 | 17,11,3 | 1,76125 |
| Citrinin | 251,164 | 251,1642 | 14,98 | 12 | 1,95085 |
| Clemastine | 344,196 | 344,1950 | 11,96 | 12 | 1,94401 |
| Clibucaine | 337,179 | 337,1814 | 11,69 | 12 | 1,9190 |
| Climbazol | 293,106 | 293,1058 | 17,02 | 12 | 1,93284 |
| Clobazam | 301,142 | 301,1419 | 1,60 | 12 | 1,91014 |
| Clobendazole | 330,137 | 330,1373 | 14,67 | 9 | 1,90776 |
| Clodinafop-propargyl | 350,248 | 350,2482 | 19,41 | 4 | 1,97762 |
| Clofentezine | 303,163 | 303,1561 | 1,56 | 17 | 1,92601 |
| Clomipramine | 315,161 | 315,1614 | 16,54 | 13,16,11 | 1,7209 |
| Clonidine | 230,154 | 230,1541 | 17,90 | 15,12,8 | 1,73321 |
| Corticosterone | 347,184 | 347,1845 | 1,41 | 17,10,12,4,9 | 1,42151 |
| Cotinine | 177,102 | 177,1022 | 10,73 | 10 | 1,91386 |
| Crotetamide | 227,175 | 227,1753 | 13,74 | 14,17,15,11,1,2,4,5,6,7 | 1,22091 |
| Curvularin | 291,197 | 291,1963 | 1,37 | 12 | 1,91708 |
| Cyclizine | 267,174 | 267,1742 | 17,30 | 17 | 1,9363 |
| Cyclobenzaprine | 276,145 | 276,1443 | 1,51 | 2 | 1,90127 |
| Cyclovalone | 367,320 | 367,3207 | 1,26 | 3 | 1,91846 |
| Cyhalofop-butyl | 392,374 | 392,3741 | 19,73 | 2,4,9 | 1,81147 |
| Cyprodinil | 226,217 | 226,2173 | 1,75 | 4,5 | 1,89571 |

| Library Detection | Theoretical m/z | Detected m/z | RT (min) | Samples | VIP |
|------------------------------------|--------------------|-----------------|-------------|--|---------|
| Cytochalasin E | 496,449 | 496,4436 | 19,92 | 17,11,12,4,9 | 1,55721 |
| DEET | 192,139 | 192,1384 | 15,93 | 13 | 1,91091 |
| Delorazepam | 337,237 | 337,2371 | 17,06 | 6 | 1,94854 |
| Demeton-O | 291,196 | 291,1963 | 18,74 | 1,2,10,8 | 1,70727 |
| Deoxynivalenol-3-glucoside | 459,306 | 459,3062 | 1,51 | 12 | 1,94334 |
| Desmethylcitalopram | 311,146 | 311,1548 | 14,47 | 5 | 1,94378 |
| Desmethyl-formamido- pirimicarb | 253,159 | 253,1621 | 18,56 | 12 | 1,92795 |
| Detomidine | 187,096 | 187,0962 | 8,41 | 13,14,15,16,17,11,1,2,3,4,5,6,7,8,9,10 | 1,03726 |
| Dexamethasone | 393,226 | 393,2246 | 16,93 | 11 | 1,96466 |
| Diacetylmorphine | 370,368 | 370,3687 | 18,77 | 7 | 1,92043 |
| Dibutyl phthalate sodiated | 301,141 | 301,1415 | 18,26 | 13,17,14,12,1,2,6,7,8 | 1,55739 |
| Dienestrol-D2 | 269,175 | 269,1746 | 14,15 | 5,7,4 | 1,71973 |
| Dihydrolysergol | 257,138 | 257,1414 | 1,50 | 12 | 1,90485 |
| Dilazep | 605,145 | 605,1452 | 11,31 | 12 | 1,90725 |
| Dimethomorph | 388,132 | 388,1316 | 16,81 | 17 | 1,97567 |
| Dioxacarb | 224,118 | 224,1129 | 11,18 | 17,15,16,11,12 | 1,55673 |
| Dipyridamole | 505,334 | 505,3340 | 1,40 | 12 | 1,90718 |
| Diuron | 233,025 | 233,0245 | 9,98 | 9 | 1,90512 |
| DMT | 189,128 | 189,1276 | 15,63 | 15 | 1,93161 |
| D-Norpseudoephedrine | 169,097 | 169,0985 | 1,50 | 14 | 1,90875 |
| Dodemorph | 282,205 | 282,2076 | 10,76 | 16 | 1,91828 |
| DOEt | 224,128 | 224,1282 | 1,50 | 15 | 1,96997 |
| DOM | 210,135 | 210,1355 | 1,46 | 11,10 | 1,92882 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|----------------------|---------------------------|------------------------|-------------|--------------------------------|---------|
| Doramectin | 921,690 | 921,6905 | 19,24 | 7 | 1,96182 |
| Doxepin | 280,199 | 280,1988 | 1,40 | 7 | 1,99683 |
| Emamectin B1b | 872,704 | 872,7008 | 1,45 | 11,1,3 | 1,79217 |
| EMDP | 264,196 | 264,2006 | 1,64 | 2 | 1,94601 |
| Ephedrine | 166,086 | 166,0866 | 1,38 | 13 | 1,95787 |
| Epioxandrolone | 329,175 | 329,1757 | 1,36 | 13,14,15,16,17,1,6,8,2,3,7,10 | 1,2253 |
| Eprinomectin B1a | 936,686 | 936,6846 | 19,80 | 12,9 | 1,89869 |
| Eprosartan | 425,215 | 425,2162 | 18,76 | 13 | 1,97473 |
| Erginine | 268,263 | 268,2652 | 19,58 | 1,4 | 1,91029 |
| Ethambutol | 237,149 | 237,1489 | 11,31 | 17 | 1,96684 |
| Ethiofencarb-sulfone | 258,182 | 258,1849 | 1,53 | 16 | 1,9927 |
| Ethofumesate | 287,270 | 287,2694 | 1,48 | 2 | 1,98802 |
| Ethylone | 222,149 | 222,1489 | 1,47 | 16,1 | 2,00484 |
| Etilefrine | 182,082 | 182,0812 | 1,33 | 13,1 | 1,94503 |
| Etorphine | 412,349 | 412,3488 | 13,61 | 2 | 1,93064 |
| fenoxy carb | 302,246 | 302,2322 | 17,42 | 17,10 | 1,96871 |
| Fenproporex | 189,075 | 189,0743 | 1,43 | 2,12,5 | 1,85652 |
| Flecainide | 415,145 | 415,1483 | 14,62 | 9 | 1,97004 |
| Flubendazole | 314,269 | 314,2696 | 1,52 | 2,7 | 1,94074 |
| Flumethasone | 411,215 | 411,2221 | 11,72 | 12,4,11,16 | 1,62781 |
| Foramsulfuron | 453,344 | 453,3445 | 2,12 | 15 | 1,95191 |
| Fumonisin B1 | 722,507 | 722,5071 | 1,87 | 8,3,11 | 1,73928 |
| Furosemide | 331,001 | 331,0024 | 1,40 | 8 | 1,92939 |
| Fusaproliferin | 445,284 | 445,2837 | 1,50 | 1,2,3,4,5,6,7,8,10,13,16,14,15 | 0,90517 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|-----------------------|---------------------------|------------------------|-------------|----------------------------------|---------|
| Gabapentin | 172,133 | 172,1332 | 1,58 | 9,6,1,16,5,7,12,3,8 | 1,09125 |
| Gallopamil | 485,382 | 485,3818 | 19,55 | 3 | 1,9310 |
| Glufosinate | 182,080 | 182,0814 | 1,68 | 7 | 1,99065 |
| Glutethimide | 218,211 | 218,2110 | 1,48 | 8,5,10,1 | 1,74446 |
| Glycopyrrolate | 318,240 | 318,2405 | 1,33 | 8 | 1,94281 |
| Haloxypop-etotyl | 434,267 | 434,2493 | 18,77 | 2 | 1,96252 |
| Helvolic acid | 569,435 | 569,4349 | 1,68 | 8 | 1,91681 |
| Hexaconazole | 314,083 | 314,0828 | 17,70 | 12 | 1,92966 |
| Hexazinone | 253,166 | 253,1659 | 9,38 | 11 | 1,90187 |
| Histamine | 112,050 | 112,0502 | 1,51 | 9 | 1,98965 |
| Histidine | 156,102 | 156,1019 | 1,50 | 3,7 | 1,94454 |
| Hordenine | 166,123 | 166,1226 | 1,47 | 3,14,7 | 1,80238 |
| HT-2 Toxin | 447,294 | 447,2944 | 1,41 | 3,7,8,2,15,6,11,10,17,5,16,1 | 1,11109 |
| Hydralazine | 161,071 | 161,0712 | 11,45 | 4,9 | 1,95659 |
| Hydroxychloroquine | 336,252 | 336,2515 | 1,42 | 8 | 1,98085 |
| Hydroxymethylpyridine | 110,060 | 110,0600 | 1,91 | 9 | 1,92264 |
| Inabenfide | 356,315 | 356,3158 | 18,73 | 12 | 1,97593 |
| Infectopyrone | 265,144 | 265,1443 | 1,47 | 16,15,10,13,17,6,3,11,14,2,5,7,8 | 1,0226 |
| Ipconazole | 334,204 | 334,2039 | 18,84 | 12 | 1,95642 |
| Irbesartan | 429,241 | 429,2414 | 16,50 | 12 | 1,97948 |
| isoprocarb | 211,133 | 211,1331 | 11,31 | 14,11,9,4,3 | 1,55031 |
| Isoproturon | 207,149 | 207,1490 | 10,02 | 4,9 | 1,92176 |
| Isothipendyl | 286,238 | 286,2373 | 18,12 | 12 | 1,9925 |
| Josamycin | 828,677 | 828,6765 | 1,40 | 6 | 1,9629 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|-----------------------------|---------------------------|------------------------|-------------|---------------------------------|--------|
| Ketorolac | 256,133 | 256,1333 | 15,01 | 9,4 | 1,8889 |
| Ketotifen | 310,311 | 310,3112 | 19,65 | 15,14,8,10 | 1,5185 |
| Kresoxim-methyl | 314,159 | 314,1616 | 10,82 | 4 | 1,9958 |
| Lenacil | 235,144 | 235,1448 | 10,09 | 14,10 | 1,9853 |
| Levamisole | 205,105 | 205,1051 | 1,65 | 5,12,6,13,15,14,16 | 1,0289 |
| Levocabastine | 438,379 | 438,3792 | 18,09 | 5 | 1,9085 |
| Levodopa | 198,185 | 198,1852 | 1,75 | 10,12,13,16,15,6,8 | 1,1777 |
| Lidocaine | 235,181 | 235,1809 | 11,54 | 12 | 1,9303 |
| Lincomycin | 407,222 | 407,2221 | 10,68 | 12 | 1,9016 |
| Lonazolac | 313,274 | 313,2750 | 12,98 | 14 | 1,9122 |
| Lorazepam | 321,169 | 321,1684 | 14,88 | 12,15,17 | 1,6405 |
| Lysergol | 255,175 | 255,1770 | 17,55 | 3,17 | 1,8006 |
| Malaoxon | 315,301 | 315,3013 | 18,88 | 5 | 1,9014 |
| Maleic hydrazide | 113,107 | 113,1069 | 2,08 | 11,7,12,3 | 1,7003 |
| Maprotiline | 278,190 | 278,1902 | 1,40 | 7,8,4,3,1,5,2 | 1,3077 |
| Mazindol | 285,076 | 285,0763 | 17,26 | 9,12,13 | 1,7039 |
| Mebeverine | 430,151 | 430,1512 | 1,83 | 1 | 1,9667 |
| Medetomidine | 201,107 | 201,1131 | 11,71 | 1 | 1,9453 |
| Medroxyprogesterone acetate | 387,285 | 387,2842 | 1,55 | 8,1,3,6,16,13,15,7,10,8,11 | 1,2471 |
| Mefenamic acid | 242,144 | 242,1535 | 17,70 | 3 | 1,9852 |
| Mepenzolate | 340,246 | 340,2472 | 18,86 | 2,11 | 1,9140 |
| Mesoridazine | 387,199 | 387,1989 | 14,61 | 5,2,12 | 1,8480 |
| Metaxalone | 222,149 | 222,1488 | 1,45 | 11,8,5,7,3,17,15,1,10,6,16,2,12 | 1,3399 |
| Metformin | 130,109 | 130,1083 | 1,39 | 4,9 | 1,9707 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|---------------------|---------------------------|------------------------|-------------|--------------------------------|--------|
| Methacrifos | 241,217 | 241,2168 | 1,35 | 4,9 | 1,9025 |
| Methamphetamine | 150,128 | 150,1372 | 19,89 | 9 | 1,9186 |
| Methaqualone | 251,126 | 251,1263 | 11,34 | 4 | 1,9057 |
| Methazolamide | 237,148 | 237,1481 | 14,30 | 9,14,11 | 1,8009 |
| Methfuroxam | 230,175 | 230,1750 | 14,28 | 13,15 | 1,8273 |
| Methocarbamol | 242,154 | 242,1441 | 1,45 | 1,2,6,7,8,10,11,12,13,14,15,16 | 1,2046 |
| Methoprotrolyne | 272,259 | 272,2590 | 1,76 | 1 | 1,9636 |
| Methyl Prednisolone | 375,217 | 375,2167 | 1,39 | 15 | 1,9348 |
| Methylprednisolone | 375,347 | 375,3471 | 19,81 | 10,13,9,12,16,14 | 1,5102 |
| Methyltestosterone | 303,180 | 303,1795 | 1,51 | 8,5 | 1,9098 |
| Metolachlor | 284,141 | 284,1412 | 11,20 | 11,12,9,4 | 1,7763 |
| Metolcarb | 166,123 | 166,1226 | 1,51 | 7,2,15,1,5,13,12,11 | 1,1001 |
| Mexacarbate | 245,079 | 245,0790 | 16,09 | 2 | 1,9079 |
| Minoxidil | 210,135 | 210,1349 | 12,70 | 14 | 1,9373 |
| Mitragynine | 399,309 | 399,3088 | 12,84 | 5 | 1,9800 |
| Molinate | 188,111 | 188,1109 | 10,86 | 8 | 1,9878 |
| Molsidomine | 243,122 | 243,1224 | 11,50 | 14 | 1,9952 |
| Monocerin | 309,134 | 309,1341 | 15,91 | 12 | 1,9435 |
| Monuron | 199,169 | 199,1699 | 12,10 | 10,3 | 1,8895 |
| Moxisylyte | 280,164 | 280,1636 | 8,40 | 17,10,14 | 1,7851 |
| Moxonidine | 242,248 | 242,2482 | 19,41 | 9,10,8 | 1,7069 |
| Nandrolon | 275,166 | 275,1661 | 1,53 | 6,15,4,7,13,16,5,3,2,8,14 | 1,1203 |
| Nandrolone | 275,201 | 275,2014 | 1,45 | 5,10,16,11,3,8,1,10 | 1,3110 |
| Naphazoline | 211,149 | 211,1479 | 18,04 | 9,12 | 1,8014 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|-------------------------|---------------------------|------------------------|-------------|---|--------|
| Naproxen | 230,891 | 230,8904 | 2,28 | 5 | 1,9993 |
| Neburon | 275,147 | 275,1465 | 10,88 | 4,9 | 1,9349 |
| N-Ethyl Hexedrone Hexen | 220,096 | 220,0969 | 1,49 | 7,3,1 | 1,7474 |
| N-Ethyl Pentylone | 250,146 | 250,1486 | 1,34 | 13,16 | 1,8112 |
| Nicotinamide | 123,055 | 123,0546 | 3,03 | 4,9,17 | 1,8109 |
| Nicotine | 163,133 | 163,1337 | 11,56 | 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17 | 1,0086 |
| Niflumic acid | 283,153 | 283,1528 | 15,54 | 17 | 1,9976 |
| N-Isopropylsalicylamide | 180,102 | 180,1020 | 2,01 | 11,17,1,2,8,3,4,9,7 | 1,0084 |
| N-methyltryptamine NMT | 175,123 | 175,1228 | 9,79 | 17 | 1,9254 |
| Nonivamide | 294,207 | 294,2068 | 1,47 | 10 | 1,9154 |
| Norbuprenorphine | 414,270 | 414,2724 | 19,79 | 11 | 1,9048 |
| Norcotinine | 163,039 | 163,0392 | 14,33 | 14,17 | 1,9547 |
| Norephedrine | 194,098 | 194,0918 | 10,88 | 13 | 1,9705 |
| Norethisterone | 299,220 | 299,2199 | 1,39 | 15,11 | 1,9498 |
| Norethisterone acetate | 358,241 | 358,2413 | 1,53 | 5,8,7,16,6,3,10,17,11,3,12 | 1,0076 |
| Norgesterel | 313,236 | 313,2384 | 12,07 | 13,12,10,15,9 | 1,5191 |
| Norhydrocodone | 286,311 | 286,3088 | 18,21 | 9 | 1,9062 |
| Norsertaline | 292,227 | 292,2203 | 19,63 | 12 | 1,9105 |
| Noscapine | 414,155 | 414,1547 | 1,48 | 2,3,7,1,4,5,12 | 1,1552 |
| N-propylamphetamine | 178,058 | 178,0581 | 16,07 | 9,12 | 1,9081 |
| Obidoxime | 144,066 | 144,0731 | 1,49 | 3 | 1,9035 |
| Ochratoxin alpha | 295,095 | 295,0948 | 12,39 | 17,12,9,4 | 1,6196 |
| Omethoate | 214,125 | 214,1256 | 1,53 | 11 | 1,9685 |
| Ophiobolin A | 401,123 | 401,1214 | 14,21 | 17,3,11 | 1,7438 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|-------------------------|---------------------------|------------------------|-------------|--|--------|
| Oxadixyl | 279,194 | 279,1944 | 15,96 | 9,7 | 1,8347 |
| Oxamyl | 237,149 | 237,1486 | 14,92 | 17 | 1,9087 |
| Oxitriptan | 221,118 | 221,1177 | 1,42 | 16,13,15,2 | 1,8403 |
| Oxydemeton-methyl | 247,133 | 247,1334 | 14,27 | 17 | 1,9534 |
| Oxymorphone | 305,175 | 305,1751 | 14,90 | 14 | 1,9030 |
| Oxypendyl | 371,102 | 371,103 | 7,45 | 6 | 1,9543 |
| Oxyphencyclimine | 345,337 | 345,3364 | 2,23 | 11 | 1,9727 |
| Pacllobutrazol | 294,243 | 294,2428 | 12,53 | 17 | 1,9268 |
| Palitantin | 287,147 | 287,1477 | 11,20 | 2,3,5,6,7,8,10,11,12,13,14,15,16,17 | 1,0079 |
| Papaverine | 340,264 | 340,2645 | 18,86 | 3,5 | 1,9169 |
| Para-Methoxyamphetamine | 166,086 | 166,0851 | 1,40 | 14,7 | 1,8214 |
| Paraoxon-methyl | 248,149 | 248,1483 | 14,14 | 12 | 1,9198 |
| Penconazole | 284,072 | 284,0719 | 11,35 | 3,1 | 1,9382 |
| Penicillic acid | 171,101 | 171,1014 | 12,66 | 9,7,10,1,2,5,6,3,12,8,11 | 1,1704 |
| Pentedrone | 209,165 | 209,1648 | 1,47 | 17,16,11,10,7,15 | 1,0552 |
| Pentylene-tetrazole | 156,102 | 156,1015 | 1,51 | 5,6,15,3,16,8,10,17,2,7 | 1,1770 |
| Perazine | 340,264 | 340,2641 | 18,95 | 7,13 | 1,9225 |
| Pestalotin | 215,093 | 215,0931 | 1,85 | 1,2,4,5,6,7,8,10,11,12,13,14,15,16,17, | 1,0487 |
| Phenelzine | 137,096 | 137,0961 | 1,51 | 1,2,3,5,6,7,8,10,11,12,15,16,17 | 1,0238 |
| Pheniramine | 241,144 | 241,1439 | 14,71 | 17 | 1,9983 |
| Phenmedipham | 301,141 | 301,1427 | 12,30 | 12 | 1,9379 |
| Phentermine | 150,027 | 150,0269 | 1,61 | 13,15,6,11,7,16,8,5,3,10,12,2 | 1,0836 |
| Phenylephrine | 168,063 | 168,0645 | 1,59 | 16,8,12,6,5,7,1,11,2,4,3,17,6 | 1,1054 |
| Picoxystrobin | 368,241 | 368,2415 | 17,34 | 12,4,9 | 1,0757 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|--------------------------------------|---------------------------|------------------------|-------------|--------------------------------------|--------|
| Piperacetazine | 411,346 | 411,3453 | 11,20 | 3,8,6,5,2,10,7 | 1,0091 |
| Piperonyl-butoxide | 356,244 | 356,2437 | 18,68 | 12 | 1,9291 |
| Pirimiphos-methyl | 306,243 | 306,2429 | 14,63 | 17 | 1,9104 |
| Pizotifen | 296,295 | 296,2954 | 19,58 | 9,17 | 1,8254 |
| PMMA | 180,16 | 180,1603 | 18,21 | 14,17,16,15,12,1 | 1,3604 |
| Prenylamine | 330,338 | 330,3373 | 1,82 | 1,8,7,6,10,5 | 1,1116 |
| Progesterone | 347,298 | 347,2933 | 19,15 | 7,1,3 | 1,6392 |
| Prohexadione | 213,112 | 213,1123 | 1,49 | 13,12,8,7,10,15,5,6,1,11 | 1,4528 |
| Prometon | 226,142 | 226,1433 | 1,44 | 15 | 1,9330 |
| Propoxycarbazone | 438,379 | 438,3794 | 18,02 | 8 | 1,9121 |
| Prosulfocarb | 252,046 | 252,0459 | 1,26 | 13,14,15,17,11,12,10,1,2,3,4,5,6,7,8 | 1,0698 |
| Prothioconazole Desthiometabolite | 312,327 | 312,3263 | 5,39 | 14 | 1,9709 |
| Pymetrozine | 218,211 | 218,2111 | 14,18 | 9 | 1,9036 |
| Pyrenocine A | 209,129 | 209,1289 | 1,50 | 3 | 1,9492 |
| Pyridoxine | 170,117 | 170,1173 | 1,50 | 14 | 1,9967 |
| Pyrifenox | 295,153 | 295,1606 | 11,98 | 14 | 1,9094 |
| Pyrilamine | 286,238 | 286,2368 | 18,25 | 1 | 1,9001 |
| Pyrimethamine | 249,185 | 249,1854 | 1,76 | 16,17 | 1,8831 |
| Pyrimethanil | 200,118 | 200,1182 | 10,48 | 7 | 1,9449 |
| Pyrvinium | 383,314 | 383,3137 | 19,44 | 6 | 1,9043 |
| RCS-8 | 376,260 | 376,2602 | 1,37 | 14 | 1,9549 |
| Selegiline | 188,128 | 188,1282 | 1,79 | 14 | 1,9540 |
| Serotonin | 177,092 | 177,0919 | 1,51 | 11 | 1,9288 |

