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1 **Opinion paper on strategies for toxicity assessment of organic trace pollutants in wastewater**

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1 ***Abstract***

2 This opinion paper focuses on the role of eco-toxicological tools in the assessment of possible impacts  
3 of emerging contaminants on the aquatic ecosystem, hence, on human health. Indeed, organic trace  
4 pollutants present in raw and treated wastewater are pivot targets: a multidisciplinary approach allows  
5 defining the basic principles for managing this issue, from setting a proper monitoring campaign up to  
6 evaluating the optimal process treatment. Giving hints on trace pollutants fate and behaviour, attention  
7 has been focused on the choice of bioassay(s), by analysing the meaning of biological answers. Data  
8 interpretation and exploitation are detailed with the final goal of providing criteria to be able to select the  
9 best targeted treatment options. The manuscript deals with conventional and innovative analytical  
10 approaches for assessing toxicity, by reviewing laboratory and field assays; illustrative real scale and  
11 laboratory applications integrated and exemplified the proposed approach.

12

13 ***Keywords***

14 Aquatic ecosystem; bioassays; ecotoxicity; micro-pollutants; risk assessment; wastewater treatment

15

16 ***Highlights***

- 17 • Bioassays-derived information in assessing organic trace pollutants toxicity is explored
- 18 • Meaning, role and choice of proper bioassays are discussed
- 19 • Hints on sampling raw and treated wastewater, which are a complex matrix, are given
- 20 • Interpretation and use of results of bioassays are suggested
- 21 • Real and laboratory scale experiences are presented as case studies

22

23

## 1 **1. Introduction**

2 Ecological risk assessment is the scientific decision supporting process for gauging risks based on the  
3 occurrence of physical or biological agents or the amount of chemical/mixture of chemicals/emission  
4 discharged into the environment, on the exposure of an ecological receptor (e.g. plants, fish, birds, ...)  
5 and on the inherent toxicity of the agent itself. The awareness to investigate the effects of pollutants  
6 exposure throughout the whole lifespan of an organism (or during specific phases of its development)  
7 demands a new approach. This comes up beside and integrates conventional tests, issued accordingly  
8 with established guidelines and performed on specific laboratory organisms, generally aimed to assess  
9 short to mid-term effects.

10 Together with the monitoring of different effects on a single organism (from the first life periods),  
11 nowadays it is clear that also cross-generational, ecological and ethological aspects must be taken into  
12 account [1], [2], [3]. Ecotoxicity testing strategies are developed worldwide and supported by  
13 international organizations. Risk characterization/assessment schemes are tiered, enabling a progressive  
14 refinement of exposure/effect ratios. Nevertheless, it is not possible to specify the number of tiers  
15 generally required since they depend on each specific situation, as the complexity of community  
16 structures and relationships among different populations should be considered.

17 This opinion paper aims to gather the main findings obtained by various research groups involved in the  
18 COST Action ES1202 Conceiving Wastewater Treatment in 2020-Energetic, environmental and  
19 economic challenges (Water\_2020). The final goal is to present the strength of a multi-tiered approach  
20 within the risk assessment of pollutants/whole effluents detailing potentials and weighing pros and cons  
21 of conventional and innovative bioassays. The work considers some exemplifying case studies of  
22 wastewater treatment and, finally, examines the socio-economic aspects of the whole procedure of  
23 ecotoxicity evaluation.

24

## 1 **2. Principal well-established knowledge and open issues**

2 As far as chemical pollution is concerned, substances are prioritized for action based on the risk to or via  
3 the aquatic environment, according to the Water Framework Directive (2000/60/EC; WFD) [4], and  
4 included in article 16 for the “Strategies against pollution of water”. Compounds must be identified by a  
5 targeted risk-based assessment focusing solely on aquatic ecotoxicity and human toxicity via the aquatic  
6 environment. According to WFD, “hazardous substances” are defined as substances or groups of  
7 substances that are toxic, persistent and liable to bio-accumulate, and other substances or groups of  
8 substances, which give rise to an equivalent level of concern. In the Risk Assessment Process, the initial  
9 step would be the hazard identification. Further, the primary investigation should already include the  
10 possible health problems caused by the pollutants. This process uses the intrinsic properties of chemicals  
11 (i.e. persistence, solubility,  $K_{ow}$ , volatilisation, ...) to determine expected adverse effects, and on the other  
12 hand, to estimate the probability of adverse effects to occur. In addition, the physical-chemical data  
13 provide information about the relevance of some exposure paths. As the next step, and already partly  
14 depending on the nature of the substance(s) under scrutiny, proper analytical tools capable of providing  
15 deeper information on exposure and effects are required: the most commonly applied are acute, sub-  
16 chronic and chronic toxicity, abiotic and biotic degradability, and bioconcentration or bioaccumulation.  
17 Consequently, these tests can be used for toxicants and hydrophobic chemicals with low solubility in  
18 water that may accumulate and concentrate in lipids of aquatic organisms. The storage of these toxicants  
19 within the organism demonstrates cumulative toxicity to exposed organisms. These chemicals are usually  
20 considered in bio-concentration tests, which use bio-concentration factors (BCF) to predict the  
21 accumulation of hydrophobic contaminants. BCFs may be obtained either from calculations ( $\log K_{ow}$ ,  
22 (Q)SAR, or other computational-based model), or from experiments. This parameter represents the  
23 partitioning of the chemicals between an organism and the surrounding environment [5] based on the  
24 observation of the hydrophobic character. Thus, values of  $BCF > 1$  indicate for hydrophobic character,

1 not bioaccumulative if the BCF in the range 1000–5000. Further, in case BCF is > 5000, the substance  
2 is very bioaccumulative [6].