| Library Detection | Theoretical m/z | Detected m/z | RT (min) | Samples | VIP |
|--------------------------|--------------------|-----------------|-------------|----------------------------------|--------|
| Simazine-2-hydroxy | 184,169 | 184,1697 | 14,68 | 13,14,16,6,2,11,10,5,8,3,12,9,17 | 1,0447 |
| Spinetoram A | 748,542 | 748,5414 | 18,72 | 12,6 | 1,8430 |
| Spiromesifen | 371,102 | 371,1023 | 7,28 | 4,12 | 1,8459 |
| Stanozolol | 329,157 | 329,1572 | 12,58 | 2 | 1,9408 |
| Strychnine | 335,222 | 335,2179 | 17,44 | 8 | 1,9466 |
| Sulfadiazine | 251,060 | 251,0603 | 9,18 | 12,2 | 1,9018 |
| T-2 triol | 383,208 | 383,2076 | 1,49 | 12,4 | 1,8392 |
| Tapentadol | 222,186 | 222,1856 | 12,77 | 12 | 1,9980 |
| Tebuconazole | 308,153 | 308,1527 | 17,60 | 16,17,15,11 | 1,7711 |
| Tebuthiuron | 229,108 | 229,1083 | 16,30 | 12 | 1,9056 |
| Telmisartan | 515,244 | 515,2436 | 17,60 | 9,12 | 1,9416 |
| Temazepam | 301,217 | 301,2168 | 19,06 | 12,9,4 | 1,8437 |
| Terbumeton | 225,936 | 225,9364 | 2,27 | 12,4,9,10,2,5,11,7,6 | 1,1006 |
| Terbuthylazine | 230,117 | 230,1165 | 16,69 | 4,6 | 1,9204 |
| Terbuthylazine-2-hydroxy | 212,151 | 212,1508 | 14,07 | 4,9,13 | 1,8557 |
| Terbuthylazine-desethyl | 202,086 | 202,0855 | 15,06 | 12,9 | 1,9765 |
| Terbutryn | 242,144 | 242,1439 | 17,27 | 12 | 1,9442 |
| Theobromine | 181,072 | 181,0720 | 8,67 | 12,9,4 | 1,8813 |
| Theophylline | 181,072 | 181,0717 | 9,77 | 12,9,17,9,14,4 | 1,7538 |
| Thiabendazole | 202,181 | 202,1806 | 1,49 | 12,7 | 1,8655 |
| Thiacloprid | 253,031 | 253,0310 | 8,39 | 12,9 | 1,7944 |
| Thionazin | 248,872 | 248,8716 | 1,24 | 9 | 1,9379 |
| Thioproperazine | 447,294 | 447,2932 | 1,43 | 17,2 | 1,9158 |
| Thioridazine | 371,290 | 371,2912 | 17,98 | 1 | 1,9759 |

| Library Detection | Theoretical <i>m/z</i> | Detected <i>m/z</i> | RT (min) | Samples | VIP |
|-------------------|---------------------------|------------------------|-------------|------------------------------|--------|
| Thymopentin | 680,516 | 680,5163 | 1,52 | 5,16,15,14,13,17,6 | 1,4462 |
| Tiocarlide | 401,287 | 401,2877 | 1,78 | 9,5,16 | 1,7633 |
| Tocainide | 193,072 | 193,0723 | 9,44 | 5 | 1,9626 |
| Tramadol | 264,196 | 264,1964 | 12,09 | 1,2,6,3,11,8,7,16,11,15,9,12 | 1,0015 |
| Tranexamic acid | 158,027 | 158,0269 | 1,56 | 9 | 1,9701 |
| Triamcinolone | 412,218 | 412,2193 | 12,47 | 12 | 1,9960 |
| Triamterene | 254,248 | 254,2478 | 18,12 | 8 | 1,9043 |
| Triazoxide | 248,237 | 248,2377 | 18,94 | 9,15,2,13,3,8,10,17 | 1,0539 |
| Triflupromazine | 353,304 | 353,3033 | 19,81 | 1,10,9,5,7,2,3,12,6,8 | 1,0236 |
| Trihexyphenidyl | 302,246 | 302,2455 | 5,79 | 2,3,5,7,8,10 | 1,0914 |
| Tritoqualine | 501,377 | 501,3772 | 18,69 | 2,13,3,10,5 | 1,0211 |
| Tryptamine | 161,096 | 161,0962 | 1,39 | 2 | 1,9944 |
| Tryptophol | 162,076 | 162,0763 | 1,47 | 11,2,7,13,14,15,16,17 | 1,0001 |
| Tylosin A | 916,733 | 916,7327 | 3,41 | 13,14,16 | 1,7227 |
| Valsartan | 436,234 | 436,2347 | 16,25 | 3 | 1,9809 |
| Vardenafil | 489,359 | 489,3590 | 1,76 | 3 | 1,9972 |
| Verapamil | 455,372 | 455,3719 | 3,11 | 8 | 1,9150 |
| Vincamine | 355,012 | 355,0129 | 1,13 | 12 | 1,9717 |
| Xylazine | 221,154 | 221,1538 | 17,34 | 13,14,15,16,17,11 | 1,3044 |
| Xylometazoline | 245,247 | 245,2268 | 19,50 | 1,7,12 | 1,7033 |
| Zinniol | 267,123 | 267,1211 | 16,77 | 10 | 1,9218 |
| Ziprasidone | 413,267 | 413,2670 | 19,87 | 12 | 1,9806 |

3. Conclusions

In this chapter, suspect and non-target screening analyses were combined in order to obtain a more holistic view of the quality of European water bodies. Analyses were carried out with HRMS due to its high sensitivity especially at low concentration levels. Pollution patterns among samples originating from different surface water sources were identified using multivariate statistical analysis tools. PCA models were used for their identification among the samples while PLS-DA models were used in order to show which features (micropollutants) were more important for the discrimination between the water samples. Identification of the compounds showed the wide range of compounds present in water sources, indicating the high occurrence of pharmaceuticals, pesticides and PFAS. The results of this study will be implemented in the NORMAN database, and can make an important contribution in decision making for more target monitoring assessments – in order to evaluate the pollution levels and plan the appropriate treatment activities, as well as for prioritization of compounds for policy making.

Section IV

Degradation Processes

Chapter 6 Degradation of glyphosate

Firstly, the degradation rate of glyphosate after direct photolysis with UV irradiation was examined. The major aim of this study was to determine the effectiveness of UV irradiation in decreasing the toxicity of glyphosate and identify the relevant UV exposure regimes (UV wavelength, UV dose). A test battery including different aquatic organisms was used in order to observe changes in activity and growth of the target organisms before and after UV irradiation. The test organisms were selected from different trophic levels for better assessing the biological effects of all the bioactive compounds in the samples after UV treatment including glyphosate's degradation byproducts. Suspect-screening analyses of the samples before and after treatment with LC-HRMS provided valuable information for understanding the degradation mechanism of glyphosate.

1. Materials and Methods

1.1 Chemicals and materials

Stock standards of *N*-(phosphonomethyl)glycine, monoisopropylamine salt solution (CAS 38641-94-0), and *N*-(phosphonomethyl)glycine (CAS 1071-83-6) were purchased from Sigma-Aldrich (Saint Louis, MO, USA). For the irradiation a 4W UVP UVGL-25 lamp equipped with separate tubes for UV-A (365nm) and UV-C (254nm) (Analytic Jena US), and an 8W UVP 3UV™ lamp equipped with separate tubes for UV-A (365nm), UV-B (302 nm) and UV-C (254nm) were used. Some initial experiments with the two UV lamps gave comparable results for comparable UV doses and in order to shorten exposure times the 8W lamp with greater intensity was selected to continue with. The irradiation intensity at 15 cm distance from the UVP 3UV™ lamp was 970 $\mu\text{W}/\text{cm}^2/\text{sec}$ for UV-A, 1900 $\mu\text{W}/\text{cm}^2/\text{sec}$ for UV-B, and 327 $\mu\text{W}/\text{cm}^2/\text{sec}$ for UV-C, and it was measured with an Extech SDL470 Light meter equipped with UV-AB and UV-C sensors.

1.2 Sample preparation and treatment

Stock solutions of glyphosate were prepared in both autoclaved distilled deionized water and real water matrices, using the purchased stock standards and the commercial herbicide solutions, at a final concentration of 1 g/L. All stock solutions were stored in the dark at 5°C. As real water matrices, drinking water samples from Aalborg, Aarhus, Skagen, and Sønderborg municipalities in Denmark were used. Moreover, samples from the influent (raw water) and effluent (treated water) of a drinking water treatment plant in Elsted, Denmark were included in this study as well. All real water samples originated from groundwater sources, and regional differences in organic and inorganic

constituents among them were observed, based on the information found at the national Danish well database for water quality <https://eng.geus.dk/products-services-facilities/data-and-maps/national-well-database-jupiter>:

The following parameters were tested during the experiments in order to examine the photodegradation and biotoxicity of glyphosate under UV irradiation:

- a) effect of UV wavelength (UV-A, UV-B, UV-C);
- b) effect of UV dose (range between 2,3 and 70 J/cm²);
- c) effect of glyphosate's concentrations (range between 0,18 and 100 mg/L);
- d) importance of water matrices.

More specifically, solutions of glyphosate and glyphosate-based herbicides with concentrations of 50 and 100 mg/L were prepared in different water matrices. These samples were exposed to different doses of UV-A, UV-B and UV-C irradiation (aim a) using quartz cuvettes (10 mm 3,5 mL, Science Outlet Optical Quartz QS10 and Hellma Precision Quartz SUPRASIL[®] QS10) at room temperature (22°C). Moreover, control samples were included in the study in order to evaluate any toxicity caused from active oxygen species generated during the irradiation experiments and apart from glyphosate. Different UV doses (J/cm²) were calculated from the measured UV irradiation intensity (μW/cm²/sec) and the exposure time (sec) and were achieved by alternating the exposure times and the distance of the solution from the UV lamp (aim b). For example, a UV dose of 20 J/cm² was achieved by using the same exposure time but different distances from the UV lamp for the different wavelengths (20 cm for UV-A, 35 cm for UV-B, and 5 cm for UV-C).

Moreover, diluted glyphosate concentrations were prepared in UV bottom - transparent 96 well microplates (Nunc 96-well UV microplates, Thermo Scientific) and exposed to UV irradiation in order to examine the effects of glyphosate's concentrations on the outcome (aim c). The exposure was done from the top or bottom of the 96-well UV microplates using similar UV doses as the quartz cuvettes, while they were placed on a cooling plate in order to maintain the temperature at around 22°C and avoid evaporation of the small samples' volumes used (100 μL) due to overheating. Control samples without any UV exposure were covered with aluminum foil and stored in the dark. Finally, solutions of glyphosate at a concentration of 100mg/L were prepared in drinking water and exposed to UV irradiation (aim d).

The addition of different oxygen radical probes, helped us to identify the presence of active oxygen species in the aqueous samples after the treatment. Superoxide radicals ($\text{O}_2^{\cdot-}$), were detected by measuring chemiluminescence after post-treatment addition of 1 mM luminol. Similarly, hydroxyl radicals (OH^{\cdot}) were detected by measuring fluorescence after pretreatment addition of 1 mM coumarin, terephthalic acid, and benzoic acid. A Victor X2 Multilabel Plate Reader (Perkin Elmer) was used in order to measure chemiluminescence and fluorescence originating from oxygen radical probes after reactions with active oxygen species.

1.3 Toxicity tests

1.3.1 *Aliivibrio fischeri*

Toxicity screening of glyphosate samples was examined with a standard inhibition test using the luminescent bacterium *Aliivibrio fischeri* (ISO 11348-1, 2009) [172]. *A. fischeri* DSM 7151 was incubated in white 96-well plates (CulturPlates, Perkin Elmer), and exposed to different glyphosate concentrations (0,098, 0,195, 0,390, 0,780, 1,560, 3,130, 6,250, 12,5, 25, 50 mg/L) before and after UV irradiation. Changes in bioluminescence were quantified after 30 minutes of bacteria's exposure to glyphosate, using a Victor X2 Multilabel Plate Reader (Perkin Elmer).

1.3.2 *Bacillus subtilis*

A newly developed inhibition test with *Bacillus subtilis* was used for glyphosate's toxicity on bacteria screening. The endpoint was inhibition of growth and hydrolase activity after 18 h. *Bacillus subtilis* DSM 10 (German Collection of Microorganisms and Cell Cultures) was grown at 30 °C in Davis Minimal Broth (Sigma-Aldrich) supplemented with: 25 μM FeSO₄, 0,5 μM ZnCl₂, 0,5 μM Na₂MoO₄, 0,5 μM MnCl₂, 0,5 μM H₃BO₃, 0,5 μM CoCl₂, 0,5 μM NiCl₂, and 2,0 μM CuSO₄. Serial dilutions of glyphosate were made in 96-well clear microplates (Nunclon, Thermo Scientific) starting from 100 μL glyphosate stock solutions of 100 mg/L and serially diluted in 100 μL autoclaved distilled water resulting in different glyphosate concentrations. After the dilution, 50 μL of 4 x strength Davis Minimal Broth was added to each well, followed by the addition of 50 μL of diluted *B. subtilis* culture (1:1000 dilution in 0,9% NaCl). This resulted in a final sample volume of 200 μL in each well and 10 different concentrations of glyphosate: 0,098, 0,195, 0,390, 0,780, 1,560, 3,130, 6,250, 12,5, 25, 50 mg/L. Four replicates were included for blanks (medium only), controls (no glyphosate), and each glyphosate concentration. Sealed plates were left for 18 h incubation at 30 °C while shaking at 250 rpm (PST-60HL-4 Plate Shaker Thermostat,

Biosan). The absorbance at 620 nm was then measured for each well (Thermo Multiskan Plate Reader). Finally, hydrolase activity in *B. subtilis* was measured by adding 20 µL fluorescein diacetate stock solution (5 mM) to each well to obtain a final concentration of 5 µM. After 60 minutes incubation at 30 °C and shaking at 250 rpm, fluorescence was quantified in each well using a Victor™ X2 Multilabel Plate Reader with a 485 nm excitation and 535 nm emission filter (Perkin Elmer).

1.3.3 *Raphidocelis subcapitata*

Toxicity of glyphosate to phytoplankton was examined by the unicellular green microalgae *Raphidocelis subcapitata* (ISO 8692, 2012) [173] inhibition tests. The toxicological endpoint was inhibition of growth measured after 72 h of incubation (ISO 8692, 2012) [173]. *R. subcapitata* (MicroBioTests Inc) was cultivated in a test medium at 23 ± 2°C and under continuous illumination at 6500 lux (ISO 8692, 2012) [173]. Diluted culture was exposed in 96-well clear microplates (Nunclon, Thermo Scientific) to the following concentrations of glyphosate with and without prior UV irradiation: 0,098, 0,195, 0,390, 0,780, 1,560, 3,130, 6,250, 12,5, 25, 50 mg/L. Eight replicates were included for blanks (medium), controls (no glyphosate), and each glyphosate concentration. Plates were incubated at 23 °C shaking at 70 rpm under continuous illumination (6500 lux) for 72 h. Growth was measured after 0, 24 h, 48h, and 72h as absorbance at 450 nm using a Thermo Multiskan Plate Reader. Growth measurements for selected samples were done by measuring cell sizes (µm) and cell abundance (cells/mL) using a Multisizer 4e Coulter Counter (Beckman Coulter).

1.3.4 *Daphnia magna*

Glyphosate's toxicity to zooplankton was examined with inhibition tests of the crustacean *D. magna* (ISO 6341, 2012) [174]. The toxicological endpoint was inhibition of mobility and was determined by visual inspection of the animals (ISO 6341, 2012) [174]. *D. magna* STRAUS was cultivated from a laboratory clone originating from pure-culture ephippia [175]. Each treatment consisted of 20 juvenile animals distributed among 4 glass vials containing 5 animals and 10 mL freshwater medium in each. The mobility of each animal was determined after 24 h and 48 h (ISO 6341, 2012) [174].

1.4 Chromatographic Analysis

Analyses of the samples before and after UV treatment were done using an Ultimate 3000 High-Pressure Liquid Chromatography coupled to a high resolution LTQ-Orbitrap mass spectrometer (Thermo Scientific, Bremen, Germany) equipped with an ESI source.

Chromatographic separation was achieved with a reversed-phase C18 column (Phenomenex Luna, 150 × 2 mm, 3 μm, 110 Å; Phenomenex, Castel Maggiore, BO, Italy) by injecting a sample volume of 10 μL at a mixture of 0,1 mM Formic Acid (A) and Acetonitrile (B) used as the mobile phase. The elution followed a gradient profile which started from 5% B, increased up to 100% B in 40 minutes and arrived to 100% A after 10 minutes. Mass spectra were acquired in both Positive and Negative ESI modes. Nitrogen was used as sheath and auxiliary gas in order to deliver the LC effluent to the ESI ion source with the following parameters: auxiliary gas 15 arb, sheath gas 34 arbitrary unit (arb), capillary voltage 4.48 kV, and capillary temperature 270°C. The elaboration of the data followed a suspect screening technique and was carried out using the MZmine 2.52 [167] software in order to achieve peak alignment, peak grouping, chromatogram deconvolution and isotope removal (Table A8). The online databases ChemSpider and METLIN were used in order to identify the transformation byproducts [176-178].

1.5 Data analysis and statistics

The toxic response measured for all the endpoints was expressed as inhibition (I) relative to control samples:

$$I = 1 - (R_i / R_c),$$

where R_i is the response measured for inhibited samples and R_c is the one measured for control samples.

The concentration-response curves were fitted to a log-logistic model using iterative non-linear regression [179]:

$$\text{Response} = 1 / (1 + 10^{(\log EC_{50} - \log C) * \text{Slope}}) \quad (13)$$

where C is the concentration of the toxicant (mg/L), EC_{50} is the median effective concentration (mg/L), and Slope is the parameter that represents the slope of the curve. Non-linear regression models and calculation of 95% confidence limits for EC_{50} values were performed using the GraphPad Prism 8.0.1 (Graphpad Software).

Relative Effect Potency (REP) [179] was used in order to estimate the toxicity of a sample before and after UV treatment:

$$\text{REP} = EC_{50(\text{before})} / EC_{50(\text{after})} \quad (14)$$

where $EC_{50(\text{before})}$ is the median effective concentration (mg/L) before UV irradiation and $EC_{50(\text{after})}$ is the median effective concentration (mg/L) after treatment.

The nonparametric Mann-Whitney U test (Wilcoxon rank-sum test) was used to perform statistical analyses of the results and evaluate differences between treatments, that have a significance level of $p < 0,05$ (KaleidaGraph 4.5.4; Synergy Software).

2. Results

2.1 Test organisms

Initial experiments were conducted in order to identify which test organisms were responsive to glyphosate exposure (Figure 15). A battery of non-target organisms including *Bacillus subtilis*, *Aliivibrio fischeri*, *Raphidocelis subcapitata*, and *Daphnia magna* was tested. The results showed that the traditional toxicity screening organism *A. fischeri* was the least responsive with a median effective concentration (EC₅₀) value of 25,0 mg/L. On the other hand, the crustacean *D. magna*, the bacterium *B. subtilis*, and the green microalgae *R. subcapitata* responded at the exposure of much lower glyphosate's concentrations with EC₅₀ values of 0,990 mg/L, 3,670 mg/L, and 1,130 mg/L, respectively (Figure 15).

Subsequently, we decided to focus on *D. magna*, *B. subtilis* and *R. subcapitata* for our experiments, examining the changes in toxicity before and after UV irradiation of aqueous glyphosate.

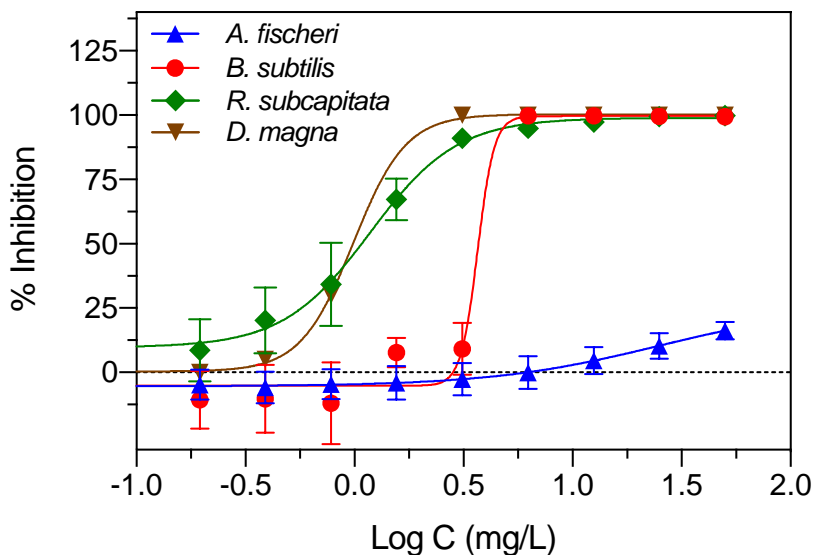


Figure 15: Concentration-response curves showing the toxicity of glyphosate to different aquatic test organisms. Data points represent means \pm standard deviation.