3 During an exposure assessment, the following questions must be answered: 1) To which pollutant doses  
4 are humans and ecosystems exposed throughout a given lapse of time? and 2) How many individuals,  
5 species or populations are exposed? In case of dose-response assessments, quantitative data regarding  
6 biological effects under different situations and types of exposure must be supplied. Either finally, risk  
7 assessment can be carried out, comparing exposure and effects, quantitatively or qualitatively, thus  
8 determining the probability of each effect occurring. Both hazard and risk assessments are mandatory to  
9 guarantee scientific support for regulations [7]. In the last decade there has been a growing interest about  
10 the impact on the whole ecosystem exerted by chemicals, on aquatic and terrestrial organisms and  
11 wildlife [8], [9], [10], [11], [12].

12 Ecotoxicity tests can be classified with regards to design (e.g. field, laboratory, computational), level of  
13 biological organization (population, assemblage/community, ecosystem), exposure period (acute, sub-  
14 chronic, chronic) and endpoint (lethal, sublethal). Short-term (“acute”) tests are generally used first, with  
15 survival as the most common endpoint. Long term (“chronic”) tests (with the observation of sublethal  
16 effects on organism growth or reproduction) are then used when results from short-term tests combined  
17 with large safety factors suggest that there may be risks to the environment.

18 The use of acute and chronic tests in ecotoxicology has been proposed in reports from EU’s Registration,  
19 Evaluation, Authorization and Restriction of Chemicals (REACH) aiming to improve the protection of  
20 human health and the environment through better and earlier identification of the inherent properties of  
21 chemical substances. Moreover, in order to reduce the use of animals in laboratory experiments, REACH  
22 suggests a number of possibilities to adapt the testing requirements and to use existing data, as well as  
23 alternative assessment approaches [13].



1 Since the Seventies, the progressive awareness of hazards linked with specific chemicals has been  
2 increasingly consolidated by epidemiology findings, the long-term follow up of environmental disasters  
3 and the availability of new technological tools to isolate and quantify a huge range of analytes from  
4 complex matrices also at risible concentrations [14]. For instance, it has been possible to carry out  
5 investigations on metal speciation. In addition, almost every class of organic compounds has been  
6 considered, starting from reaction by-products (*e.g.*, among the firstly studied, the disinfection by-  
7 products, such as the trihalomethanes), to persistent organic pollutants (POPs), and, finally, to the  
8 thousands of substances derived from the everyday use, such as PPCPs (pharmaceuticals and personal  
9 care products) and their metabolites and transformation products. Furthermore, research has focused on  
10 the size of pollutants released into the environment, by considering, *inter alia*, micro-plastics and  
11 nanomaterials. It is now well known that the size of chemical agents strongly affects both the  
12 bioavailability and the effects on the organisms. So far, the scientific literature numbers lot of remarkable  
13 works focusing on the detection of (trace) pollutants, both organic and inorganic, the study of their fate  
14 and behavior into the environment, their toxicity and the feasibility of their replacement and removal  
15 from the contaminated areas [15], [16], [17], [18], [19].

16 The present knowledge indicates that thousands of organics in trace quantities are widespread in  
17 ecosystems, aquatic organisms being important targets, as they are exposed to wastewater residues over  
18 their entire life.

19 Concentrations of pharmaceuticals in the range pg/L - µg/L have been reported worldwide in effluents  
20 from sewage treatment plants, surface, ground and even drinking water [20]. Since now, over 3,000  
21 different agents in human pharmaceuticals are used and continuously released into the environment, it is  
22 urgent to increase our knowledge about the potential combined effects of these substances on the biota.

23 Most research regarding the ecotoxicological effects of pharmaceuticals is generally focused on aquatic  
24 organisms and their food webs in standardized tests. However, despite the presence of mixtures of

1 multiple compounds in environmental media and samples, theoretical considerations and experimental  
2 findings suggest that the overall risk may be driven by only few components in these mixtures [21], [22],  
3 [23]. Furthermore, routinely detected chemicals often cannot explain observed biological responses [24],  
4 which point to a mismatch in the generally applied assessment approaches.

5 Wastewaters, due to their nature and origin (i.e. municipal, industrial, runoff, grey), evidently collect and  
6 concentrate a multitude of chemicals, that make up complex mixtures including microbial consortia.

7 Therefore, when assessing the overall impact of any given wastewater (either raw or treated), it is  
8 essential to consider both: i) the removal/discharge of specific micropollutants and ii) the residual toxicity  
9 of parent compounds, metabolites, transformation products and by-products after treatment or, in any  
10 case, the discharge. For this reason, to gain insight not only into the impact of individual micropollutants,  
11 but also into the effects exerted by wastewater as a whole, bio-analytical tools are compulsory. The  
12 removal of a target xenobiotic compound does not necessarily mean that the treatment process is  
13 detoxifying, because adverse effects may be a result of the conversion of chemicals into metabolites or  
14 breakdown products more toxic than the parent compounds. In Figure 1, the temporal and logical  
15 structure of the above-mentioned process has been depicted: after identifying and characterizing the  
16 chemicals, it is necessary to consider their integrated effects towards both humans and all levels of food  
17 webs.

18

19

20 Figure 1. Towards a global risk assessment of aquatic pollution: present and future scenarios.

21

22

23 **3. Which responses should be measured? A focus on “real life”**

24 **3.1 Principles**

1 From a “real life” perspective, a lot of possible responses can be expected and observed in the organisms  
2 exposed to the mixture of pollutants present in raw and treated wastewaters.

3 According to [25], ecotoxicology is the science devoted to the study of the contaminants and their effects  
4 induced on all parts of the biosphere. Thus, all species, including human beings, must be considered as  
5 potential targets of the toxic action of the pollutants. Therefore, upon exposure to chemicals, different  
6 responses can be observed, ranging from the molecular to the ecosystem level.

7 The tools available to assess effects on ecosystems [7] can be related to the different steps of the  
8 hierarchical ecotoxicology described by [25]: organismal, population, community, and ecosystem. None  
9 of them must be considered the most or the least relevant in the evaluation process, as each of them  
10 represents a specific target.

11 At the organismal level, different single-species bioassays (of which some are standardized) can be  
12 applied to identify hazards through all relevant routes of exposure: soil, water, air and food. Several of  
13 these tests can be run using diverse species providing data on chemically induced effects, which can be  
14 applied in ecological risk assessments (ERA) to yield Species Sensitivity Distributions (SSDs). SSDs are  
15 models of the variation in sensitivity of species to chemicals or other stressors [26]. In following steps,  
16 long-term single-species tests and laboratory multi-species tests can be used to predict or evaluate  
17 population dynamics. Finally, mesocosm and field assays can offer information about real ecological  
18 effects [7].

19 Lots of well-known biochemical and molecular mechanisms enable to explain the toxic action of  
20 contaminants and their subsequent effects. Enzyme dysfunctions (inhibition, activation or induction),  
21 DNA alterations, oxidative stress and generation of reactive oxygen species (ROS), oxidative  
22 phosphorylation inhibition, heme biosynthesis inhibition are typical mechanisms associated to toxicants  
23 [25]. It is also relevant that some products of detoxification reactions may be more toxic or reactive than  
24 the parent compound and, therefore, specific tests on drug metabolism have been developed. In general,

1 the biochemical and molecular responses express morphological (microscopic and macroscopic scale)  
2 modifications on cells and tissues, and/or functional failures of organs and systems.

3 Frequently, the biochemical mechanisms of toxic action are unknown, and therefore the effects on cells,  
4 tissues, organs and systems can be identified as the only target in the risk assessment. Different  
5 morphological findings related to the exposure to organic pollutants have been described in cells and  
6 tissues, such as inflammation, necrosis, apoptosis, and sometimes hypertrophy or hyperplasia. Genetic  
7 alterations can explain pathologic cell proliferation and cancer.

8 Tissues, organs and systems involved in vertebrate toxicokinetic, and the effects displayed on respective  
9 targets are important subjects in both toxicology and ecotoxicology. Integument, respiratory and  
10 digestive organs are first subjected to organic pollutant exposure, due to their direct interface with the  
11 outside. The first route of exposure to organic pollutants regards undoubtedly the derma, which is also  
12 the target of their toxicant action. In addition, dermal structures play a crucial role also in case of  
13 excretion processes (e.g., by means of fish and reptile scales, bird feathers and hair of mammals) and  
14 bioaccumulation phenomena. Secondly, respiratory exposure is included, while organs associated with  
15 digestion represent the third major exposure route, *i.e.*, ingestion. In addition, liver/hepatopancreas and  
16 kidney, and other analogous organs of invertebrates can be relevant as target of the toxic action,  
17 influencing the kinetic metabolism and excretion of the chemicals.

18 Nervous, immune, and endocrine are mostly studied; a wide range of toxicants affects adversely nervous  
19 system cells and tissues, through both structural and functional alterations, expressing also a wide range  
20 of manifestations at the organismal level [25]. Concerning the immune system, research focuses mainly  
21 on either cellular or humoral immune dysfunction in adult and developing vertebrates. The relevance of  
22 these immune alterations is related to the ability to induce infection, infestation, or other diseases like  
23 cancer; with consequences, not only at organismal level, but also at population, and community levels.