2.2 Effect of UV irradiation on glyphosate toxicity

The toxicity of glyphosate before and after exposure to UV irradiation under three different wavelengths - UV-A (365 nm), UV-B (302 nm), UV-C (254 nm) - was evaluated using *B. subtilis*, *R. subcapitata* and *D. magna* as test organisms (Figure 16, Figure 17, Table 17).

Exposure of glyphosate to UV-A and UV-B at a UV dose of 20 J/cm² did not have any noticeable effect on the toxicity to *B. subtilis* and *R. subcapitata* (Figure 16).

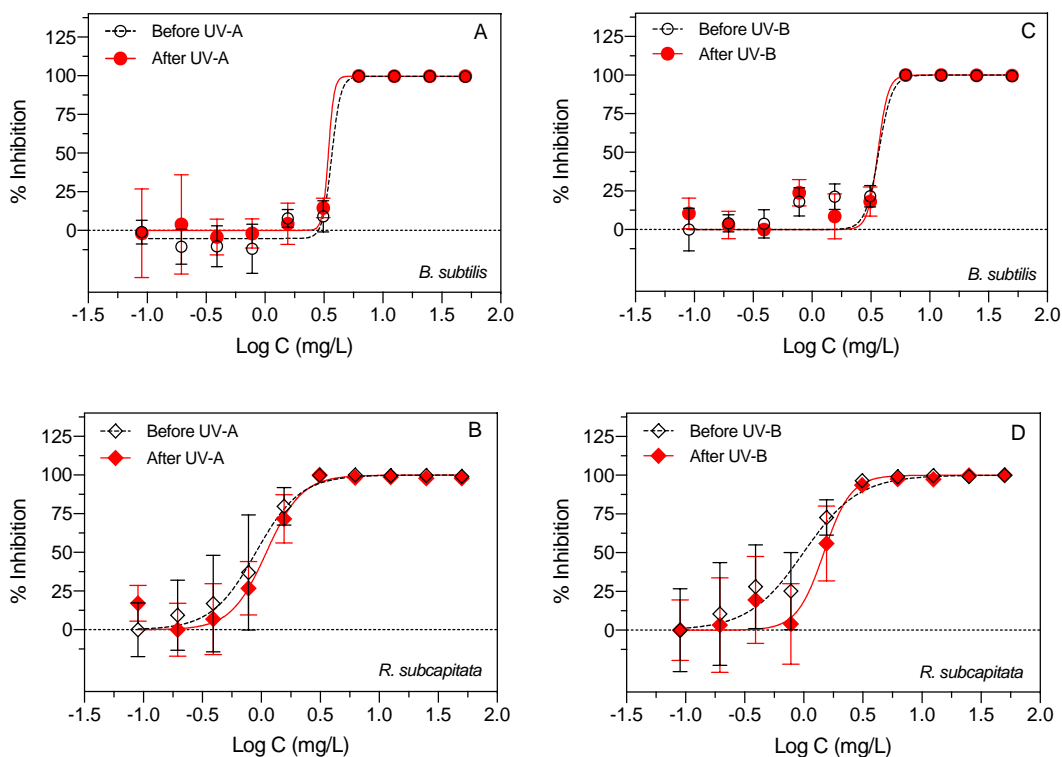


Figure 16: Effect of UV-A irradiation (panel A and B) and UV-B irradiation (panel C and D) at a UV dose of 20 J/cm² of aqueous glyphosate on toxicity to *B. subtilis* (panel A and C) and *R. subcapitata* (panel B and D). Data points represent means ± standard deviation.

The same result was achieved also after increasing the UV-A dose to 70 J/cm² (Figure 17D). However, exposure of glyphosate to UV-B at a dose of 70 J/cm² resulted to a significant toxicity decrease comparing the dose of 20 J/cm² (Mann-Whitney test; p=0,028). On the other hand, UV-C exposure at UV dose of 20 J/cm² clearly decreased the toxicity of aqueous glyphosate to *B. subtilis* and *R. subcapitata* (Figure 17A-C). Furthermore, when the UV-C dose was increased from 20 J/cm² to 70 J/cm² showed a significant decrease of the toxicity of glyphosate (Mann-Whitney; p=0,029). This suggests that UV-C and also UV-B irradiation are able to decrease the ecotoxicity of glyphosate if the UV dose is sufficiently high.

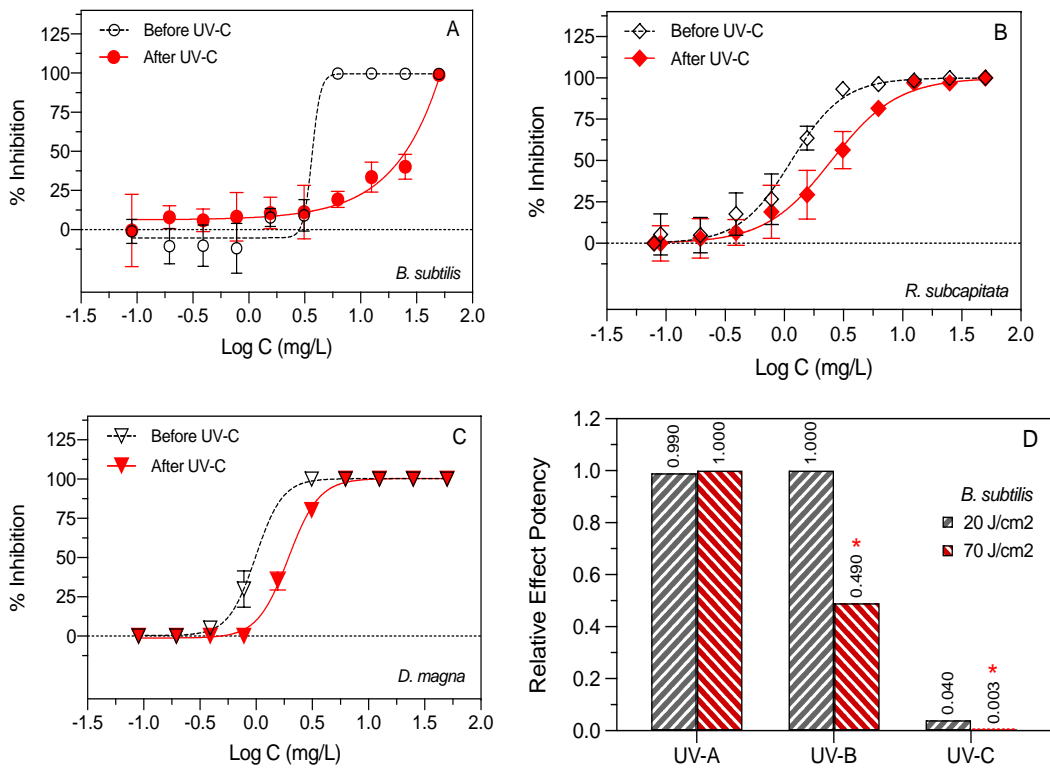


Figure 17: Effect of UV-C irradiation (A, B, and C) of aqueous glyphosate at a UV dose of 20 J/cm² on toxicity to *B. subtilis* (A), *R. subcapitata* (B), and *D. magna* (C). Data points represent means \pm standard deviation. D shows the effect of an increased UV irradiation dose of 70 J/cm² on the relative effect potency of glyphosate to *B. subtilis*. The asterisk (*) indicates the significant difference between 20 J/cm² and 70 J/cm² (Mann-Whitney test, p<0.05).

Changes in the growth of *R. subcapitata* after UV-C treatment of glyphosate were measured as changes in the absorbance as described in international standards (ISO 8692, 2012)[173]. The results were also confirmed by counting and sizing individual algae cells using a Multisizer Coulter Counter (Figure 18). Important changes in the cell numbers of *R. subcapitata* were observed after its 72 h exposure and growth in solutions which had and had not undergone UV-C irradiation (Figure 18). The results of the Mann-Whitney U test, which was done for each of the four glyphosate concentrations, showed that the difference between non-irradiated and UV-C irradiated solutions was significant ($p=0,026$; $p<0,001$; $p<0,001$; $p<0,001$, respectively).

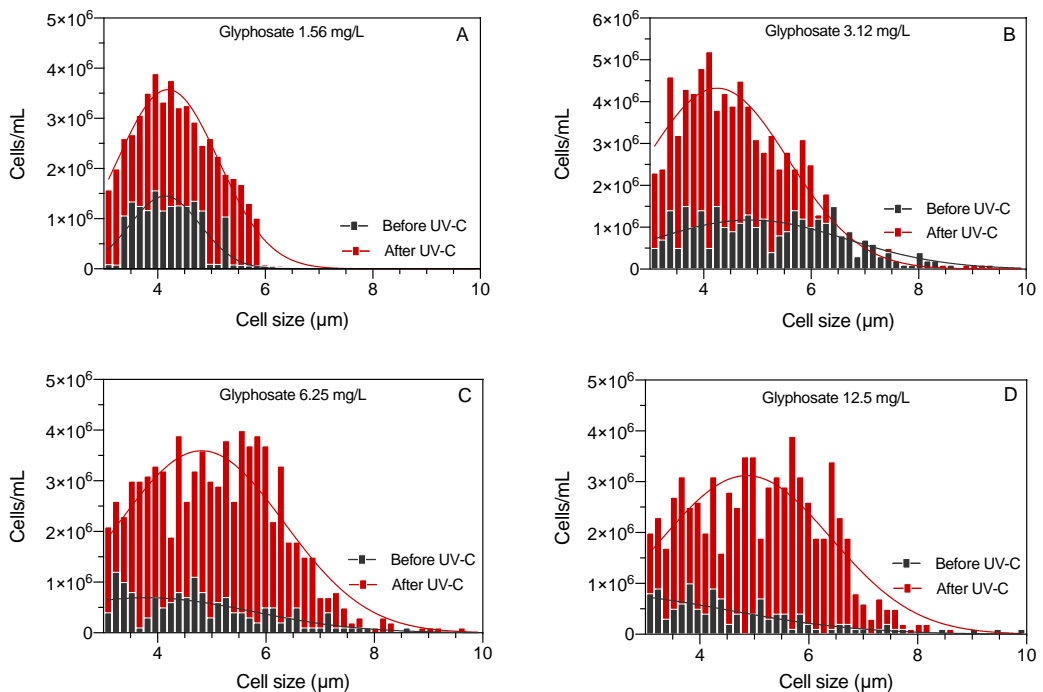


Figure 18: Effect of UV-C irradiation (20 J/cm^2) of aqueous glyphosate on toxicity to *R. subcapitata* measured as differences in cell numbers and cell sizes after 72 h of growth in the presence of glyphosate before and after UV treatment.

In the present study, the EC₅₀ value for *D. magna* exposed to glyphosate was found to be 0,99 mg/L, a value comparable to those reported in literature [175,180,181], while for *R. subcapitata* was 1,13 mg/L and for *B. subtilis* 3,67 mg/L (Table 17) -values also

within the range reported in other studies [182-186]. The median effective concentration (EC50) values of glyphosate samples were increased after UV-C irradiation at a UV dose of 20 J/cm² for all the three test organisms, a result that suggests lower toxicity (Table 17). A 2-fold decrease in toxicity of glyphosate to *D. magna*, a 5-fold decrease for *R. subcapitata* and a 23-fold decrease *B. subtilis* were observed.

Table 17. Median effective concentrations (EC50) values for three aquatic test organisms before and after exposure of glyphosate to UV-A, UV-B, or UV-C at comparable UV doses (20 J/cm²). *ND: not determined.*

| Test organism | EC50 values (mg/L) | | | |
|-----------------------|--------------------|------------|------------|------------|
| | Before UV | After UV-A | After UV-B | After UV-C |
| <i>B. subtilis</i> | 3,67 | 3,45 | 3,54 | 85,41 |
| <i>R. subcapitata</i> | 1,13 | 1,19 | 1,53 | 5,70 |
| <i>D. magna</i> | 0,99 | ND | ND | 1,93 |

3.4 Effect of different UV-C exposure conditions on glyphosate toxicity

Increase of irradiation time and UV dose resulted to an exponential decrease of the toxicity of glyphosate to *B. subtilis* and *R. subcapitata* (Figure 19). Increasing UV doses and decreasing toxicity, a relationship calculated as $\log(1/EC50)$ suggests that a loss of 90% of the initial glyphosate to the bacterium after UV-C irradiation of 23,4 J/cm² and of 23,7 J/cm² for the green algae occurs. Therefore, toxicity was mitigated for both organisms in a dose-dependent ratio and at comparable grades, suggesting that the test organisms responded equally to attenuation of UV-C glyphosate inhibition (Figure 19).

Mainly, the exposure of aqueous solutions of glyphosate to UV irradiation was carried out in sealed quartz samples. Examining though the different conditions that could affect the degradation, a different exposure method using UV-C plastic transparent microplates was considered as well. However, for reasons of eliminating sample loss due to the small amount of sample treated in the microplates (100µL) and based on the results that lower of 10 J/cm² UV doses may affect the toxicity, we focused on a comparison between the two exposure techniques at a UV-C dose of 5,4 J/cm². The results obtained did not show great differences in median effective concentrations for

B. subtilis and *R. subcapitata* for the different exposure regimes (quartz cuvettes vs. transparent plastic microplates) (Table 18).

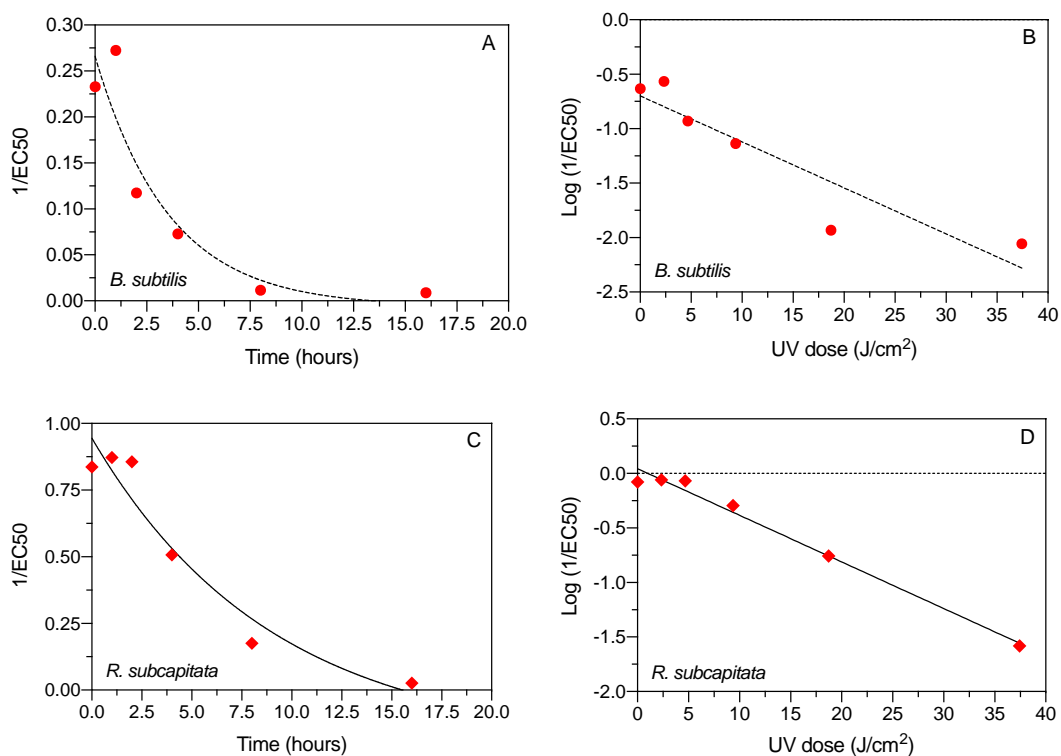


Figure 19: Effect of UV-C irradiation time (h) and UV dose (J/cm²) on the toxicity of glyphosate to *B. subtilis* (A and B) and *R. subcapitata* (C and D).

Table 18. Median effective concentrations (EC50) for *B. subtilis* and *R. subcapitata* before and after exposure of glyphosate to 5,4 J/cm² UV-C, in two different UV exposure techniques

| Exposure technique | <i>B. subtilis</i> EC50 values (mg/L) | | <i>R. subcapitata</i> EC50 values (mg/L) | |
|--------------------|--|------------|---|------------|
| | Before UV-C | After UV-C | Before UV-C | After UV-C |
| Quartz Glass | 3,67 | 7,92 | 1,13 | 4,18 |
| UV plate | 2,34 | 7,12 | 1,47 | 3,65 |

In this study, control samples for all the exposure experiments using blank solutions without glyphosate were included, in order to assess potential toxicity originating from the generated reactive radicals. The results showed no apparent inhibition of test organisms, an outcome that can be explained by the fact that these products are short-lived.

3.5 Effect of UV irradiation on glyphosate's toxicity in real water samples

The effect of UV irradiation on mitigating glyphosate's toxicity was also studied in real water matrices, in order to simulate more realistic conditions, and examine the effects from unknown water constituents, such as organic and inorganic contaminants, on degradation. Initially, the tests were conducted in a test matrix prepared with deionized distilled water and artificial freshwater. After some promising initial results, natural drinking water samples spiked with glyphosate concentrations were used for UV-C irradiation experiments. The obtained results (Figure 20) showed clear differences in toxicity before and after UV-C irradiation of 20 J/cm². In some cases, the decrease in toxicity due to UV-C treatment of aqueous glyphosate was slightly larger for the natural drinking water samples compared to parallel experiments conducted in distilled water (Figure 17 vs. Figure 20). The UV effect was also greater for glyphosate added to drinking water compared to groundwater (raw water) (Figure 20F). The raw water was slightly colored and contained elevated concentrations of natural elements such as iron, manganese, and ammonia because it was sampled before filtration at the Drinking Water Treatment Plant. For glyphosate irradiated in drinking water, the EC50 values before UV-C varied between 2,03 mg/L and 7,30 mg/L whereas the EC values after UV-C varied between 17,57 mg/L and >100 mg/L. The differences in EC50 before and after UV-C irradiation were significantly different (Mann-Whitney; p=0,002). The Relative Effect Potency after UV-C treatment was 0,02-0,40 corresponding to a 3 to 44-fold reduction in toxicity to the test organism *B. subtilis*. Hence, the effect of UV on glyphosate was not inhibited by constituents in the drinking water samples and may even be stronger in some water matrices than in deionized water suggesting that natural drinking water may even facilitate the process.

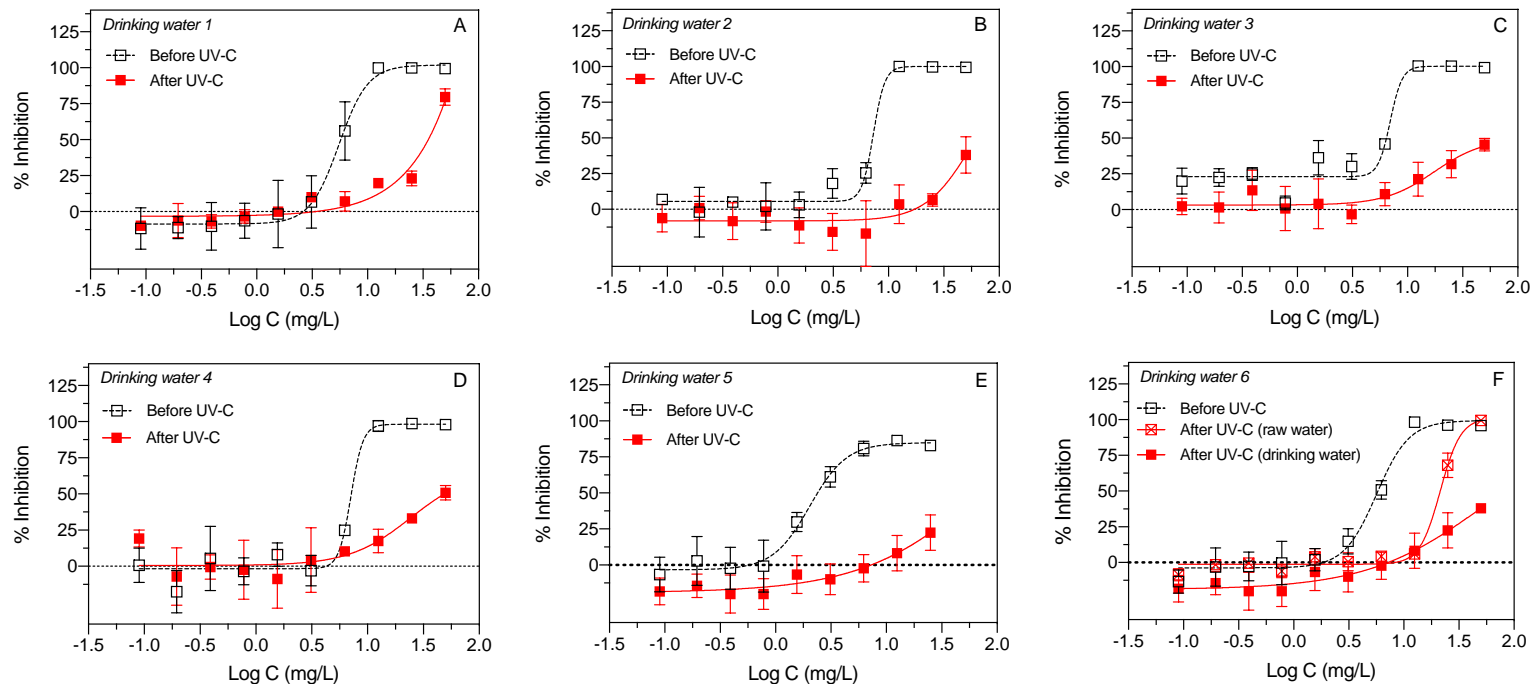


Figure 20: Effect of UV-C irradiation (20 J/cm²) of glyphosate in municipal drinking water on toxicity to *B. subtilis*. Drinking water produced from groundwater was collected at six locations in three Danish municipalities: Aalborg Municipality (A – Aalborg East; B - Aalborg Center; C – Aalborg West) Sønderborg Municipality (D); Aarhus Municipality (E), and Elsted drinking water treatment plant in Aarhus Municipality sampled before and after water treatment (F). Data points represent means ± standard deviation.