1 Endocrine disruption was historically investigated especially in terms of estrogenic/androgenic activities,  
2 but, so far, plenty of hormone-mediated mechanisms have been explored.

3 At the border between organismal and population levels, alterations on growth and development,  
4 reproduction, and behavior, are issues of concern to public and environmental health that justify the  
5 implementation and development of specific tests to evaluate these alterations induced by the exposure  
6 to chemicals. The studies of pollutant effects on growth are common because the collection of data is  
7 easy, and they have been implemented on amphibians, fish, birds, terrestrial and aquatic invertebrates,  
8 terrestrial and aquatic macrophytes, and microscopic plants. An exhaustive list of plants for toxicity tests  
9 has been compiled by [27] and [28]. In relation to reproductive effects, there are standardized tests on  
10 aquatic and terrestrial invertebrates' species, such as *Daphnia magna* or collembolans, as well as on  
11 vertebrates such as birds. Finally, modifications and alterations of the behavior are also commonly  
12 selected endpoints to evaluate the risks associated to environmental polluted scenarios. These endpoints  
13 range from individual alterations, such as movement or foraging behavior, to social interactions, such as  
14 social structuring or mating behaviors.

15 Figure 2 illustrates the core message of this paragraph: there exists an indissoluble connection between  
16 each target phenomenon and the biological hierarchy. It must be underlined that the investigator must be  
17 aware of the inevitable partiality of any obtained information, which just accounts for an element of a  
18 complex framework.

19

20 Figure 2. Approaching the bioassays: the conceptual framework.

21

22

23

### 24 **3.2 Biological responses**

1 Among the variety of biological effects currently studied in environmental research, this section focuses  
2 on the responses that can be used to assess the activity of wastewater.

3 A basic principle in toxicology research is the study of a toxicant fate in an exposed organism. To assess  
4 this, the Absorption, Distribution, Metabolism and Excretion (ADME) concept has been developed  
5 decades ago. Each of the mentioned stages of pollutant impact on an organism is characterized by specific  
6 and quantitative measures, like distribution coefficients that assess uptake potential into certain organs  
7 or cell types. Metabolism is characterized by the enzymatic activation, cleavage or conjugation of parent  
8 compounds, and excretion by the velocity of transport from the target back into the environment. For  
9 compounds or conditions where clearance from the body cannot be reached, toxicology tries to assess  
10 the biological activity by describing the mode of action (see 3.2.1 Baseline toxicity). However, reliable  
11 information on mode of action is still lacking for many chemicals. Currently, there is neither an agreed  
12 inventory of modes of action, nor a defined set of criteria on how to characterize or predict a mode of  
13 action for data-poor chemicals or how to group chemicals into assessment groups [29].

14 Two approaches can generally be followed to assess the toxicity of pollutants and effluents, based on  
15 different methodological criteria: Toxicity Identification Evaluation (TIE), which focuses on the whole  
16 organisms (therefore, bioavailability and baseline toxicity are the core issues) and Effects-Directed  
17 Analysis (EDA), which relies on *in vitro* assays, by considering specific endpoints, *i.e.*, specific modes  
18 of action [10].

19

### 20 **3.2.1 Baseline toxicity**

21 As far as chemical substances are concerned, every mechanism of toxicity is initiated by the interaction  
22 between the chemical(s) and the organism and can be described accordingly to the following sequence:  
23 exposure, bioavailability and formation of a bond with the ligand. Consequently, two opposite scenarios  
24 can reveal either alteration or adaptation, both driven by complex pathways. Sub-lethal responses can be

1 evaluated by quantifying proper physiological condition indices, linked to morphometry, biochemistry  
2 and growth.

3 Baseline toxicity involves the interaction between the substances and the cell membrane; the  
4 hydrophobicity affects the capacity of the molecules to react and pass through these barriers, whose  
5 fluidity can thus be deeply modified. Therefore, several biochemical pathways can be impaired, such as  
6 the electron transfer chain in photosynthesis, in case of vegetal cells, or specific enzymatic activities,  
7 linked for instance with the electrical signal transmission and the transport mechanism in case of animal  
8 cells.

9 Regarding the field of environmental research, increasing attention has been paid to *in situ* toxicological  
10 studies. Laboratory scale assays (*in vitro* and *in vivo*) may not allow a relevant simulation of real cases,  
11 due to the presence of other stressing factors in environmental conditions that influence the biological  
12 response. Thus, laboratory scale experiments might dramatically stray from real situation. In terms of  
13 baseline toxicity, it is even more difficult to assess the impact of pollutants on populations and  
14 communities, represented by for example biomass decrease or change in the total numbers of individuals  
15 can be a possible outcome.

16 Finally, another critical aspect inherent to toxicity assessment is the multi-faceted scenario of responses,  
17 which can present non-monotonic trends and can be affected by hormesis and adaptation phenomena  
18 [30].

19

### 20 **3.2.2 Endocrine disruption**

21 As pointed out by [31], different organizations have issued their own definition of EDCs, taking into  
22 account growing levels of biological effects. The well-known and often cited definition of United States  
23 Environmental Protection Agency (USEPA). considers the effects on “*maintenance of the homeostasis*  
24 *and the regulation of developmental processes*” [32], while World Health Organization (WHO) refers to

1 “*adverse effects in an intact organism, or its progeny, or (sub-)population*” [33]. The Endocrine Society  
2 [34] more strictly states that EDCs “*interfere with any aspect of hormone action*”. Consequently, the  
3 evaluation of possible endocrine activity caused by a substance, a mixture or any other complex matrix  
4 (such as sewage, sludge, etc.) must follow the precise definition of the final goal.

5 Secondly, it is mandatory to select the mechanisms to investigate, by taking into consideration the  
6 biological complexity of the target organism and its physiology. Several mechanisms account for  
7 endocrine impairment: the most commonly studied are the bonds with nuclear receptors (this super  
8 family includes 48 types, in case of humans), and the interactions with membrane receptors, cytosolic  
9 receptors, orphan nuclear receptors. Moreover, epigenetic changes as well as regulation cascade  
10 processes, effects on hormones and oxidative metabolism can be numbered among the modes of action  
11 of EDCs. The modes of action concern all the biological levels, from single cells to the whole organisms,  
12 both with acute and chronic effects, including reproductive, immunological and neurological disorders,  
13 cancer, diabetes, and obesity.

14 EDCs exhibit multiple modes of actions, resulting in dose-effect relationships not always following a  
15 monotone trend and changing entirely as a function of concentrations and depending on the final target.  
16 The case of bisphenol A (BPA) is emblematic: it behaves as a relatively weak estrogen towards estrogen  
17 receptor alpha ( $ER\alpha$ ) in comparison with the natural hormone estradiol, while it is equipotent towards  
18 membrane receptors [35], [36]. Furthermore, the effects of EDCs can differ based on the developmental  
19 stage of the organisms (e.g., pre-natal, post-natal and adult forms), as pointed out by [31] and [37].

20 The most studied mechanisms are related to the interference with sexual hormones (androgens and  
21 estrogens), thyroid and adrenal disorders and glucose metabolism impairment [38]; therefore, assays  
22 have been designed basically aiming at their assessment.

23 Many biological pathways are based on nuclear receptors that migrate into the nucleus and regulate gene  
24 transcription after hormone binding, despite their location. The main pathways include: thyroid signaling,



1 estrogen signaling, glucocorticoid pathway, renin-angiotensin-aldosterone system, leptin and insulin  
2 signaling. The main elicitors of estrogenic activity are micropollutants such as: the natural (estrone,  
3 estradiol, estriol) and synthetic (ethynil-estradiol) estrogens, alkylphenols (octyl- and nonyl-phenol) and  
4 the bisphenol A. The main adverse impact related to this kind of biological activity is represented by  
5 impairment of reproductive performance in wildlife. Routledge et al. [39] documented that steroidal  
6 hormones can induce feminization in fish and other aquatic organisms at concentrations as low as 1 ng/L  
7 or less estradiol equivalents. Witters et al. [40] reported that significant reproduction effects in male and  
8 female fish might appear at levels above 10 ng/L of estradiol equivalents. Moreover, any interference in  
9 their homeostasis is related to several human diseases (breast, ovary and prostate tumours, cardiovascular  
10 diseases, infertility, etc.), due to the physiological role of estrogens.

11

### 12 **3.2.3 Genetic toxicity**

13 The toxicity towards the DNA and the genetic processes exhibits a wide spectrum of effects, and,  
14 therefore, can be investigated by means of several complementary tests. The observed phenomena  
15 include genotoxicity (not directly transmissible), mutagenicity (heritable change in a genotype),  
16 mechanisms of DNA repair, carcinogenesis, and developmental toxicity. Beside chemical agents, which  
17 are the core object of this paper, also biological and physical stressors can cause DNA damages (e.g.,  
18 viruses, ionizing radiations, UV radiations); furthermore, the overall conditions of the organism, in terms  
19 of feeding regime, disruption of regulatory pathways and cellular checkpoints dramatically favors  
20 adverse events to DNA. Consequently, the assessment of possible genetic toxicity caused by trace  
21 pollutants contained in wastewater should be considered the coexistence of the aforementioned factors,  
22 together with the fact that wastewater are mixtures of chemicals, whose concentrations are extremely  
23 variable (as explained in paragraphs 4.1 and 4.2). DNA damage can be related to alkylation (i.e H- bonds  
24 alteration, errors in base-pairing), hydroxylation (hence, errors in base-pairing), deamination (bringing