3.6 Byproducts' identification

Liquid Chromatography tandem with High Resolution Mass Spectrometry (LC-HRMS) was used in order to analyze the aqueous solutions of glyphosate before and after UV exposure in order to identify the transformation products and understand the degradation mechanism. Samples after UV-C irradiation at doses of 20 J/cm² and 70 J/cm² were used. The identification followed a suspect screening approach focusing on transformation products likely to be generated from glyphosate, and was done using the ChemSpider and METLIN databases (<http://metlin.scripps.edu/>) [176-178]. However, it was a challenging procedure due to lack of analytical standards. More than 20 byproducts were detected (Table 19) after UV-C exposure at 20 J/cm² and 70 J/cm². Among them, there were Sarcosine (C₃H₇NO₂), Glycine (C₂H₅NO₂), Glyoxylic acid (C₂H₂O₃), Aminomethylphosphonic acid (CH₆NO₃P; AMPA), Acetic acid (C₂H₄O₂) and Phosphoric acid (H₃PO₄), transformation byproducts already mentioned in other studies employing different Advanced Oxidation Processes for glyphosate's remediation, as well as other potentially intermediates.

Table 19. Detected transformation products after UV-C irradiation in negative ESI (-) and positive ESI (+).

| Compound | <i>m/z</i> | RT (min) | ESI | Compound | <i>m/z</i> | RT (min) | ESI |
|---|------------|----------|-----|---|------------|----------|-----|
| H ₃ O ₄ P | 96,968 | 25,94 | - | C ₅ H ₉ NO ₂ | 116,069 | 15,66 | + |
| CH ₆ NO ₃ P | 110,001 | 1,75 | - | C ₃ H ₃ NO ₂ | 84,009 | 27,82 | - |
| C ₂ H ₅ NO ₂ | 74,020 | 1,86 | - | C ₃ H ₄ O ₄ | 103,003 | 1,69 | - |
| C ₂ H ₂ O ₃ | 72,908 | 16,64 | - | C ₃ H ₆ O ₄ | 105,017 | 26,14 | - |
| C ₃ H ₇ NO ₂ | 90,054 | 29,62 | + | C ₃ H ₆ O | 58,080 | 11,80 | + |
| C ₂ H ₇ NO ₂ | 77,084 | 1,77 | - | C ₃ H ₉ NO ₂ | 92,069 | 1,66 | + |
| C ₂ H ₅ NO | 59,070 | 35,32 | - | C ₄ H ₁₀ O ₂ | 91,074 | 28,37 | + |
| C ₂ H ₆ N ₂ O ₄ | 91,016 | 28,99 | - | C ₄ H ₈ O | 73,028 | 4,43 | + |
| C ₂ H ₄ O ₂ | 59,015 | 8,52 | - | C ₄ H ₈ O ₄ | 119,033 | 6,85 | - |
| C ₃ H ₈ N ₂ O | 89,070 | 30,13 | + | C ₄ H ₁₀ O ₃ | 106,120 | 29,42 | - |
| C ₆ H ₁₁ NO ₂ | 130,085 | 28,58 | + | C ₅ H ₅ N | 80,048 | 32,84 | + |

The proposed degradation mechanisms of Glyphosate after UV photolysis are shown in Figure 21. Both pathways are following those proposed from previous studies and are resulting in less toxic transformation products confirming the toxicity test results [109]. The identified byproducts, show that in UV photolysis treatment, degradation of Glyphosate follows the two mechanisms related to the C-P and C-N bonds, known as the “C-P pathway” and the “C-N pathway”. In the first one, Glyphosate’s (A) molecule is attacked by the generated hydroxyl radicals resulting to the cleavage of the C-P bond, the formation Sarcosine (B) and the release of Phosphate which after hydrogenation results to Phosphoric acid (C) production. Sarcosine after further treatment is subsequently transformed to Glycine (F), which can be further degrade to Glyoxylic acid (G) and Acetic acid (E). The second pathway involves the breakdown of the electrophilic C-N bond, or directly by UV irradiation, or after attack by the generated hydroxyl radicals, and results to AMPA (D) and Acetic acid (E), or directly to Glyoxylic acid (G). These two mechanisms can exist alone or together during Glyphosate’s oxidation process, and are the same as those occurring through biodegradation processes. Hence, the results of this study suggest a potential combination of UV treatment with bioremediation to increase the remediation of Glyphosate further. Detection of other compounds was done as well, for which a

Moreover, LC-HRMS analyses showed that concentrations of glyphosate after UV-C irradiation at 20 J/cm^2 were no longer detected after treatment at a dose of 70 J/cm^2 . Subsequently, the concentrations of AMPA and glycine were increasing after 70 J/cm^2 , while sarcosine’s increased with the treatment at 20 J/cm^2 and later decreased. Phosphoric acid was detected at both high and low UV-C doses while glyoxylic and acetic acid were only observed after UV-C irradiation at 70 J/cm^2 . Concentrations were not quantified, since the analyses followed a full scan acquisition for identifying transformation products. However, differences in peak areas were considered for evaluating the differences in the samples before and after treatment.

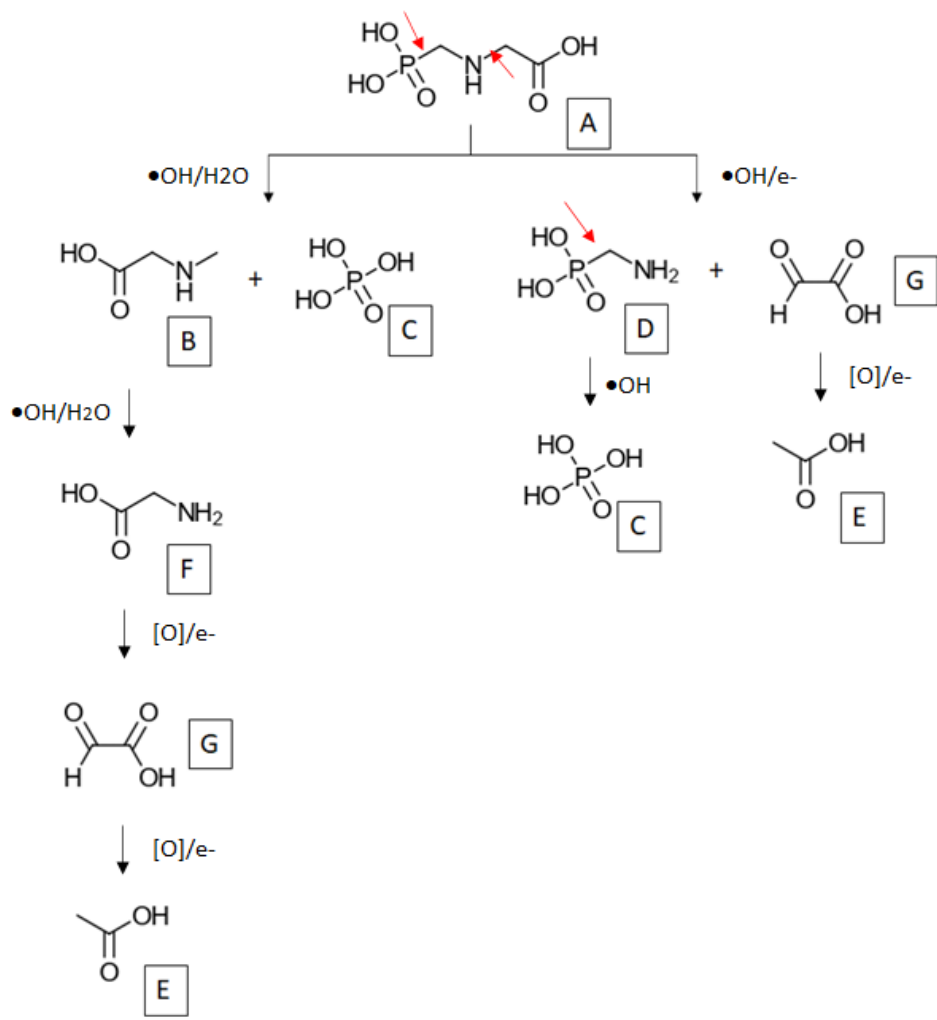


Figure 21: Potential pathways for UV mediated photolysis of Glyphosate (A) through two main routes resulting in Sarcosine (B), and Phosphoric acid (C), and AMPA (D), Acetic acid (E) and Glyoxylic acid (G). Sarcosine (B) may subsequently be transformed into Glycine (F).

3. Conclusions

Glyphosate is the most frequently detected herbicide in the aquatic environment, posing threats to the ecosystem and human health. Thus, efficient remediation techniques for its removal are fundamental. A vast variety of studies employing different methods have reported satisfactory results. However, towards a sustainable and toxic-free environment the implementation of fast, cost-efficient and environmental friendly techniques that don't generate waste or more toxic byproducts are necessary. In this way, the aim of this study was to investigate the degradation of glyphosate after UV-A, UV-B, and UV-C irradiation. Different parameters that could affect the process were taken into account and their effects on toxicity to aquatic organisms from different trophic levels were studied. The effect-based monitoring approach for the evaluation of the method's efficiency was combined with chemical analyses in order to identify transformation byproducts. Toxicity assays are an important supplement to chemical analyses in order to assess water quality, as bioassays can integrate changes in water chemistry and bioactivity before and after water treatment.

The results suggest that UV-C and to some extent UV-B photolysis of glyphosate in water could decrease concentrations of this pesticide and reduce overall ecotoxicity by generating less toxic transformation products. Even if, UV doses used in treatment plants are mainly used for disinfection processes, and they are not adequate to degrade organic contaminants, UV photolysis represents one cost-efficient, and green remediation process that doesn't produce waste and should be considered for industrial scale applications.

Chapter 7 Degradation of PFAS

Per and polyfluoroalkyl substances (PFASs) represent one of the most problematic classes of compounds - whose occurrence in different water sources has been reported worldwide - due to their ubiquitous physicochemical properties, that make them extremely difficult to be degraded and subsequently very persistent in the environment. At present, activated carbon and ion exchange are mainly used as removal technologies of PFASs from water, facing efficiency problems with shorter chain compounds [187-189] and generating waste that need to undergo further treatment, producing additional costs. For this reason, search of new remediation technologies is necessary. AOPs have been reported of being able to efficiently degrade PFASs, with promising defluorination of the compounds being reported after following electrochemical methods [190]. However, further study in order to identify the produced byproducts for understanding the degradation mechanisms and the toxicity of these compounds is still required. For this reason, in this work we evaluated the degradation mechanism of PFASs after non-thermal plasma treatment and identified the produced byproducts in individual compounds' solutions and in a mixture. The effect of different matrices was also examined, by preparing solutions in both MilliQ water and real groundwater matrices.

1. Materials and Methods

1.1 Selection of compounds

The compounds included in this study were selected based on the study reported in Chapter 3 with the highest abundance in the environment and after following the trends in literature. In this way, 3 substances were chosen, two perfluoroalkyl acids, Perfluorooctanoic acid (PFOA) and Perfluorohexanoic acid (PFHxA), and one perfluorosulfonic acid, Perfluorooctanesulfonic acid (PFOS). Two of the compounds had a carbon chain with 8 atoms while a shorter chain compound, including 6 atoms of carbon was included in order to test the efficiency of the thermal plasma treatment to degrade also shorter chain compounds.

1.2 Reagents and Chemicals

Stock compounds of PFOA (Perfluorooctanoic acid, CAS 307-24-4), PFOS (Perfluorooctanesulfonic acid, CAS 1763-23-1) and PFHxA (Perfluorohexanoic acid, CAS 335-67-1) (linear chain only) were purchased from Sigma-Aldrich (Saint Louis, MO, USA). MilliQ was obtained from MilliPore (MA, USA), UHPLC-grade Methanol was purchased from Sigma-Aldrich (Saint Louis, MO, USA), LC-MS grade Water LiChrosolv® and

Ammonium acetate for LC-MS LiChropur® were purchased from Merck KGaA (Darmstadt, Germany).

1.3 Sample preparation and treatment

Solutions of the three selected compounds, PFOA, PFOS and PFHxA, were prepared in MilliQ water and a real water matrix (groundwater) at a concentration of 1 µg/L. Treatment of the different solutions was done using a custom-built Marx generator powered with 220V AC equipped with a pulse-width modulation circuit, a high-voltage transformer and four 990 pF capacitors. A small continuous flow of compressed air with relative humidity around 14% was fed to the spark-gaps' chamber to stabilize the generator's internal atmosphere. The discharge's peak voltage was typically 100-130 kV with peak current values of 20-40 A. The pulse duration was approximately 250 ns and the frequency of discharge could be manually adjusted between 5 and 17 Hz. Electrical measures have been performed using a BK Precision 2190D oscilloscope. Total absorbed power of the generator laid between 299 and 322 W. Water was treated using two different reactors: a 20 mL cylindrical polypropylene (PP) reactor was used in the first part of the work aimed at the optimization of working parameters while a 50 mL Pyrex glass (PG) reactor was used for all other experiments. The PP reactor on one site limited the risk of damage of the reactor wall as a consequence of the generated shock waves, whilst on the other side caused Total Organic Carbon (TOC) and Total Nitrogen (TN) potential contamination from plastics and rubber parts. For both reactors, the same electrodes have been used consisting in one stainless steel bar having 10 mm diameter and a tungsten sintered electrode with a diameter of 3 mm on the other side. In the PG reactor the distance between the electrodes determined whether the discharge was forming underwater (distance < 2 mm) or at the water surface (distance > 2 mm). For both reactors, the headspace was in connection with the room air. Schematic representation of the experimental setup is shown in Figure 22. Both the reactors and the plasma generator were placed inside a Faraday cage. The optimization of its functional parameters was done in another study from our research group [191] using experimental design.

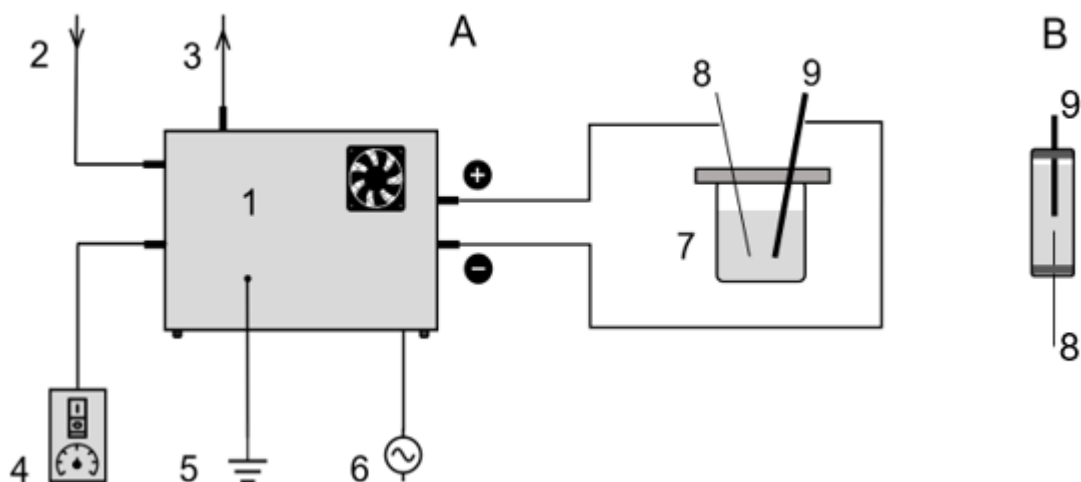


Figure 22: Schematic representation of the plasma generator connected to the PG reactor (A): 1) plasma generator, 2) air inlet, 3) air outlet, 4) plasma switch and frequency control, 5) ground, 6) 220V power supply, 7) PG reactor, 8) sintered tungsten electrode, 9) stainless steel electrode. PP reactor (B).

At the beginning of the experiments the reactor was filled with 50ml of each solution for treatment. The treatment time was 30 minutes and samples to be analyzed were taken at different time intervals (0min, 1min, 2,5min, 5min, 7,5min, 10min, 15min, 20min, 30min) in order to evaluate the degradation profiles and understand the generation of by-products. A mix of the compounds was prepared as well, in the same matrices and with a final concentration of 1 $\mu\text{g/L}$ for each compound. The treatment time was extended to 60 minutes with samples taken at different time intervals (0min, 1min, 2,5min, 5min, 7,5min, 10min, 15min, 20min, 30min, 40min, 50min, 60min) in order to observe the differences in the degradation mechanisms, understand the potential cocktail effects and overall simulate a more realistic situation, since these compounds usually exist in the environment as mixes. Caution throughout the experiments – as reported in Chapter 3- was taken in order to minimize as much as possible the background contamination of the samples, that could potentially interfere with the results. In this way, Teflon materials were avoided throughout the whole treatment and analysis time.

1.4 Instrumental Analysis

Samples before and after treatment were analyzed with a SCIEX X500R QTOF system coupled to a Shimadzu ExionLC UHPLC system. The chromatographic separation was achieved using a Luna® Omega Polar C18 100 LC Column (3µm particle size, 100 x 2.1mm) –heated at 40°C- by injecting a 50µL sample volume into a mixture of 5mM Ammonium Acetate in H₂O (A) and 5mM Ammonium Acetate in MeOH (B) as mobile phase at a flow of 0,350 mL/min. The elution followed a gradient profile starting from 95% A and 5% B, keeping this ratio for 1 minute and then gradually changing to 100% B within 10 minutes, and keeping it for the next 2 minutes. Finally, the conditions returned to the initial within 15 minutes of elution. The samples were kept at a temperature of 4°C throughout the whole analysis time. The X500R QTOF system operated in Negative Electrospray Ionization Mode (ESI) with the following parameters: Ion source gas 1: 45 psi, ion source gas 2: 55psi, curtain gas: 30 psi, gas temperature: 500°C, ionspray voltage: -4500V, in both MRM and SWATH acquisition modes.

The samples were analyzed with the MRM acquisition mode in order to follow a target screening, which allowed the estimation of the degradation rate of the molecules. Quantification was achieved with six points calibration curves of final concentrations of 10, 50, 100, 500, 1000 and 5000 ng/L for each compound. Good coefficient results with $R^2 > 0,990$ were obtained for all the compounds.

On the other hand, SWATH was used for a qualitative screening, in order to identify degradation byproducts and it consisted of a full scan acquisition, followed by a Q1 isolation. The parameters used in full-scan MS mode were as follows: accumulation time: 0,05 sec; declustering potential: -80 V; TOF start mass: 100 Da; TOF stop mass: 1000 Da. A generic collision energy spread of $(-35) \pm 15$ was used. For the Q1 isolation strategy (MS/MS) the parameters were: TOF start mass: 50 Da; TOF stop mass: 1500 Da; total number of windows: 24; window accumulation time: 0,035 sec. An external calibration of the instrument was performed, using a mixture of 10 compounds with a mass range between m/z 68,99 and m/z 2233,91. This mixture was automatically injected every 5 samples in order to maintain the mass accuracy below 2 ppm. SCIEX OS 1.7 software (SCIEX, Massachusetts, USA) was used for data acquisition and elaboration.

1.5 Toxicity tests

Three different methods were used in order to examine the toxicity of the three individual compounds' samples before and after treatment. Two *In vitro* bioassays based

on the acute toxicity effects on the crustacean *Thamnocephalus platyurus* and the bioluminescent bacteria *Aliivibrio fischeri*, and one *In silico* method were followed.

1.5.1 *Thamnocephalus platyurus*

The commercially available Thamnotoxkit FTM microbiotest [192] was used for determining the lethal effects of toxicants on *Thamnocephalus platyurus* after freshly hatched larvae's exposure to samples before and after 30 minutes treatment, for 24h. The assays were performed according to the standard operational procedures declared in the form ISO 14380:2011(E) [193]. The lethal responses of the microorganisms were measured after 24h of incubation at 25°C.

1.5.2 *Aliivibrio fischeri*

Toxicity screening of the three PFAS samples was examined with a standard inhibition test using the luminescent bacterium *Aliivibrio fischeri* (ISO 11348-1, 2009) [172]. Freeze-dried bacteria, reconstitution solution, diluent (2% NaCl) and an adjustment solution (non-toxic 22% sodium chloride) were obtained from Azur (Milan, Italy). Samples were prepared in a medium containing 2% sodium chloride. Changes in bioluminescence were recorded after 5, 15 and 30 minutes of incubation at 15 °C, using a Microtox Model 500 toxicity analyzer.

1.5.3 *In silico*

In silico toxicity was evaluated regarding LC50 values on fish (96h LC50 values), daphnids (48h LC50 values) and green algae (96h LC50 values) using the ECOSAR tool (developed by EPA) -based on Quantitative Structure-Activity Relationship (QSAR) mathematical models.

2. Results

2.1 Degradation rate

Individual solutions of PFOA, PFOS and PFHxA both in MilliQ and in groundwater matrices were treated using surface plasma discharge, performing all the experiments in triplicates for better evaluating the degradation rate. Figure 23 summarizes the degradation rates of the three compounds in individual solutions of both matrices. For all the substances, the degradation started immediately after treatment but followed different rates. More specifically, the best results were obtained for PFOS, showing a 100% removal in MilliQ and 85% in groundwater matrix after 30 minutes of treatment. PFHxA showed a 40% removal after 30 minutes of treatment in MilliQ and a 35% removal in real water, while PFOA showed a 50% removal at the end of the treatment time, regardless the matrix.

When the compounds were treated in a mixture, the degradation rate was slower for all of them, as shown in figure 23D, due to the competition among the substrates for the produced reactive species. For that reason, the treatment time was extended to 60 minutes. In general, the degradation profiles were similar to those obtained for individual treatments. More specifically, degradation of PFOS showed again the best results, with 85% of removal in MilliQ and 79% in groundwater after the 30 minutes of treatment, with a complete removal after 40 minutes of treatment in both matrices. PFOA in MilliQ water showed a 40% removal after 30 minutes of treatment, which arrived to 50% after 1 hour, and 32% removal after 30 minutes when prepared in groundwater, which eventually achieved a 44% removal after 1 hour. Finally, PFHxA showed the lowest removal when treated in the mix as well, with 26% after 30 minutes of treatment in MilliQ and 21% in groundwater, arriving to 45% and 32% removal after 1 hour respectively.