1 on changes from cytosine to uracil, then errors in base-pairing and base substitution), formation of base  
2 analogues (for instance by replacement of H atoms with halogens) leading to errors in base-pairing and  
3 base substitutions, strand breaks, and intra/interstrand cross links. Large planar molecules can intercalate  
4 within the double helix, without reacting but disrupting replication, recombination and repair. The  
5 mutations consist in point mutations (referred to nucleotide substitutions), yielding to errors in amino  
6 acids coding, and chromosomal mutations (consisting in deletion or insertion of several contiguous  
7 genes, inversion of genes on a chromosome, exchange of large segments of DNA between non-  
8 homologous chromosomes) which lead to several mistakes in amino acids coding and, thus, to major  
9 phenotypic consequences. Mechanisms of DNA repair are often exploited in bioassays; they are aimed  
10 to restore its pristine function or to destroy the damaged cell by means of apoptosis. They are based on  
11 the action of multiple enzymatic reactions, which, for instance, allow repairing base excision, nucleotide  
12 excision, double strand breaks and mismatches.

13 Decades of research have shown that mutagenesis is a critical component and that there is a strong  
14 correlation between carcinogenicity and mutagenicity [41].

15 Recent developments in genomics, together with the decrease of equipment prices and the wide  
16 availability of sophisticated tools (such as DNA micro-arrays) have contributed to a tremendous  
17 exploitation of molecular techniques. Starting from genome sequencing, this led to the study of  
18 expression profiling (m-RNA transcripts, miRNA, ncRNA), the so-called transcriptomics, until the  
19 characterization of protein (proteomics), peptide (peptidomics) and metabolic profiles (metabolomics).

20 The application of these analyses to toxicology (toxicogenomics) has rapidly spread to the impact  
21 assessment of chemicals, mixtures and effluents towards the whole ecosystems (with particular regard to  
22 water matrices), thus turning the research field into ecotoxicogenomics. This novel discipline, by  
23 investigating transcripts, proteins and metabolites, overcomes several gaps inherent to the traditional  
24 approach, such as long response time and relationships between exposure duration and possible adverse

1 effects. Meanwhile, it is possible to gain information on basic biology of organisms, also highlighting  
2 common patterns of modes of action [42], [43].

3 The identification of gene expression mechanisms due to stimulation of natural hormones and  
4 xenobiotics has been studied by means of DNA microarrays. Research has been focused mainly on the  
5 estrogen nuclear receptors, which behave as transcription factors, *i.e.*, they interfere with the DNA  
6 transcription process. On the contrary, knowledge of the response elements in gene promoter regions is  
7 still lacking [44], [45]. Transcriptome differs from proteome, due to post- translational modifications;  
8 each environmental stimulus affects these mechanisms, as well as gene expression. Thus, the challenge  
9 is finding the link between the “protein expression signatures”, which are constituted by biomarker  
10 patterns and the modes of actions of chemicals. At the same time, however, the physiological levels,  
11 from the sub-cellular up to the organism, must be scrutinized to investigation, to avoid collecting a huge  
12 amount of protein sequences without getting any relative response [9], [46].

13

#### 14 **3.2.4 Oxidative stress**

15 A detrimental response generated by pollutants on a cellular level includes the production of reactive  
16 oxygen species (ROS) and free radicals. These “basic” processes can be associated with carcinogenesis,  
17 immunotoxicity, teratogenesis and genotoxicity.

18 Although oxidative processes and the subsequent generation of free radicals are normal in the cellular  
19 metabolism of organisms [47], oxidative stress is a condition of imbalance between the antioxidant  
20 defense and the production of ROS, so that the defense is overcome by the formation of radicals [48].

21 This process may cause oxidative damage to membrane lipids, DNA and proteins, and lead to cellular  
22 dysfunction and tissue injury [49], [50].

23 Oxidative stress can be induced through different mechanisms. They may affect the redox cycle by  
24 donating electrons to or withdrawing electrons from cell components. During metabolism, they may

1 deplete glutathione (endogenous antioxidant) or even inactivate other endogenous antioxidants [51]. In  
2 short, oxidative stress can act either through overproduction of free radicals or alteration in antioxidant  
3 homeostasis [52]. Indeed, a close relationship was described between metal cytotoxicity, total GSH  
4 content and the dissociation energy of the sulphur-metal bonds, confirming the involvement of  
5 antioxidant defense mechanisms in the toxic action of these ions [53]. Oxidative stress is also due to the  
6 alteration of antioxidant enzymes such as glutathione peroxidase (GPx), glutathione reductase (GR),  
7 glutathione-S-transferase (GST), catalase (CAT) and superoxide dismutase (SOD), which may lead to  
8 elevated lipid peroxidation [54], [55], [56]. Increased concentration of plasma and red blood cell  
9 thiobarbituric acid reactive substances (TBARs), changes in the antioxidant status, and altered activities  
10 of cellular enzymes such as superoxide dismutase (SOD) and catalase (CAT) indicated higher oxidative  
11 stress in pesticides sprayers. Hence, many researchers have associated exposure to pesticides with  
12 oxidative stress [57].