In all cases, an exponential decay of the concentration following a pseudo first order kinetics was observed, with those in groundwater showing a lower degradation rate, indicating effects from unknown constituents in the water. The results obtained from this study are in accordance with those reported in literature, with longer chain compounds (with more than 8 atoms of carbon) showing higher degradation rates, and those with shorter chains (with less than 8 atoms of carbon) being more recalcitrant.

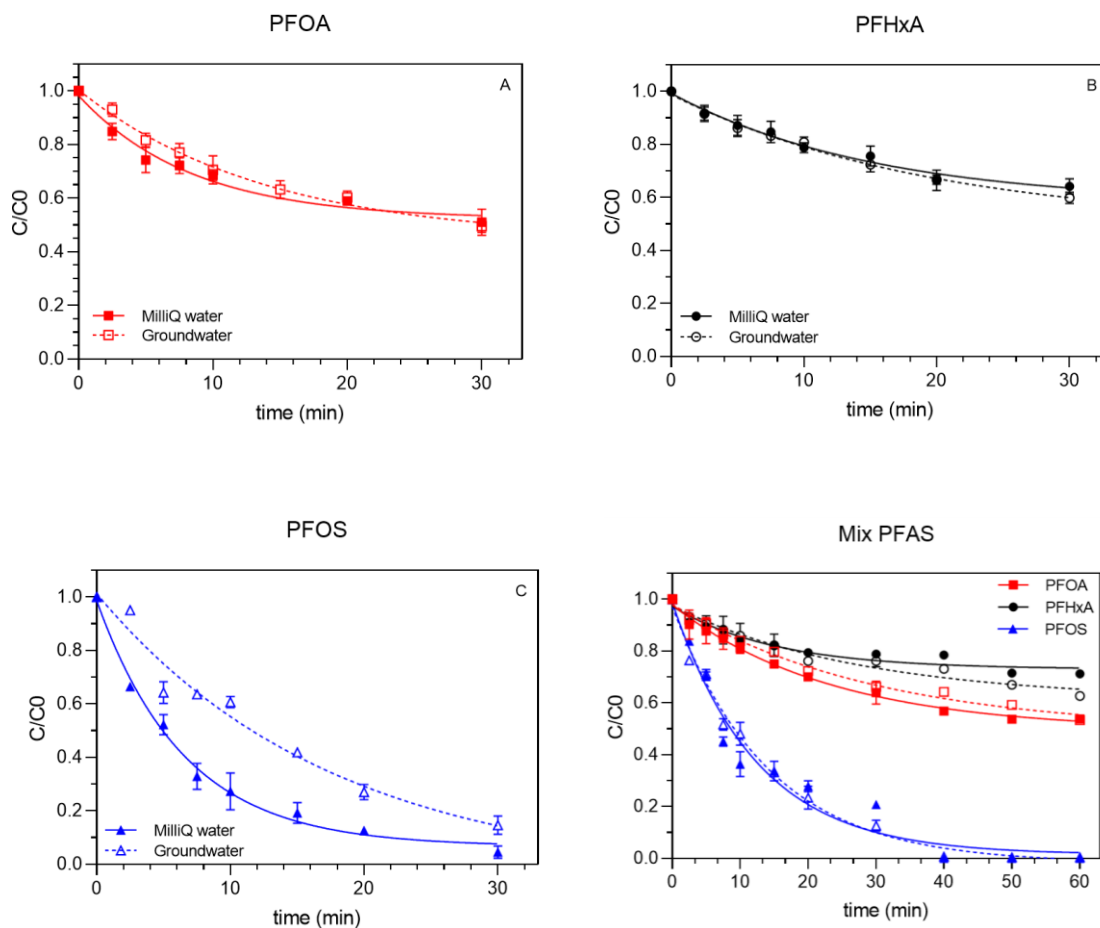


Figure 23: Degradation rate of PFOA (A), PFHxA (B) and PFOS (C) treated in individual solutions prepared in MilliQ water and groundwater matrices. Degradation of PFAS treated as mixture (D) in MilliQ (solid symbols, solid lines) and groundwater (open symbols, dashed lines).

2.2 Generation of byproducts

The identification of byproducts was challenging due to the lack of commercially available standards. However, after an HRMS analysis using the SWATH acquisition mode, identification of potential byproducts was achieved, based on mass accuracy of the measured molecular ions $[M-H]^-$ within 5 ppm, isotopic patterns, retention times and MS/MS spectra comparison between theoretical and found m/z features. A suspect screening method was followed, focusing on compounds with compatible elemental composition to PFASs. More specifically, only compounds with a maximum number of $C_8O_4F_{18}N_5H_{18}S_2$ were considered. For those that matches with online libraries were not found, *In silico* fragmentation was performed using ChemSpider.

Previous studies have shown that plasma water treatment technologies are able to degrade longer chain PFASs into short-chain Perfluorocarboxylic acids (PFCAs) and Perfluorosulfonic acids (PFSAs) [194]. The most common byproducts from PFOA and PFOS degradation are found to be the following: Perfluoroheptanoic acid (PFHpA), Perfluorohexanoic acid (PFHxA), Perfluoropentanoic acid (PFHPeA), Perfluorobutanoic acid (PFBA), Perfluorobutane sulfonate (PFBS) and Perfluorohexane sulfonate (PFHxS), with the last two being reported only for PFOS.

In this study, the above mentioned shorter chain compounds were detected as well (Table 20), confirming the results found in literature. More specifically, after treatment of PFOA's solution treatment individually, all the four previously mentioned compounds were detected. PFHpA and PFHxA were firstly detected after 2,5 minutes of treatment, while for PFPeA and PFBA the first time was after 5 minutes of treatment. A quantification study of the byproducts was not possible – mainly due to lack of analytical standards - and hence specific information about their concentrations cannot be provided. However, the byproducts' peak areas were considered in order to understand their formation rate over time. A peak area increase was observed for all four byproducts until the end of the experiment (30 minutes) (Figure 24). These results confirm PFOA's already known degradation pathway, based on the deterioration of the carbon chain: PFOA (C8) > PFHpA (C7) > PFHxA (C6) > PFPeA (C5) > PFBA (C4).

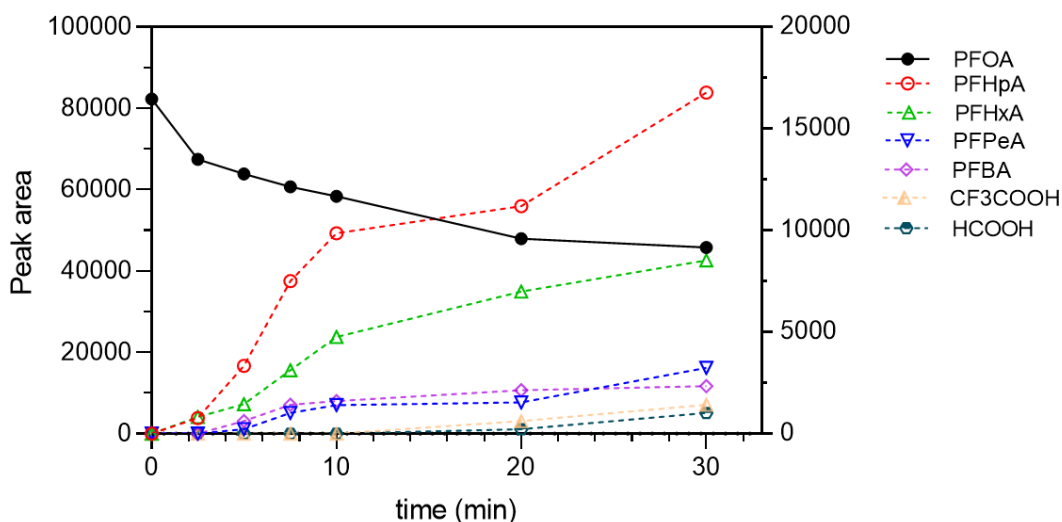


Figure 24: Generation of byproducts, after individual treatment of PFOA in MilliQ water matrix.

Concerning PFOS, the first byproduct was PFOA and was detected after 2,5 minutes, while after 5 minutes PFHpA, PFHpS and PFHxS were detected, and finally PFBA, PFPeS

and PFBS firstly occurred after 7,5 minutes. These results indicate a gradual degradation of the carbon chain as well, starting from eight atoms of carbon and arriving at four. However, PFHxA and PFPeA were not detected at all during PFOS treatment. The peak areas for all the identified byproducts were increasing with the treatment time, arriving at a highest point at 30 minutes of treatment (Figure 25). Finally, in treated solutions of both individual compounds, further mineralization of the byproducts was observed, since CF_3COOH and HCOOH were detected after 20 minutes of treatment, with their peak areas increasing until the end of the treatment time (Figure 24,25).

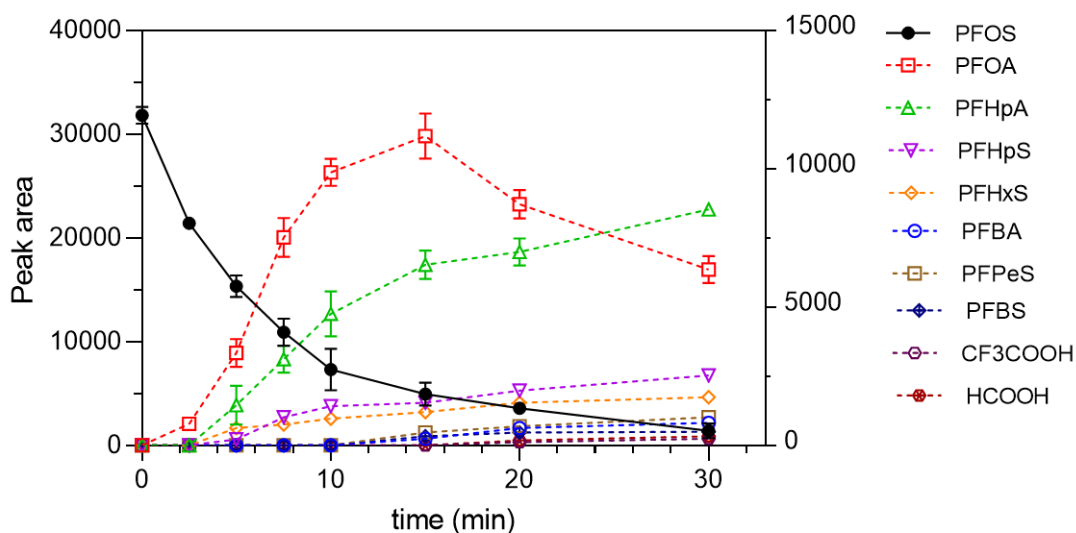


Figure 25: Generation of byproducts, after individual treatment of PFOS in MilliQ water matrix.

Concerning PFHxA - the shorter chain target molecule, which resulted to be also the most recalcitrant - PFBA and PFPeA were identified as its byproducts, molecules with carbon chains containing four and five atoms of carbon respectively. Both molecules were detected after 2,5 minutes of treatment and their peak areas were increasing with the treatment time as well. In this case CF_3COOH was detected after 20 minutes of treatment. Interestingly, the moiety $\bullet\text{C}_4\text{F}_9$ was detected in the treated PFHxA solution after the 30 minutes interval, indicating the fact that further treatment could potentially result to a higher degradation rate.

Table 20. Detected byproducts after the individual treatment of PFOA, PFOS and PFHxA, in both MilliQ water and groundwater matrices.

| Molecular formula | RT (min) | Observed <i>m/z</i> | Library Identification |
|--|-----------------|----------------------------|-------------------------------|
| PFOA byproducts | | | |
| C ₇ HF ₁₃ O ₂ | 9,91 | 363,997 | PFHpA |
| C ₆ HF ₁₁ O ₂ | 6,89 | 313,992 | PFHxA |
| C ₅ HF ₉ O ₂ | 3,96 | 263,952 | PFPeA |
| C ₄ HF ₇ O ₂ | 1,53 | 214,002 | PFBA |
| C ₂ HF ₃ O ₂ | 1,99 | 114,939 | CF ₃ COOH |
| CH ₂ O ₂ | 1,16 | 46,028 | HCOOH |
| C ₂ HO ₂ | 1,02 | 57,031 | - |
| C ₃ H ₅ F ₃ O | 4,76 | 114,029 | - |
| C ₃ HF ₇ O ₆ | 1,03 | 265,967 | - |
| C ₄ H ₅ F ₃ O ₂ | 1,09 | 142,025 | - |
| C ₅ H ₆ F ₆ | 2,64 | 180,038 | - |
| C ₅ HF ₉ O | 1,05 | 247,989 | - |
| C ₆ H ₁₁ F ₃ | 9,13 | 140,150 | - |
| C ₆ H ₂ F ₁₀ | 7,01 | 279,996 | - |
| C ₆ H ₂ F ₁₃ O ₂ | 5,03 | 352,060 | - |
| C ₆ HF ₉ O ₃ | 9,15 | 291,060 | - |
| C ₇ H ₆ F ₁₀ O | 1,03 | 296,011 | - |
| C ₇ HF ₁₅ | 9,41 | 369,986 | - |
| C ₈ H ₃ F ₉ O ₄ | 9,06 | 333,989 | - |
| C ₈ H ₅ F ₉ O ₃ | 10,38 | 320,011 | - |
| PFOS byproducts | | | |
| C ₈ HF ₁₅ O ₂ | 10,44 | 413,985 | PFOA |
| C ₇ HF ₁₃ O ₂ | 9,93 | 365,001 | PFHpA |
| C ₇ HF ₁₅ O ₃ S | 13,26 | 450,790 | PFHpS |
| C ₆ HF ₁₃ O ₃ S | 12,72 | 401,127 | PFHxS |
| C ₅ HF ₁₁ O ₃ S | 11,74 | 351,163 | PFPeS |
| C ₄ HF ₇ O ₂ | 1,53 | 215,076 | PFBA |

| Molecular formula | RT (min) | Observed <i>m/z</i> | Library Identification |
|--|----------|---------------------|------------------------|
| C ₄ HF ₉ O ₃ S | 6,62 | 301,056 | PFBS |
| C ₃ HF ₅ O ₃ S | 2,06 | 212,014 | PFPrS |
| C ₂ HF ₃ O ₂ | 1,85 | 114,522 | CF ₃ COOH |
| CH ₂ O ₂ | 1,13 | 45,998 | HCOOH |
| C ₂ HO ₂ | 1,02 | 56,998 | - |
| C ₈ H ₇ F ₁₁ O ₃ | 8,76 | 360,012 | - |
| C ₈ H ₇ F ₁₁ O ₂ S | 7,03 | 376,019 | - |
| C ₈ H ₆ F ₈ O ₃ S | 9,00 | 334,018 | - |
| C ₆ H ₂ F ₆ O ₂ S ₂ | 6,09 | 283,941 | - |
| C ₅ H ₂ F ₈ O ₃ S | 2,64 | 293,960 | - |
| C ₄ H ₅ F ₃ O ₂ | 6,05 | 142,083 | - |
| C ₃ H ₇ FO ₂ S | 3,01 | 126,015 | - |
| C ₂ H ₄ O ₄ S ₂ | 1,03 | 156,018 | - |
| PFHxA byproducts | | | |
| C ₅ HF ₉ O ₂ | 3,96 | 264,050 | PFPeA |
| C ₄ HF ₇ O ₂ | 1,53 | 215,076 | PFBA |
| C ₆ H ₆ F ₅ O ₂ | 6,87 | 205,103 | - |
| C ₅ HF ₁₁ | 8,98 | 270,004 | - |
| C ₅ H ₂ F ₈ O ₂ | 6,23 | 246,016 | - |
| C ₄ H ₅ F ₃ O ₂ | 5,32 | 142,084 | - |
| C ₂ HF ₃ O ₂ | 1,69 | 114,022 | CF ₃ COOH |
| C ₃ H ₅ F ₃ O | 3,85 | 114,071 | - |

Moieties like •C₈F₁₇, •C₇F₁₅, •C₆F₁₃, •C₅F₁₁, and •C₄F₉ were observed in all the treated samples. Their existence can be explained by the fact that plasma or aqueous electrons – which are the main species responsible for degradation in the non-thermal plasma treatment technique, and those generating the hydroxyl radical species as well- attack the -COOH functional group of PFCAs, which may transform to the more stable alkane form (C₈HF₁₇, C₇HF₁₅, C₆HF₁₃, C₅HF₁₁, C₄HF₉), or after addition of the •OH⁻ radical may result to the formation of alcohols (C₈HF₁₇O, C₇HF₁₅O, C₆HF₁₃O, C₅HF₁₁O, C₄HF₉O) (Figure 26 a,b). Thermally unstable alcohols could be transformed into their more stable ketone

forms (-C=O), after elimination of an -HF, caused by e^- attack. Furthermore, hydrolysis of the ketones could yield the formation of carboxylic acids ($C_7HF_{13}O_2$, $C_6HF_{11}O_2$, $C_5HF_9O_2$, $C_4HF_7O_2$, $C_2HF_3O_2$) with a loss of another -HF molecule. In this way, chain deterioration reactions for longer chain PFCAs (PFOA, PFHxA) result into shorter chain compounds (Figure 26). The degradation pathway of PFSA's seems to be similar to the one of PFCAs. More specifically, the degradation mechanism of PFOS initiates with the attack of electrons to the C-S bond forming the $\bullet C_8F_{17}$ moiety and the SO_3^- group. The chain propagation reactions of $\bullet C_8F_{17}$ follow the mechanisms described above, resulting to shorter chain PFCAs. Moreover, the moieties such as $\bullet C_7F_{15}$, $\bullet C_6F_{13}$, $\bullet C_5F_{11}$, and $\bullet C_4F_9$, may react with the SO_3^- group and form shorter chain of PFSA's (PFHpS, PFHxS, PFPeS, PFBS). The byproducts from all the three (common) degradation mechanisms, were further mineralized to CF_3COOH , $HCOOH$, and CO_2 . Figures 26a and 26b summarize these degradation pathways. The identified short-chain PFAS compounds were detected in both studied matrices, in both states – individual substances and their mix.

Moreover, chemical reactions of PFASs with plasma electrons, can result to the formation of many transient or stable compounds. A number of novel byproducts (Table 20) was detected after PFOA, PFOS and PFHxA degradation, with no spectral matches in the libraries. Their chemical formulas were confirmed based on accurate mass measurements and after comparison with results obtained from their *in silico* fragmentation with ChemSpider. However, such information was not adequate for building structural formulas for all the detected byproducts, due to for example isomeric patterns. These novel byproducts, showed the degradation of the strong C-F bond (Figure 26 a,b), followed by a substitution with a H atom ($C_3H_5F_3O$, $C_5H_6F_6$, $C_6H_{11}F_3$, $C_6H_2F_{10}$, $C_6H_2F_{13}$, $C_7H_6F_{10}$, $C_8H_5F_8O_3S$, $C_5H_2F_8O_3S$, $C_5H_2F_8O_3S$, $C_4H_4F_3O_2$, $C_6H_6F_5O_2$, $C_5H_2F_8O_2$, $C_4H_5F_3O_2$) providing us with satisfactory results about the bond's stability elimination, which would mean less bioaccumulative compounds in the water bodies. Analysis of total fluorine mass in the liquid phase was carried out, but the low initial concentrations of the solutions (1 $\mu g/L$) were significantly lower than the limits of detection of the method used, and thus confirmation of these results was not done.

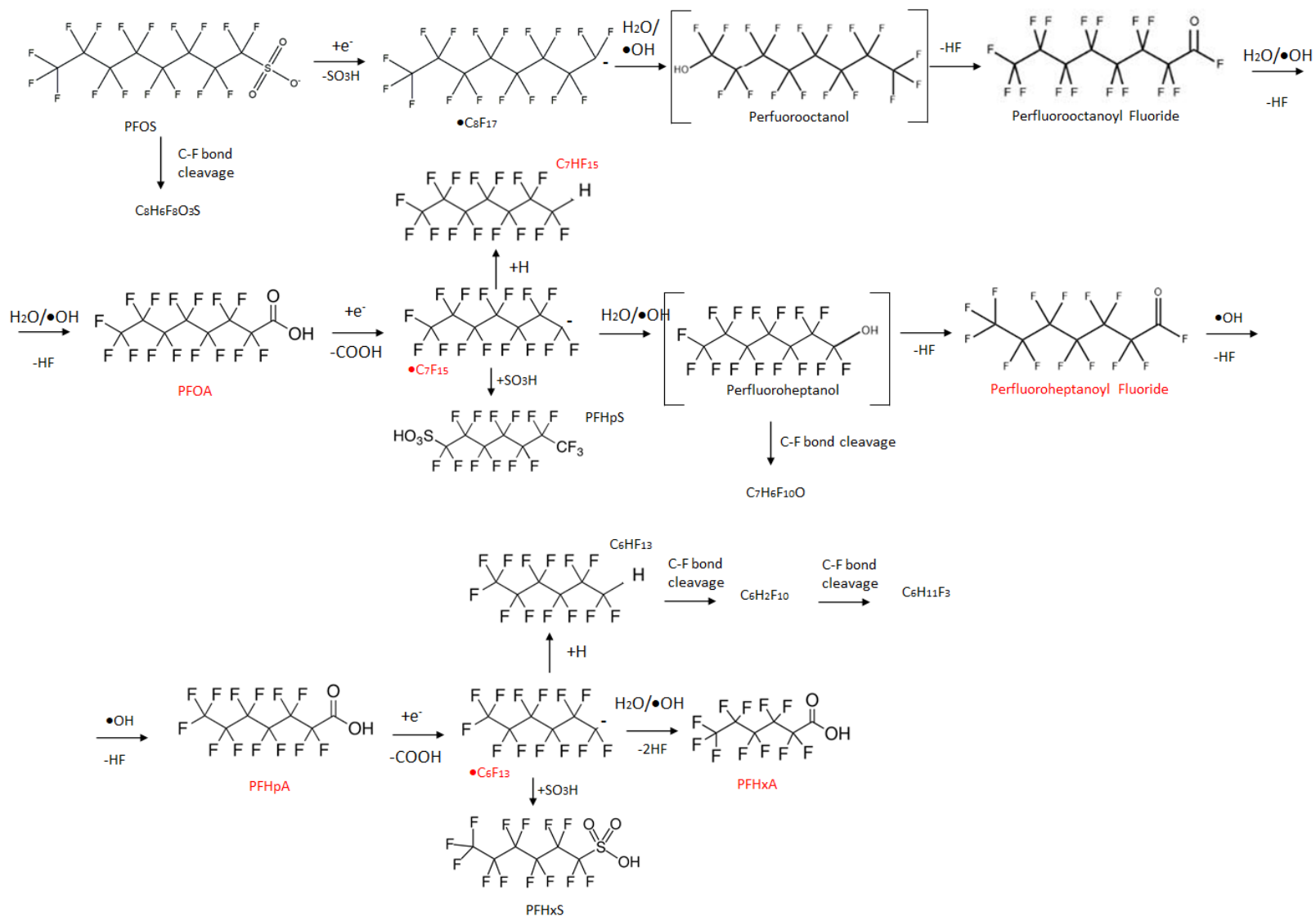


Figure 26(a): Proposed degradation mechanism of PFOS, PFOA and PFHxA, as red are marked the common byproducts, between parenthesis are the unstable byproducts.

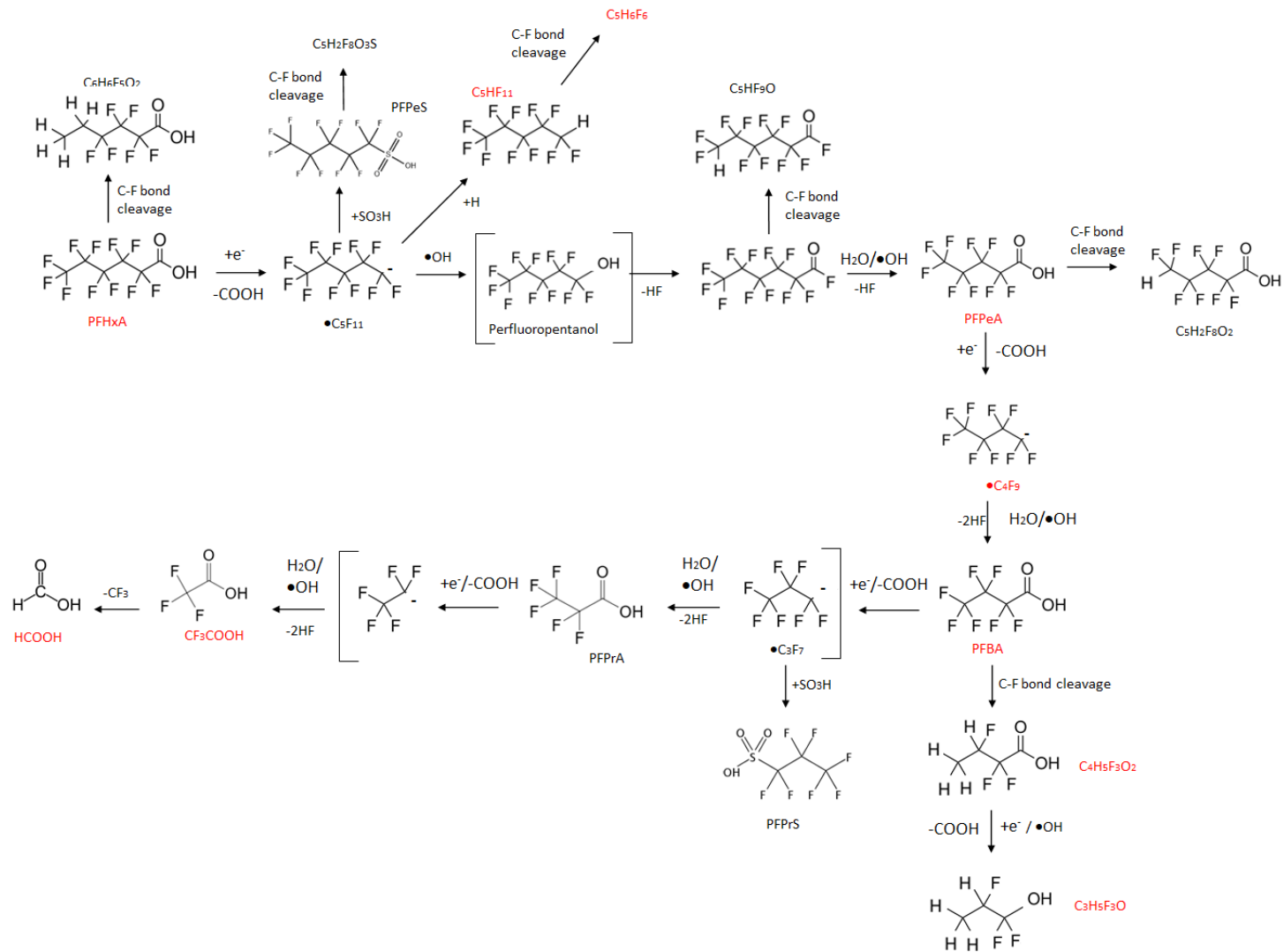


Figure 36(b): Proposed degradation mechanism of PFOS, PFOA and PFHxA, as red are marked the common byproducts, between parenthesis are the unstable byproducts.

2.3 Toxicity of byproducts

Toxicity of the samples before and after treatment was firstly evaluated using two different bioassays targeting to the crustacean *Thamnocephalus platyurus* and the bioluminescent bacteria *Alivibrio fischeri*. However, due to metals' release from the electrodes after the treatment [191], toxic effects on the two organisms after treatment was not possible to evaluate. For this reason, the potential toxicity of the 3 target analytes and their generated byproducts was predicted using the ECOSAR 2.0 program.

From the three target compounds, PFOA has been predicted as the most toxic to aquatic organisms with LC50 values of 10,1 mg/L for fish, 7,44 mg/L for Daphnid and 16,2 mg/L for Green Algae species (Table 21). The second most toxic compound was found to be PFOS, with LC50 values of 23,7 mg/L, 16,9 mg/L and 32,6 mg/L for fish, Daphnid and Green Algae species respectively (Table 21). The least toxic compound among the three target analytes was PFHxA, with LC50 values of 122 mg/L for fish, 79,3 mg/L for Daphnid and 104 mg/L for Green Algae species (Table 21). All these results are in accordance with those already published in the bibliography and are confirming the fact that shorter chain compounds have less harmful effects [46].

Moreover, a toxicity decrease was observed alongside the degradation of the carbon chain of the three target compounds, confirming once again the less harmful effects of short-chain PFAS (Table 21). In general, the toxic effects of all the detected PFASs to aquatic organisms were predicted to be lower than their equivalent PFCAs. Interestingly, the shorter chain compound that was identified in PFOA and PFOS degradation after 20 minutes of treatment – HCOOH – was predicted to have more harmful effects than CF₃COOH. Finally, concerning the novel byproducts, toxicity information was difficult to be predicted due to data limitations in literature. Optimization of the reactor parameters for eliminating metals' release and analytical standards for identifying the novel byproducts are necessary for a more comprehensive evaluation of the toxic effects before and after treatment.

Table 21. Toxicity predictions for PFOA, PFOS, PFHxA and their detected byproducts using the ECOSAR software.

| Molecular formula | Library Identification | LC50 (mg/L) Fish | LC50 (mg/L) Daphnid | EC50 (mg/L) Green Algae |
|--|-------------------------------|-------------------------|----------------------------|--------------------------------|
| C ₈ HF ₁₅ O ₂ | PFOA | 10,10 | 7,44 | 16,20 |
| C ₇ HF ₁₃ O ₂ | PFHpA | 35,40 | 24,50 | 41,40 |
| C ₆ HF ₁₁ O ₂ | PFHxA | 122 | 79,30 | 104 |
| C ₅ HF ₉ O ₂ | PFPeA | 409 | 250 | 254 |
| C ₄ HF ₇ O ₂ | PFBA | 1,32E+3 | 761 | 597 |
| C ₂ HF ₃ O ₂ | CF ₃ COOH | 2,07E+4 | 1,02E+4 | 4,31E+3 |
| CH ₂ O ₂ | HCOOH | 6,13E+3 | 2,77E+3 | 807 |
| C ₂ HO ₂ | - | 11,80 | 46,70 | 2,32 |
| C ₃ H ₅ F ₃ O | - | 1,40E+3 | 705 | 319 |
| C ₃ HF ₇ O ₆ | - | - | - | - |
| C ₄ H ₅ F ₃ O ₂ | - | 50 | 112 | 53,60 |
| C ₅ H ₆ F ₆ | - | 5,94 | 3,89 | 5,25 |
| C ₅ HF ₉ O | - | 32,80 | 394 | 283 |
| C ₆ H ₁₁ F ₃ | - | 3,40 | 2,26 | 3,22 |
| C ₆ H ₂ F ₁₀ | - | 3,79 | 2,58 | 4,06 |
| C ₆ H ₂ F ₁₃ O ₂ | - | - | - | - |
| C ₆ HF ₉ O ₃ | - | 232 | 548 | 385 |
| C ₇ H ₆ F ₁₀ O | - | - | - | - |
| C ₇ HF ₁₅ | - | 0,76 | 0,57 | 1,28 |
| C ₈ H ₃ F ₉ O ₄ | - | - | - | - |
| C ₈ H ₅ F ₉ O ₃ | - | 41,30 | 85,70 | 36,30 |
| C ₈ HF ₁₇ O ₃ S | PFOS | 23,70 | 16,90 | 32,60 |
| C ₇ HF ₁₅ O ₃ S | PFHpS | 85 | 57,10 | 85,40 |
| C ₆ HF ₁₃ O ₃ S | PFHxS | 301 | 190 | 220 |
| C ₅ HF ₁₁ O ₃ S | PFPeS | 1,05E+3 | 625 | 560 |
| C ₄ HF ₉ O ₃ S | PFBS | 3,60E+3 | 2,01E+3 | 1,40E+3 |
| C ₃ HF ₅ O ₃ S | PFPrS | - | - | - |
| C ₈ H ₇ F ₁₁ O ₃ | - | 49,90 | 30,70 | 31,70 |

| Molecular formula | Library Identification | LC50 (mg/L) Fish | LC50 (mg/L) Daphnid | EC50 (mg/L) Green Algae |
|--|-------------------------------|-------------------------|----------------------------|--------------------------------|
| C ₈ H ₇ F ₁₁ O ₂ S | - | - | - | - |
| C ₈ H ₆ F ₈ O ₃ S | - | - | - | - |
| C ₆ H ₂ F ₆ O ₂ S ₂ | - | - | - | - |
| C ₅ H ₂ F ₈ O ₃ S | - | 0,51 | 0,37 | 0,72 |
| C ₄ H ₅ F ₃ O ₂ | - | 50 | 112 | 53,60 |
| C ₃ H ₇ FO ₂ S | - | 8,37 | 64,70 | 54,30 |
| C ₂ H ₄ O ₄ S ₂ | - | 502 | 1,34E+3 | 832 |
| C ₆ H ₆ F ₅ O ₂ | - | - | - | - |
| C ₅ HF ₁₁ | - | 8,91 | 5,84 | 7,88 |
| C ₅ H ₂ F ₈ O ₂ | - | 1,34E+3 | 775 | 623 |
| C ₄ H ₅ F ₃ O ₂ | - | 50 | 112 | 53,60 |
| C ₃ H ₅ F ₃ O | - | 1,20E+3 | 610 | 283 |

3. Conclusions

In this chapter, the degradation of PFOA, PFOS and PFHxA, as well as their mix, in different water matrices after treatment with a non-thermal plasma generator is reported. The compounds were chosen according to their occurrence levels in groundwater sources from the Metropolitan Area of Turin. The samples were prepared in trace level concentrations, in order to examine the efficiency of this technique in realistic conditions. The best results were obtained for PFOS in both matrices, while PFHxA showed the lowest removal. Identification of byproducts was done with LC-HRMS analysis. In summary, the identified molecules showed a reduction of the carbon chain, confirming the mechanisms already reported in literature obtained after the employment of other AOP techniques. *In silico* prediction of the toxicity before and after treatment showed that the generated byproducts have less harmful effects on aquatic organisms. Moreover, in this study, defluorination of the compounds was observed as well, highlighting the ability of this technique to break one of the strongest chemical bonds.

The results obtained for the tested compounds, significantly assisted in understanding the degradation of PFASs after the non-thermal plasma treatment, and suggest that this technology could efficiently be used for the removal of PFASs found in water sources used for drinking water production. Moreover, as the degradation results were satisfactory for one of the most recalcitrant classes of CECs, this technique could be efficient also on the removal of other contaminants. Further studies in order to understand if a large scale application of this technique, for example after the granulated activated carbon (GAC) filters in a DWTP line where the concentrations of these molecules are extremely low, are needed. A human health risk assessment for the existence of these compounds in drinking water sources was not carried out, since lack of information about the effects these compounds on human health can have. Further investigation in this field is necessary.

Section V

Conclusions

This thesis is part of the AQUALity-ETN project, which aims to find solutions for the removal of Contaminants of Emerging Concern, present in water bodies in trace level concentrations. The main research activities of the individual project were carried out in the Research Center of Società Metropolitana Acque Torino, while secondment periods at the Chemistry and Biology Department of Aalborg University, Denmark and the Chemistry Department of Turin University and further collaborations within the consortium contributed to its conclusion.

The main goal of this thesis was to develop green and without waste generation, advanced analytical tools for evaluating the presence of CECs in different water sources and improve conventional monitoring approaches which resulted insufficient in evaluating water bodies' quality, towards the adoption of a circular economy approach. In order to take control measures for a safe and sustainable water supply, it is important to evaluate the holistic quality of water sources and identify the hazardous components, their occurrence areas, and the points at higher contamination risk. We achieved this goal by combining target, suspect and non-target chemical analyses with risk-based approaches and toxicological assessments.

Firstly, a new green, fast and cost-efficient method – following the principles of Green Analytical Chemistry - with high sensitivity in detecting a mix of sixteen different PFAS compounds in drinking water samples at trace level concentrations was developed and validated. The key characteristic of this method, was the absence of an extraction step and a direct injection into the analytical system. This method was used in an assessment of PFASs occurrence levels in the Metropolitan Area of Turin in Italy. A correlation of the “positive” sampling points and the potential pollution sources in the territory based on multivariate and spatial statistical tools, was done in order to understand their influence on the pollution levels and take decisions for reduction of contamination at source. The results showed that the number of point sources within a watershed significantly affects PFASs occurrence levels, providing us with significant predictors for guiding future choice of sampling points at higher risk.

Based on that “smart” monitoring tool and in order to avoid the costs, efforts and environmental impact of large-scale, blind monitoring assessments, a prioritization of the sites at major risks of pollution with pharmaceuticals and hormones was done for the second monitoring assessment in the study area included in this thesis. A new method following the principles of Green Analytical Chemistry as well, targeting at sixteen compounds was developed achieving low Quantification Limits for every analyte,

and validated. The method was applied in the analyses of raw water samples taken from the areas at higher risks, including surface, groundwater and treated water. The results confirmed the presence of the target compounds in the area, in concentrations of the ng/L scale, with those geographically closer to the considered pollution sources showing higher detection rates. Analysis of the samples after the treatment line showed sufficient removal of the target pollutants, minimizing human health risks and a risk assessment study was then carried out in order to evaluate the potential effects on human health, taking into account the cocktail effects of the compounds' occurrence in mixtures.

However, target analyses alone, have been proved insufficient in estimating the total quality of water bodies. For this reason, in this thesis we included a non-target screening of surface samples collected from three different locations in Greece and Italy. High resolution Mass Spectrometry using the SWATH-MS acquisition mode showed the occurrence of a vast variety of pollutants, mainly pharmaceuticals, pesticides and herbicides, PFAS and personal care products. Multivariate statistical analysis tools were used in order to identify pollution patterns and similarities among the samples, as well as to identify the compounds that are responsible for the discrimination among the samples. The use of non-target screening highlighted the need of including it as a first step in monitoring assessments, in order to evaluate the quality of water sources, prioritize the contaminants to be included in quantification studies and take decisions for treatment needs.

On the second part of this thesis, based on the results obtained from the three monitoring assessments, the removal efficiency of different degradation methods was studied. Towards a sustainable and toxic-free environment the implementation of fast, cost-efficient and environmental friendly techniques that don't generate waste or more toxic byproducts, are necessary. In this way, Advanced Oxidation Processes (AOPs) were studied as degradation techniques, within the context of Green Chemistry, taking into account byproducts generation and toxicity effects.

More specifically, first the effects of UV-A, UV-B, and UV-C irradiation on the degradation of glyphosate were studied. Different parameters that could affect the process were taken into account and their effects on toxicity to aquatic organisms from different trophic levels were studied. The effect-based monitoring approach for the evaluation of the method's efficiency was combined with chemical analyses in order to identify transformation byproducts. Toxicity assays were used as an important supplement to chemical analyses in order to assess water quality, as bioassays can integrate changes in

water chemistry and bioactivity before and after water treatment. The results suggested that UV-C and to some extent UV-B photolysis of glyphosate in water could decrease concentrations of this pesticide and reduce overall ecotoxicity by generating less toxic transformation products. Moreover, even if UV doses used in treatment plants are mainly used for disinfection processes, being not adequate to degrade organic contaminants, UV photolysis doesn't generate waste and could be a perfect candidate for implementation in WWTPs and DWTPs.

Finally, the degradation of PFOA, PFOS and PFHxA, as well as their mix, in different water matrices after treatment with a non-thermal plasma generator was studied. This technique basically applies on one or several very high voltage (HV) pulses of very short duration to a reactor containing the contaminated water samples, generating in this way pressure waves, UV light and formation of chemically active species such as $\bullet\text{OH}$, $\bullet\text{H}$, $\bullet\text{O}$, $\bullet\text{O}_2^-$, $\bullet\text{HO}_2$, $\bullet\text{H}_2\text{O}_2$, $\bullet\text{O}_3$ that can break the organic molecules. The results showed degradation of the compounds after 30 minutes of treatment individually, and 60 minutes as a mix. Identification of the byproducts was done with HRMS analysis and defluorination of the compounds was observed as well, highlighting the ability of this method to break one of the strongest chemical bonds. Toxicity before and after treatment was predicted with the ECOSAR software and showed less harmful effects on aquatic organisms alongside with reduction of the carbon chain.

In conclusion, moving towards a circular economy concept, a more sustainable water management is necessary, including access to tools for identification and detection of potentially hazardous compounds and efficient treatment techniques that don't generate additional waste, with the upper aim of water reuse. The results presented in this thesis highlight the importance of improving water quality monitoring assessments. Target analyses alone are not able to sufficiently evaluate the pollution of water bodies. Combination with non-target and risk-based approaches is fundamental in order to evaluate water bodies status more comprehensively. In the future, in order to take control measures for a safe and sustainable water supply, is important to identify the hazardous components, their occurrence areas and the points at higher contamination risk, including toxicological methods in order to minimize risk to human health and ecosystems.

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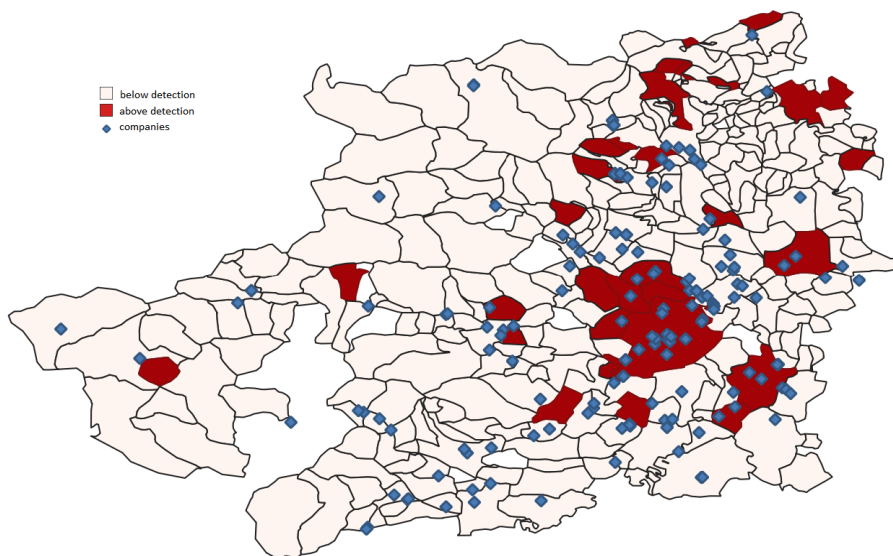
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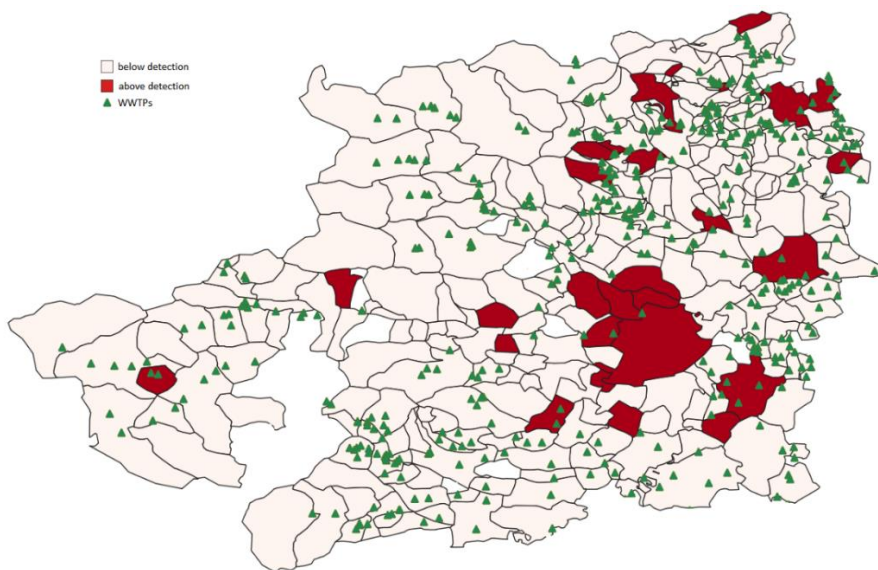
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Appendix



(a)



(b)

Figure A1: Map of the pollution levels of PFAS as a sum (limit 10 ng L^{-1}) and the selected point sources present in the studied area: (a) industrial sites and (b) waste water treatment plants, Chapter 3.