13 Biomarkers can be chosen based on the biological damage they are linked to. For instance, membrane  
14 disruption might be associated with malondialdehyde, ethylene and ethane, isoprostanes concentration;  
15 ROS production affects glutathione, photosynthetic pigments, total phenolics content. Other biomarkers  
16 may indicate more general phenomena, such ageing, decay and cell integrity (putrescine, spermidine,  
17 spermine) and undetermined stressors (proline).

18 For such a complex scenario, it is essential to use an array of biomarkers to detect oxidative stress.  
19 Different antioxidants are involved in the protection against ROS through a close cooperation between  
20 them, and antioxidant defense may respond differently depending on the species used [58]. Hence, at  
21 least one marker of oxidative damage should be measured in order to make inferences about oxidative  
22 stress [58]. Previous studies have shown that antioxidant enzymes, particularly GPx, CAT and SOD, as  
23 well as lipid peroxidation may function as useful biomarkers of metal induced effects on the antioxidant  
24 system in different bird species [59], [60], [61]. Further studies on other taxa will help to better

1 understand the mechanisms of metal toxicity in wild birds, and the concentrations prone to cause effects  
2 on the antioxidant system.

3

### 4 **3.2.5 Antibiotic resistance**

5 Diverse spectres of antibiotics (chemotherapeutic agents) are being developed to protect humans and  
6 animal husbandry against bacterial, fungal and protozoan infections. Large environmental releases are  
7 caused by their intensive use and, often, overuse or misuse. Furthermore, it is worth noting that most  
8 antibiotics can be only partially metabolized after administration, and, thus, are excreted directly into the  
9 wastewater. Main hotspots are soils fertilized with manure runoff water from farms [62], effluents of  
10 drug production units [63] [64], WWTP effluents and sludge and, consequently, the receiving  
11 waterbodies ([65], [66], [67]). Since microorganisms also naturally produce antibiotics, they are  
12 ubiquitously and perpetually present in ecosystems. Consequently, bacterial resistance to naturally  
13 occurring antibiotics is common in nature. However, the increased presence of antibiotics in the  
14 environment increases the risk of development and spreading of antibiotic resistance [68], also to human  
15 pathogens [69], which has become of great global concern in clinical environments. WWTPs are  
16 considered as important reservoirs for antibiotic resistance genes (ARGs) since sewage receives gut  
17 bacteria previously exposed to antibiotics [70].

18 Antibiotic resistance is mechanistically based on inactivation or modification of the antibiotic, an  
19 alteration in the target site of the antibiotic that reduces its binding capacity, a modification of the  
20 metabolic pathways to circumvent the antibiotic effect or a reduction in the intracellular antibiotic  
21 accumulation by decreasing the permeability and/or increasing the active efflux of the antibiotic [69].  
22 Acquisition of antibiotic resistance may occur by mutation of its own genes (vertical evolution) or by  
23 acquiring new genes from other strains or species (horizontal gene transfer) [71]. The latter is mediated  
24 by so-called mobile genetic elements (MGE) such as phages, plasmids, integrons and transposons.

1 The pool of genetic material maintained by the environmental bacterial communities, named the  
2 resistome, provides the molecular functions for protecting bacteria against most classes of clinically  
3 important antibiotics and constitutes a reservoir of ARGs that can be mobilized into human pathogenic  
4 bacteria [72], [73]. ARGs have gained increasing attention in recent years [74], [68], [69], [75]; there is  
5 still a critical lack of knowledge about the diversity, distribution and origin of resistance genes [76],  
6 especially for the unculturable majority of environmental bacteria, of which less than 1% are estimated  
7 to be culturable [77].

8

9 **4. Definition of assays for testing ecotoxicity: a focus on “lab-life”**

10 Different biological responses or stresses potentially caused by micropollutants present in wastewater  
11 (see Section 2) can be assessed in lab-scale tests. Such assays have to be able to simulate the actual  
12 conditions occurring in case of receiving waterbodies and reused waters. Due to the challenges in  
13 collecting representative samples without losses and inherent high costs for conducting proper toxicity  
14 assessments, a well thought-through sampling strategy and sample collection and preparation are of  
15 major importance. Figure 3 shows the main factors, which must be considered before planning a  
16 monitoring by adopting the bioassays.

17

18

19 Figure 3. Factors to be accounted for when selecting the proper bioassay.

20

21

22 **4.1. Sampling strategy**

23 Chemicals present in wastewater can considerably vary as a function of time, urban populations and  
24 treatment plant performance. All these factors can be dramatically different from site to site. For instance,

1 the hydraulic retention time (HRT) and sludge retention time (SRT) are highly dependent on plant design  
2 (e.g. type and size of treatment units and internal flow patterns, including sludge treatment and reject  
3 water) and changes in volumetric loading (e.g., due to storm water intrusion). When assessing acute  
4 toxicity, a worst-case scenario would usually be appropriate. However, chronic effects would better  
5 require average or median exposure conditions.