Table A1. List of the municipalities in the Metropolitan Area of Turin, Italy included in the studies reported in Chapters 3 and 4

| Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID |
|----------------------|-----------|------------------------|-----------|----------------------|-----------|------------------------|-----------|
| AGLIE' | 1 | CHIVASSO | 74 | MONCALIERI | 147 | SAN CARLO CANAVESE | 220 |
| AIRASCA | 2 | CICONIO | 75 | MONCENISIO | 148 | SAN COLOMBANO BELMONTE | 221 |
| ALA DI STURA | 3 | CINTANO | 76 | MONTALDO TORINESE | 149 | SAN DIDERO | 222 |
| ALBIANO D'IVREA | 4 | CINZANO | 77 | MONTALENGHE | 150 | SAN FRANCESCO AL CAMPO | 223 |
| ALICE SUPERIORE | 5 | CIRIE' | 78 | MONTALTO DORA | 151 | SAN GERMANO CHISONE | 224 |
| ALMESE | 6 | CLAVIERE | 79 | MONTANARO | 152 | SAN GILLIO | 225 |
| ALPETTE | 7 | COASSOLO TORINESE | 80 | NICHELINO | 153 | SAN GIORGIO CANAVESE | 226 |
| ANDEZENO | 8 | COAZZE | 81 | NOASCA | 154 | SAN GIORIO DI SUSÀ | 227 |
| ANDRATE | 9 | COLLEGNO | 82 | NOLE | 155 | SAN GIUSTO CANAVESE | 228 |
| ANGROGNA | 10 | COLLERETTO CASTELNUOVO | 83 | NOMAGLIO | 156 | SAN MARTINO CANAVESE | 229 |
| ARIGNANO | 11 | COLLERETTO GIACOSA | 84 | NONE | 157 | SAN MAURIZIO CANAVESE | 230 |
| AVIGLIANA | 12 | CONDOVE | 85 | NOVALESA | 158 | SAN MAURO TORINESE | 231 |

| Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID |
|---------------------------------|----------------------|---------------------------------|----------------------|---------------------------------|----------------------|---------------------------------|----------------------|
| BAIRO | 13 | CORIO | 86 | OGLIANICO | 159 | SAN PIETRO VAL LEMINA | 232 |
| BALANGERO | 14 | COSSANO CANAVESE | 87 | ORBASSANO | 160 | SAN PONSO | 233 |
| BALDISSERO CANAVESE | 15 | CUCEGLIO | 88 | ORIO CANAVESE | 161 | SAN RAFFAELE CIMENA | 234 |
| BALDISSERO TORINESE | 16 | CUMIANA | 89 | OSASCO | 162 | SAN SEBASTIANO DA PO | 235 |
| BALME | 17 | CUORGNE' | 90 | OSASIO | 163 | SAN SECONDO DI PINEROLO | 236 |
| BANCHETTE | 18 | DRUENTO | 91 | OULX | 164 | SANGANO | 237 |
| BARBANIA | 19 | EXILLES | 92 | OZEGNA | 165 | SANT'AMBROGIO DI TORINO | 238 |
| BARDONECCHIA | 20 | FAVRIA | 93 | PANCALIERI | 166 | SANT'ANTONINO DI SUSA | 239 |
| BARONE CANAVESE | 21 | FELETTO | 94 | PARELLA | 167 | SANTENA | 240 |
| BEINASCO | 22 | FIANO | 95 | PAVAROLO | 168 | SAUZE DI CESANA | 241 |
| BIBIANA | 23 | FIORANO CANAVESE | 96 | PAVONE CANAVESE | 169 | SAUZE D'OULX | 242 |
| BOBBIO PELLICE | 24 | FOGLIZZO | 97 | PECCO | 170 | SCALENGHE | 243 |
| BOLLENGO | 25 | FORNO CANAVESE | 98 | PECETTO TORINESE | 171 | SCARMAGNO | 244 |
| BORGARO TORINESE | 26 | FRASSINETTO | 99 | PEROSA ARGENTINA | 172 | SCIOLZE | 245 |
| BORGIALLO | 27 | FRONT | 100 | PEROSA CANAVESE | 173 | SESTRIERE | 246 |

| Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID |
|-----------------------------|------------------|-----------------------------|------------------|-----------------------------|------------------|-----------------------------|------------------|
| BORGOFRANCO D'IVREA | 28 | FROSSASCO | 101 | PERTUSIO | 174 | SETTIMO ROTTARO | 247 |
| BORGOMASINO | 29 | GARZIGLIANA | 102 | PESSINETTO | 175 | SETTIMO TORINESE | 248 |
| BORGONE SUSA | 30 | GASSINO TORINESE | 103 | PIANEZZA | 176 | SETTIMO VITTONI | 249 |
| BOSCONERO | 31 | GERMAGNANO | 104 | PINASCA | 177 | SPARONE | 250 |
| BRANDIZZO | 32 | GIAGLIONE | 105 | PINEROLO | 178 | STRAMBINO | 251 |
| BRICHERASIO | 33 | GIAVENO | 106 | PINO TORINESE | 179 | SUSA | 252 |
| BROSSO | 34 | GIVOLETTO | 107 | PIOBESI TORINESE | 180 | TAVAGNASCO | 253 |
| BRUINO | 35 | GRAVERE | 108 | PIOSSASCO | 181 | TORINO | 254 |
| BURIASCO | 36 | GROSCAVALLO | 109 | PISCINA | 182 | TORRAZZA PIEMONTE | 255 |
| BUSANO | 37 | GROSSO | 110 | POIRINO | 183 | TORRE CANAVESE | 256 |
| BUSSOLENO | 38 | GRUGLIASCO | 111 | POMARETTO | 184 | TORRE PELLICE | 257 |
| BUTTIGLIERA ALTA | 39 | INGRIA | 112 | PONT CANAVESE | 185 | TRANA | 258 |
| CAFASSE | 40 | INVERSO PINASCA | 113 | PORTE | 186 | TRAUSELLA | 259 |
| CALUSO | 41 | ISOLABELLA | 114 | PRAGELATO | 187 | TRAVERSELLA | 260 |
| CAMBIANO | 42 | ISSIGLIO | 115 | PRALORMO | 188 | TROFARELLO | 261 |
| CAMPIGLIONE FENILE | 43 | IVREA | 116 | PRAMOLLO | 189 | USSEAU | 262 |
| CANDIA CANAVESE | 44 | LA CASSA | 117 | PRAROSTINO | 190 | USSEGLIO | 263 |
| CANDIOLO | 45 | LA LOGGIA | 118 | PRASCORSANO | 191 | VAIE | 264 |
| CANISCHIO | 46 | LANZO TORINESE | 119 | PRATIGLIONE | 192 | VAL DELLA TORRE | 265 |
| CANTALUPA | 47 | LEINI' | 120 | QUAGLIUZZO | 193 | VALGIOIE | 266 |
| CANTOIRA | 48 | LEMIE | 121 | QUASSOLO | 194 | VALPERGA | 267 |
| CAPRIE | 49 | LESSOLO | 122 | QUINCINETTO | 195 | VAUDA CANAVESE | 268 |

| Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID |
|-----------------------------|------------------|-----------------------------|------------------|-----------------------------|------------------|-----------------------------|------------------|
| CARAVINO | 50 | LEVONE | 123 | REANO | 196 | VENARIA REALE | 269 |
| CAREMA | 51 | LOCANA | 124 | RIBORDONE | 197 | VENAUS | 270 |
| CARIGNANO | 52 | LOMBARDORE | 125 | RIVA PRESSO CHIERI | 198 | VEROLENGO | 271 |
| CARMAGNOLA | 53 | LOMBRIASCO | 126 | RIVALBA | 199 | VESTIGNE' | 272 |
| CASALBORGONE | 54 | LORANZE' | 127 | RIVALTA DI TORINO | 200 | VIALFRE' | 273 |
| CASCINETTE D'IVREA | 55 | LUGNACCO | 128 | RIVARA | 201 | VICO CANAVESE | 274 |
| CASELETTE | 56 | LUSERNA SAN GIOVANNI | 129 | RIVAROLO CANAVESE | 202 | VIDRACCO | 275 |
| CASELLE TORINESE | 57 | LUSERNETTA | 130 | RIVAROSSA | 203 | VIGONE | 276 |
| CASTAGNETO PO | 58 | LUSIGLIE' | 131 | RIVOLI | 204 | VILLAFRANCA PIEMONTE | 277 |
| CASTAGNOLE PIEMONTE | 59 | MACELLO | 132 | ROBASSOMERO | 205 | VILLANOVA CANAVESE | 278 |
| CASTELLAMONTE | 60 | MAGLIONE | 133 | ROCCA CANAVESE | 206 | VILLAR DORA | 279 |
| CASTELNUOVO NIGRA | 61 | MAPPANO | 134 | ROLETTO | 207 | VILLAR PELLICE | 280 |
| CASTIGLIONE TORINESE | 62 | MARENTINO | 135 | ROMANO CANAVESE | 208 | VILLAR PEROSA | 281 |
| CAVOUR | 63 | MASSELLO | 136 | RONCO CANAVESE | 209 | VILLARBASSE | 282 |
| CERCENASCO | 64 | MATHI | 137 | RONDISSONE | 210 | VILLAREGGIA | 283 |
| CERES | 65 | MATTIE | 138 | RORA' | 211 | VILLASTELLONE | 284 |
| CERESOLE REALE | 66 | MAZZE' | 139 | ROSTA | 212 | VINOVO | 285 |

| Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID | Municipality Name ID | Number ID |
|-----------------------------|------------------|-----------------------------|------------------|-----------------------------|------------------|-----------------------------|------------------|
| CESANA TORINESE | 67 | MEANA DI SUSÀ | 140 | RUBIANA | 213 | VIRLE PIEMONTE | 286 |
| CHIALAMBERTO | 68 | MERCENASCO | 141 | RUEGLIO | 214 | VISCHE | 287 |
| CHIANOCCO | 69 | MEUGLIANO | 142 | SALASSA | 215 | VISTRORIO | 288 |
| CHIERI | 70 | MEZZENILE | 143 | SALBERTRAND | 216 | VIU' | 289 |
| CHIESANUOVA | 71 | MOMBELLO DI TORINO | 144 | SALERANO CANAVESE | 217 | VOLPIANO | 290 |
| CHIOMONTE | 72 | MOMPANTERO | 145 | SAMONE | 218 | VOLVERA | 291 |
| CHIUSA DI SAN MICHELE | 73 | MONASTERO DI LANZO | 146 | SAN BENIGNO CANAVESE | 219 | | |

Table A2. LC Gradient conditions of the method reported in Chapter 3.

| Time (min) | Flow Rate (mL/min) | A % | B % |
|------------|--------------------|-----|-----|
| 0,00 | 0,550 | 98 | 2 |
| 0,00 | 0,550 | 98 | 2 |
| 0,50 | 0,550 | 98 | 2 |
| 1,00 | 0,550 | 70 | 30 |
| 6,00 | 0,550 | 0 | 100 |
| 7,50 | 0,550 | 0 | 100 |
| 7,60 | 0,550 | 98 | 2 |
| 10,00 | 0,550 | 98 | 2 |

Table A3. Electrospray Ionization Mode (ESI) source parameters of the method reported in Chapter 3.

| Parameter | Value |
|------------------|----------|
| Polarity | Negative |
| Curtains Gas | 30 psi |
| Collision Gas | 30 psi |
| Ionspray Voltage | -4500 V |
| Temperature | 350 °C |
| GS1 | 50 psi |
| GS2 | 55 psi |

Table A4. Multiple Reaction Monitoring (MRM) transitions and the retention time (RT) for analytes and internal standards included in the method reported in Chapter 3.

| Compound | Q1 m/z | Q3 m/z | RT (min) |
|----------|--------|--------|----------|
| PFBA | 213 | 169 | 2,10 |
| PFPeA | 263 | 219 | 3,10 |
| PFHxA | 131 | 269 | 3,80 |
| PFHpA | 363 | 319 | 4,30 |
| PFOA | 413 | 369 | 4,60 |
| PFNA | 463 | 419 | 5,00 |
| PFDA | 513 | 469 | 6,90 |
| PFUdA | 563 | 519 | 5,50 |
| PFDoA | 613 | 569 | 5,70 |
| PFTrDA | 663 | 619 | 5,90 |
| PFTeDA | 713 | 669 | 6,00 |
| PFHxDA | 813 | 769 | 6,30 |
| PFODA | 913 | 869 | 6,50 |
| L-PFBS | 299 | 99 | 3,30 |
| L-PFHxS | 399 | 99 | 4,30 |
| L-PFOS | 499 | 99 | 5,00 |
| L-PFDS | 599 | 99 | 5,40 |
| MPFHxS | 403 | 10 | 4,30 |
| MPFOS | 503 | 99 | 5,00 |
| MPFBA | 217 | 172 | 2,10 |
| MPFHxA | 315 | 270 | 3,80 |
| MPFOA | 417 | 372 | 4,60 |
| MPFNA | 468 | 423 | 4,90 |
| MPFDA | 515 | 470 | 5,20 |
| MPFUdA | 565 | 520 | 5,50 |
| MPFDoA | 615 | 570 | 5,70 |

Table A5. MRM transitions and retention time (RT) for the target compounds included in the method reported in Chapter 4.

| Compound | Ionization mode | Q1 m/z | Q3 m/z | RT (min) |
|-------------------|-----------------|----------------------------|----------------------------|----------|
| Atenolol | ESI (+) | 266,80 266,80 | 145,00 190,00 | 5,29 |
| Azithromycin | ESI (+) | 749,50 749,50 749,50 | 591,30 158,10 573,30 | 6,51 |
| Caffeine | ESI (+) | 195,10 195,10 195,10 | 138,10 110,00 123,00 | 6,37 |
| Carbamazepine | ESI (+) | 237,10 237,10 | 194,10 191,80 | 7,78 |
| Erythromycin | ESI (+) | 734,50 734,50 | 158,30 83,00 | 7,33 |
| Sulfamethoxazole | ESI (+) | 254,20 254,20 254,20 | 155,90 107,90 146,90 | 6,52 |
| Trimethoprim | ESI (+) | 291,10 291,10 | 230,20 275,00 | 5,88 |
| Clarithromycin | ESI (+) | 748,50 748,50 748,50 | 83,00 116,10 590,20 | 7,56 |
| Ketoprofen | ESI (+) | 255,10 255,10 | 105,00 77,10 | 8,13 |
| Diclofenac | ESI (+) | 296,00 296,00 | 215,00 215,00 | 8,71 |
| Ofloxacin | ESI (+) | 362,40 362,40 362,40 | 318,00 261,00 344,00 | 6,09 |
| Ciprofloxacin | ESI (+) | 332,20 332,20 332,20 | 314,00 231,00 288,30 | 6,27 |
| Cyclophosphamide | ESI (+) | 261,30 261,30 | 139,80 106,00 | 7,37 |
| 17-beta estradiol | ESI (-) | 271,10 | 145,10 | 6,51 |
| Estrone | ESI (-) | 269,10 | 145,20 | 6,48 |
| Ibuprofen | ESI (-) | 205,20 205,20 | 161,20 177,00 | 6,88 |

Table A6. Details of the samples used in the study of Chapter 5.

| No. | Sample code | Lake/River (Country) | Sampling date | Site Location GPS | Environ. Condition | Description |
|-----|-------------|----------------------|---------------|--------------------------|--------------------|----------------------------|
| 1 | S1 | Orta (Italy) | 9/5/2020 | 45,7774133, 8,4077208 | Cloud cover | Lake beach |
| 2 | S2 | Orta (Italy) | 9/5/2020 | 45,8733481, 8,4075652 | Rainy | Lake pier |
| 3 | S3 | Orta (Italy) | 9/5/2020 | 45,8753988, 8,4092584 | Rainy | Near Toce river |
| 4 | S4 | Orta (Italy) | 9/5/2020 | 45,8744660, 8,4101274 | Cloud cover | Near Hospital |
| 5 | S5 | Orta (Italy) | 9/5/2020 | 45,7948270, 8,4156655 | Cloud cover | Lake beach |
| 6 | S6 | Comabbio (Italy) | 9/5/2020 | 45,7612867, 8,6801196 | Cloud cover | Lake beach |
| 7 | S7 | Comabbio (Italy) | 9/5/2020 | 45,7772281, 8,6868874 | Cloud cover | Lake pier |
| 8 | S8 | Comabbio (Italy) | 9/5/2020 | 45,7789047, 8,6967841 | Cloud cover | Lake beach |
| 9 | S9 | Comabbio (Italy) | 9/5/2020 | 45,7721398, 8,7001449 | Cloud cover | Lake beach |
| 10 | S10 | Comabbio (Italy) | 9/5/2020 | 45,7673782, 8,7020929 | Cloud cover | Lake beach |
| 11 | S11 | Po (Italy) | 7/31/2020 | 45,02359, 7,40508 | Sunny (~28 °C) | Near Hospital |
| 12 | S12 | Po (Italy) | 9/17/2020 | 45,02359, 7,40508 | Sunny (~26 °C) | Near Hospital |
| 13 | S13 | Pamvotis (Greece) | 9/5/2020 | 39,633149, 20,898402 | Cloud cover | Lake side, bridge |
| 14 | S14 | Pamvotis (Greece) | 9/5/2020 | 39,673641, 20,858234 | Cloud cover | Lake side, touristic area |
| 15 | S15 | Pamvotis (Greece) | 9/5/2020 | 39,689113, 20,840971 | Cloud cover | Near a bridge |
| 16 | S16 | Pamvotis (Greece) | 9/5/2020 | 39,683082, 20,878095 | Cloud cover | Lake side, recreation area |
| 17 | S17 | Pamvotis (Greece) | 9/5/2020 | 39,677277, 20,907347 | Cloud cover | Lake side |

Table A7. Details about the standard compounds that were used as suspect analytes in Chapter 5.

| Compound | CAS Number | Compound | CAS Number | |
|-------------------------|------------|----------------------|------------|------------|
| Dichlorvos | CUS 19925 | Epoxiconazole | CUS 18511 | |
| Methyl parathion | | Ethofumesate | | |
| Parathion (ethyl) | | Fenamidone | | |
| Penconazole | | Fenbuconazole | | |
| Pendimethalin | | Fenhexamid | | |
| Pethoxamid | | Hexazinone | | |
| Phosalone | | Indoxacarb | | |
| Pirimicarb | | Kresoxim-methyl | | |
| Prochloraz | | Lenacil | | |
| Procymidone | | Mepanipyrim | | |
| Prometryn | | Alachlor | | CUS 12525 |
| Pronamide | | Ametryn | | |
| Propachlor | | Atrazine | | |
| Propazine | | Atrazine-desethyl | | |
| Propiconazole | | Cyanazine | | |
| Pyraclostrobin | | Metolachlor | | |
| Pyrimethanil | | Molinate | | |
| Simazine | | Oxadiazon | | |
| Spirotetramat | | Prometryn | | |
| Spiroxamine | | Propazine | | |
| Tebufenozide | | Simazine | | |
| Terbuthylazine | | Terbuthylazine | | |
| Terbuthylazine-desethyl | | Chlorpyriphos | CUS17405 | |
| Tetraconazole | | Chlorpyriphos-methyl | | |
| Thiacloprid | | Diazinon | | |
| Thiamethoxam | | Pendimethalin | | |
| Thiobencarb | | Clarithromycin | 81103-11-9 | |
| Tolylfluanid | | Erythromycin | 114-07-8 | |
| Trifloxystrobin | | Caffeine | 58-05-2 | |
| Trifluralin | | Carbamazepine | 298-46-4 | |
| Triticonazole | | Ciprofloxacin | 85721-33-1 | |
| Vinclozolin | | Ofloxacin | 82419-36-1 | |
| Zoxamide | | Cyclophosphamide | 50-18-0 | |
| Atenolol | | 29122-68-7 | Diclofenac | 15307-79-6 |
| Azithromycin | 83905-01-5 | Ketoprofen | 22071-15-4 | |

| Compound | CAS Number | Compound | CAS Number |
|-------------------|-------------------|-----------------|-------------------|
| Ibuprofen | 15687-27-1 | Estrone | 53-16-7 |
| Sulfamethoxazole | 723-46-6 | PFOA | 307-24-4 |
| Trimethoprim | 738-70-5 | PFOS | 1763-23-1 |
| 17-beta Estradiol | 50-28-2 | PFHxA | 335-67-1 |

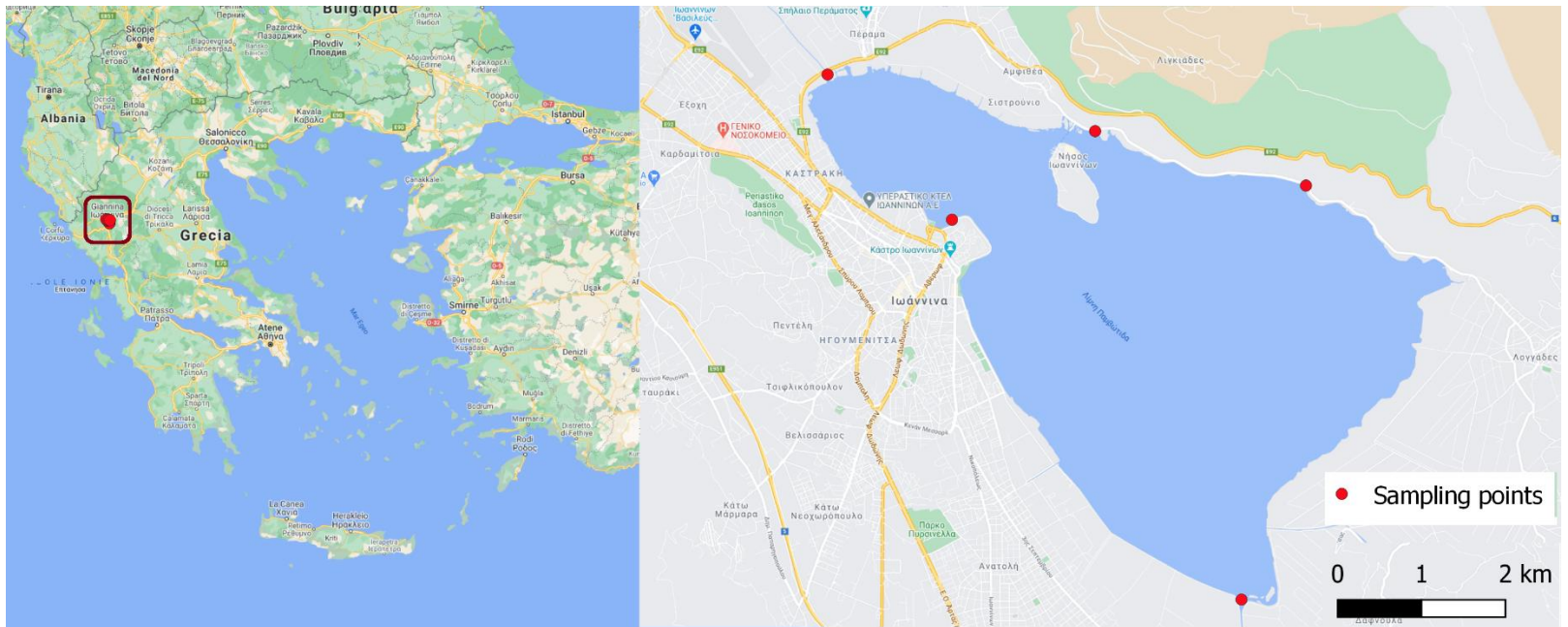


Figure A2a: A map of the sampling points from the lake Pamvotis, in Ioannina, Greece, included in Chapter 5, and its position on the national territory.

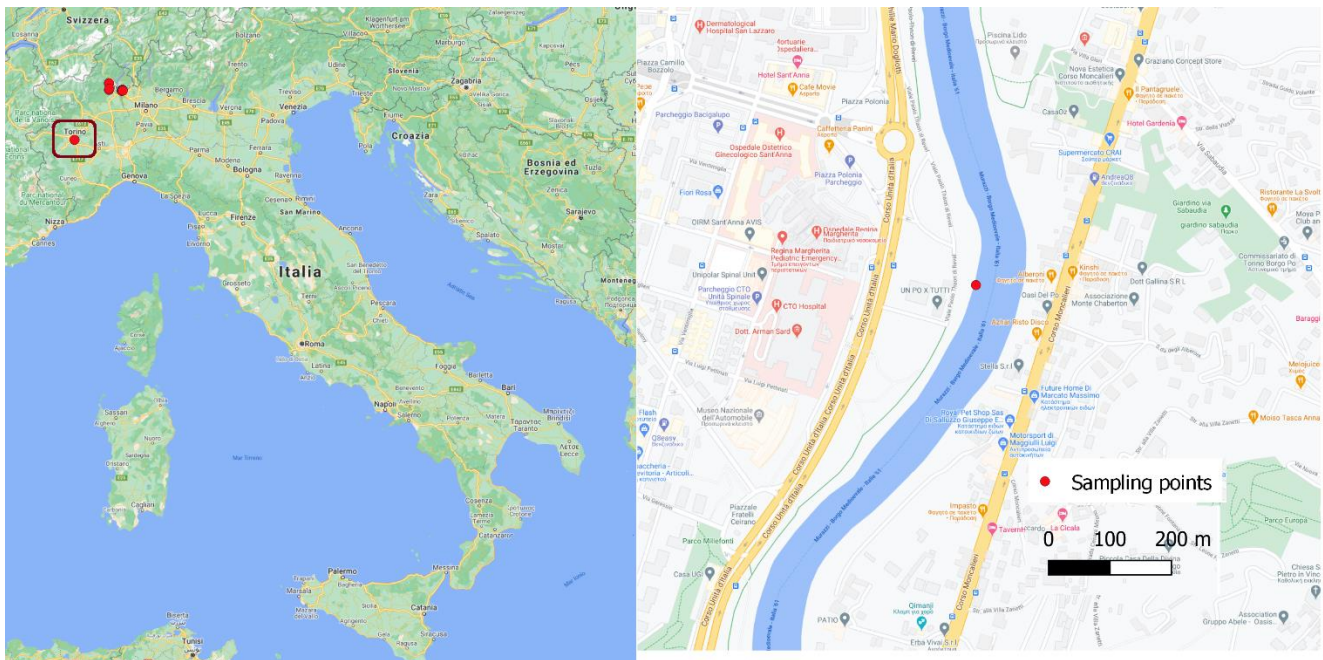


Figure A2b: A map of the sampling point from the Po river, in Turin, Italy, included in Chapter 5, and its position on the national territory.

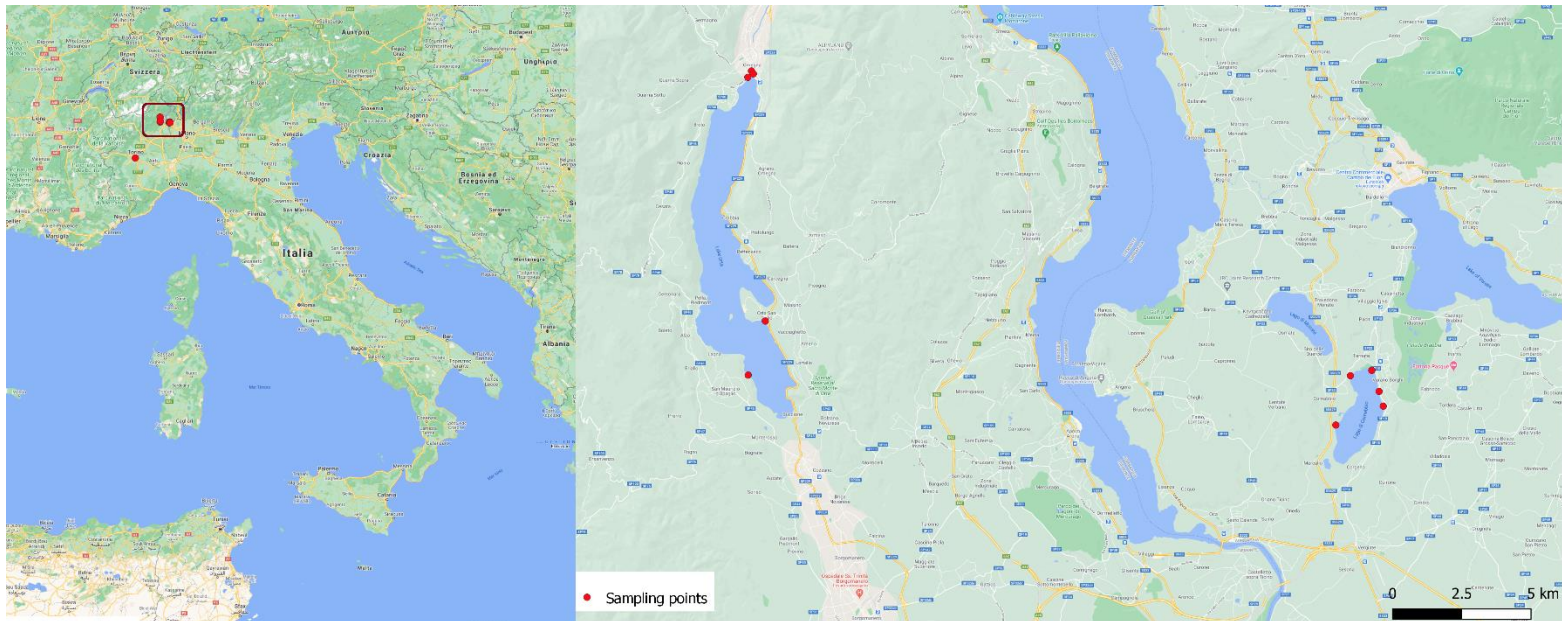


Figure A2c: A map of the sampling points from the lakes Orta and Comabbio in Italy included in Chapter 5, and their positions on the national territory.

Table A8. MzMine Parameters, used for the elaboration of the data in Chapter 5.

| Step | Parameter | Setting |
|----------------------------|--------------------------|------------------------------------|
| Mass detection | Retention Time | 0-20 min |
| | MS level | 1 |
| | Polarity | + for ESI (+) - for ESI (-) |
| | Spectrum type | Profile |
| | Mass detector | Exact mass |
| | Noise level | 2,00E+02 (ESI-) 5,00E+02 (ESI+) |
| Chromatogram builder | Retention time | 0-20 min |
| | MS level | 1 |
| | Polarity | + for ESI (+) - for ESI (-) |
| | Spectrum type | Profile |
| | Min time span (min) | 0,01 |
| | Min height | 2,10E+02 (ESI-) 5,10E+02 (ESI+) |
| | <i>m/z</i> tolerance | 0,001 <i>m/z</i> or 5 ppm |
| Smoothing | Filter width | 7 |
| Chromatogram deconvolution | Algorithm | Noise amplitude |
| | Min peak height | 3,00E+02 (ESI-) 6,00E+02 (ESI+) |
| | Peak duration range | 0,05-1,00 min |
| | Amplitude of noise | 2,00E+02 (ESI-) 5,00E+02 (ESI+) |
| Isotopic peak filter | <i>m/z</i> tolerance | 0,001 <i>m/z</i> or 5 ppm |
| | Retention time tolerance | 0,01 min |
| | Maximum charge | 1 |
| | Representative isotope | Lowest <i>m/z</i> |
| Join aligner | <i>m/z</i> tolerance | 0,001 <i>m/z</i> or 5 ppm |
| | Weight for <i>m/z</i> | 70 |
| | Retention time tolerance | 0,3 (absolute) min |

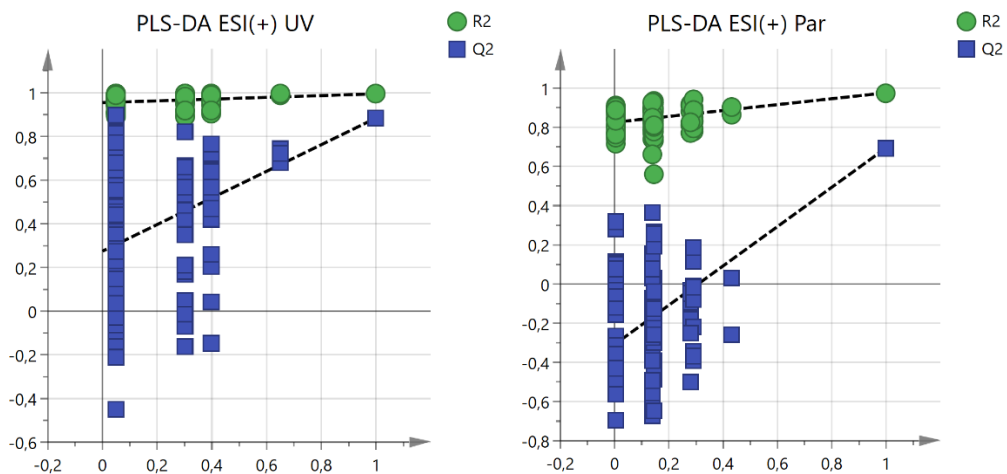


Figure A3: Validation tests for the PLS-DA model in UV scaling for the features detected in ESI (+) based on 100 permutations/replicated (left), and the PLS-DA model in Pareto scaling for the features detected in ESI (+) based on 100 permutations/replicated (right).

Training Activities, Seminars and Conferences Attended

- **14-19/03/2018, 07/05/2018:** Health and safety rules and laws in Italy, risks and hazards in the laboratory, SMAT, Turin, Italy. *24 hours*
- **27/03/2018, 18/04/2018:** UHPLC-QTRAP MS course, SMAT, Turin, Italy. *12 hours*
- **10-12/10/2018:** XENOWAC II "Challenges and Solutions related to Xenobiotics and Antimicrobial Resistance in the Framework of Urban Wastewater Reuse: Towards a Blue Circle Society' conference", Limassol, Cyprus.
- **24/01/2019, 28/01/2019:** Transizione alla norma UNI EN ISO/IEC 17025:2018 e requisiti Accredia (RT-08 per i CAB e RT-25 per i LAT): principali novità, approccio al rischio, requisiti della norma nella nuova edizione, SMAT, Turin, Italy. *8 hours*
- **10/03/2019:** Corso di Formazione Generale alla Salute e Sicurezza per i Lavoratori, Online, UniTO. *4 hours*
- **26-30/05/2019:** SETAC Europe 29th Annual Meeting, Helsinki, Finland.
- **04-06/09/2019:** International Conference on Chemical Energy and Semiconductor Photochemistry (CESCOP 2019), Trabzon, Turkey.
- **16-20/06/2019:** 17th International Conference on Chemistry and the Environment (ICCE 2019), Thessaloniki, Greece.
- **30/09/2019:** Piano di Sicurezza dell'acqua per la città di Torino, Presentazione e Pianificazione delle attività, SMAT, Turin, Italy. *4 hours*
- **17-18/02/2020:** Colloquium "Machine Learning meets Chemistry", University of Turin, Turin, Italy. *8 hours*
- **31/03/2020:** Corso Privacy per le persone autorizzate al trattamento dei dati personali, Online. *4 hours*
- **16/04/2020, 20/04/2020:** Italian Mass Spectrometry Society (IMaSS) webinar on Data Analysis, Online. *8 hours*
- **01-12/06/2020:** American Society of Mass Spectrometry (ASMS) 2020 Reboot Conference, Online.
- **12/06/2020:** Sample Preparation, Quo Vadis: Current Status of Sample Preparation Approaches (Molecules MDPI) webinar, Online. *4 hours*
- **15-16/06/2020:** ASMS Untargeted Metabolomics short course, Online. *16 hours*
- **20-24/07/2020:** UPLC-QTOF MS course, SMAT, Online. *16 hours*
- **18/01/2021:** EU versus Italian water management, Online. *3ECTS*
- **01/02/2021:** Basics of project writing: hands-on workshop session, Online. *4 hours*
- **05/02/2021:** Antibiotic Resistant Bacteria: Occurrence and removal from Urban Wastewater, Online. *2 hours*

PhD Courses Attended

| University | Course Title | Hours |
|---------------------|---|-------|
| University of Turin | Introduction in Technological Sector of Circular Economy / Water Treatment Plan | 4 |
| University of Turin | Circular Economy and Social Innovation | 3 |
| University of Turin | New Technologies and Climate Change– “Rethinking the Design of Climate Stabilization Policy” | 6 |
| University of Turin | Economic Fundamental Principles | 3 |
| University of Turin | Economics of Innovation and Green Technologies | 12 |
| University of Turin | Organization, Innovation and Value Measures | 3 |
| University of Turin | Data Science and Circular Economy | 4 |
| University of Turin | Circular Economy and Public Policies | 4 |
| University of Turin | Public Policies for the Environment | 6 |
| University of Turin | Innovation, sustainability and new business models | 6 |
| University of Turin | Resources and Raw Materials | 9 |
| University of Turin | LCA - Life Cycle Assessment – Principles, Plastic Materials in the Circular Economy Paradigm, Metallic Materials in the Circular Economy Paradigm | 4 |
| University of Turin | OpenLCA - The open source software for LCA, LCA analysis of plastic and metallic materials | 4 |
| University of Turin | Interactive laboratory to stimulate an attitude to outreach activities on basic physico-chemical phenomena | 12 |

PhD Schools Attended

| Date | Title | Place |
|-----------------------|--|--------------------------|
| 25/04/2018-27/04/2018 | Summer School on Photochemistry and Depollution | Clermont-Ferrand, France |
| 27/08/2018-29/08/2018 | International Summer School on “Micropollutant Analysis and Abatement” | Aalborg Denmark |
| 04/03/2019-05/03/2019 | International Winter School on Mass Spectrometry and Workshop on substances prioritization | Palaiseau France |
| 03/06/2019-07/06/2019 | 3rd European Summer School on Environmental Applications of Advanced Oxidation Processes | Alcoy Spain |
| 23/09/2020-24/09/2020 | Summer school on “Introduction to Basic Statistical Tools and Data Analysis in Research” | Online |

Presentations

1. Oral Presentation: **D. Papagiannaki**, P. Calza, R. Binetti, "General Introduction-ESR6", 1st AQUALity Meeting, Clermont-Ferrand, France, 23-24 April 2018.
2. Oral Presentation: **D. Papagiannaki**, P. Calza, R. Binetti, "PFAS determination in real water samples by UHPLC/MS/MS", Scientific Storm Meeting, Turin, Italy, 30 May 2018.
3. Oral Presentation: **D. Papagiannaki**, F. Barsotti, M. Fungi, R. Binetti, "PFAS determination in real water samples", 2nd AQUALity Meeting, Aalborg, Denmark, 30-31 August 2019.
4. Poster Presentation: **D. Papagiannaki**, F. Barsotti, A. Salaris, M. Fungi, R. Binetti, "Fate of Metolachlor and Terbutylazine in surface water and related drinking water treatment plant", XENOWAC II, Limassol, Cyprus, 10-12 October 2018.
5. Oral Presentation: **D. Papagiannaki**, R. Binetti, "Trace level analysis of CECs in drinking water using Mass Spectrometry", International Winter School on Mass Spectrometry, Palaiseau, France, 4-5 March 2019.
6. Oral Presentation: **D. Papagiannaki**, S. Morgillo, P. Calza, R. Binetti, "PFAS an overview", 3rd AQUALity Meeting, Palaiseau, France, 7-8 March 2019.
7. Poster Presentation: **D. Papagiannaki**, S. Morgillo, G. Costantino, M. Fungi, R. Binetti, "Perfluoroalkyl Substances Assessment in Turin Metropolitan area and correlation with potential sources of pollution according to the Water Safety Plan risk management approach", SETAC Europe 29th Annual Meeting, Helsinki, Finland, 26-30 May 2019.
8. Poster Presentation: **D. Papagiannaki**, S. Morgillo, G. Costantino, M. Fungi, R. Binetti, "Trace level analysis of perfluoroalkyl substances in drinking water and their assessment in Metropolitan Area of Turin", 17th International Conference on Chemistry and the Environment, Thessaloniki, Greece, 16-20 June 2019.
9. Oral Presentation: **D. Papagiannaki**, P. Roslev, "Effect of UV-A, UV-B and UV-C irradiation on Biototoxicity of Glyphosate in Drinking Water Samples", 4th AQUALity Meeting, Trabzon, Turkey, 2-3 September 2019.
10. Oral Presentation: **D. Papagiannaki**, P. Calza, R. Binetti, "Evaluation of CECs in drinking water including toxicological assessment of their degradation by-products", Horizon 2020 MSCA-ITN cluster event "Clean Water", Girona, Spain, 22 October 2019.

11. Oral Presentation: **D. Papagiannaki**, S. Morgillo, R. Binetti, "Screening-level risk assessment of selected pharmaceuticals in the Metropolitan Area of Turin", 5th AQUALity Meeting, Online, 6 April 2020.

12. Poster Presentation: **D. Papagiannaki**, R. Binetti, P. Calza, P. Roslev, "UV Irradiation Decreases Ecotoxicity of Glyphosate", SETAC SciCon, SETAC Europe 30th Annual Meeting, online, 3-7 May 2020.

13. Oral Presentation: **D. Papagiannaki**, Z. Varga, A. Cedrino, R. Binetti "Monitoring, understanding and predicting contamination of groundwater sources destined for drinking water supply", 6th AQUALity Meeting, Online, 22 September 2020.

14. Oral Presentation: **D. Papagiannaki**, D. Fabbri, P. Calza, R. Binetti, "Non-target screening analysis of water samples using LC-HRMS and GC-MS", 7th AQUALity Meeting, Online, 1-2 March 2021.

15. Oral Presentation: **D. Papagiannaki**, D. Palma, A. Cedrino, G. Molinari, M. Lai, M. Minella, R. Binetti "Removal Of Contaminants Of Emerging Concern From Water Using High Voltage Pulsed Electric Field Discharge", SPEA 2020 postponed, planned.

Outreach Activities

1. E. Robotti, M.H. Belay, N.P.F. Gonçalves, **D. Papagiannaki**, F.E.B Coelho, "AQUALity Project: Removal of contaminants of emerging concern", European Researcher's night 2018, Alessandria, Italy, 28 September 2018.

2. **D. Papagiannaki**, A. Pavanello "AQUALity Project: Removal of contaminants of emerging concern", Festival dell'Innovazione e della Scienza 2018, Settimo Torinese, Italy, 20 October 2018.

3. M.H. Belay, N.P.F. Gonçalves, **D. Papagiannaki**, I. Berruti, K. Janowska, F.E.B Coelho "AQUALity Lab", laboratory experiments for secondary school students, Turin, Italy, 19 February 2019.

4. **D. Papagiannaki**, R. Binetti "PFAS Assessment in Metropolitan Area of Turin, Italy", TV Interview, TG Leonardo, Rai News, 17 April 2019.

5. N.P.F. Goncalves, **D. Papagiannaki**, D. Fabbri, P. Calza, "Il trattamento dell'acqua: abbattimento degli inquinanti", European Researcher's night 2019, Turin, Italy, 27 September 2019.

6. N.P.F. Gonçalves, Z. Varga, **D. Papagiannaki**, "AQUALity Lab", laboratory experiments for secondary school students, Turin, Italy, 12 February 2020.
7. I. Berruti, M.H. Belay, **D. Papagiannaki**, "Il suolo come filtro naturale per gli inquinanti", Video presentation (translated into Italian), European Researcher's night 2020, online, 23 November 2020.
8. **D. Papagiannaki**, C. Jimenez Holgado, D. Fabbri, P. Calza, "La sfida di AQUALity: rimuovere i contaminanti di ultima generazione dalle acque", Video presentation, European Researcher's night 2020, online, 24 November 2020.
9. **D. Papagiannaki**, R. Binetti, "L'acqua", Online lesson to secondary school students, European Researcher's night 2020, online, 27 November 2020.
10. N.P.F. Gonçalves, **D. Papagiannaki**, I. Berruti, P. Calza, "Pandemia e ricerca", TV interview, TGR Piemonte, TG Leonardo, 6 May 2021.

Periods as Visiting Researcher

1. **01/05/2019-31/08/2019**: Department of Chemistry and Bioscience, Aalborg University, Aalborg, Denmark. Tutor: Prof. Peter Rolsev.
2. **01/11/2020-28/02/2021**: Department of Chemistry, University of Turin, Turin, Italy. Tutor: Prof. Debora Fabbri.

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