6 If possible, (considering the storage time constrains) composite samples are preferred, taken by means  
7 of automatic samplers, usually collecting a volume aliquot every 10 min over a certain period, typically  
8 24 hours. This would normally cover at least 1 HRT at most WWTPs, and it would be within the  
9 maximum recommended storage time (if stored refrigerated) for the most relevant compounds [78], [79].

10 In case of a longer sampling period, more 24-hours composite samples can be summed, possibly after a  
11 pre-treatment (see below). When planning the monitoring campaign, depending on the final goals, the  
12 expected weekly and seasonal variations may be taken into account [80], as well as the conditions of  
13 receiving waterbodies (see Chapter 5). For instance, the concentrations of some illegal drugs have been  
14 found to increase towards the weekends or in relation to popular events which draw the crowds [81]. The  
15 economic aspects are crucial, and often decisive, in setting the total number of samples to acquire. Hence,  
16 for the acute toxicity assessments, uncertainty is marked, if case campaigns do not cover the actual worst  
17 case, specifically in case of the investigation of the combined effects of all substances contained in  
18 sewage. Furthermore, routinely detected chemicals often cannot explain the observed biological  
19 responses (e.g. [24]) pointing to a mismatch in the assessment approaches generally applied. Therefore,  
20 it may be difficult to pinpoint the best or most appropriate term to conduct the sampling campaign.

21 Ort et al. [82] detailed and explained all these aspects, focusing on sample volumes, collection duration,  
22 storage conditions and data elaboration.

23

## 24 **4.2. Sample collection and preparation**

1 Several factors can undesirably influence the composition of wastewater from collection to analysis: i)  
2 Compounds may be adsorbed to or diffuse from the sampler tubing and container. Most of the larger  
3 WWTPs have an automatic sampling equipment installed, which should preferably be used to minimize  
4 these effects. ii) Depending on the WWTP scheme, the compounds of interest in effluent water samples  
5 will usually be in  $\mu\text{g/L}$  -  $\text{pg/L}$  concentration range together with solids, microorganisms, beside a great  
6 number of dissolved or suspended compounds in the  $\text{mg/L}$  concentration range. Biodegradation  
7 preferably occurs in raw sewage and in the effluents, with respect to the receiving waterbodies. Hence,  
8 it is important to limit both biotic and abiotic processes after sampling. Sterile filtration ( $<0.2 \mu\text{m}$ ) is an  
9 efficient way to stop biotransformation, though enzymes may still be present. It is also necessary prior  
10 to solid-phase extraction (SPE) to prevent clogging (see below).  $0.45 \mu\text{m}$  filters are more commonly used  
11 since they are less prone to blocking. Anyway, the choice of filter type is also crucial: polycarbonate or  
12 (low static charge) cellulose acetate filters may be preferred as *e.g.* nitrocellulose filters tend to bind  
13 proteins while nylon filters tend to bind proteins, DNA and RNA. Acidification with HCl or  $\text{HNO}_3$  is  
14 often adopted, alone or after filtration, to preserve the sample. However, this may alter the speciation and  
15 stability of the compounds and, therefore, should be applied with care [83]. iii) Micropollutants in the  
16 filtered sample are usually cleaned up and concentrated prior to application in the toxicity tests by step-  
17 wise SPE and elution. The composition and concentration of the eluate depend on parameters such as  
18 physico-chemical properties of the compounds themselves, type of SPE, sample volume and percolation  
19 rate, sorbent cartridge volume, type of elution solvent and elution volume (see [83] for a more detailed  
20 discussion).

21 Ideally, the whole, unaltered sample should be used in toxicity tests. However, this is not possible for  
22 practical reasons for most of the tests (see Section 3.2). Some sorts of sample clean up and concentration,  
23 as briefly described above, are usually needed. If only the filtrate is submitted to toxicity tests, the effects  
24 of hydrophobic compounds and other pollutants adsorbed to the particulate matter will probably be



1 underestimated. Moreover, if only an extract of the filtrate is tested, even more compounds may be lost.  
2 To improve the recovery of micropollutants from the original sample, the substances retained by the filter  
3 might also be extracted using the same solvent as applied for SPE elution.

#### 4 5 **4.3. From research to standards: multifaceted approach in bioassays**

6 Several national and international authorities and scientific and technical organizations are instrumental  
7 in compiling and evaluating toxicity tests such as the Organisation for Economic Cooperation and  
8 Development (OECD), WHO, Food and Agriculture Organization (FAO), International Organization for  
9 Standardization (ISO), American Society for Testing and Materials (ASTM), USEPA, United States  
10 Army Corps of Engineers (USACE), American Public Health Association (APHA), Association  
11 Française de Normalisation (AFNOR), Deutsches Institut für Normung (DIN), Italian Association for  
12 Standardisation in the Chemical Sector (UNICHIM). The level of worldwide methods harmonization is  
13 sometimes limited; thus a plethora of standard protocols exist with overlapping normalisation actions  
14 that sometimes can be conflicting in terms of sensitivity, meaning that each protocol has its own  
15 feasibility. Time-by-time authors must clearly declare which method they follow, to assure data  
16 reproducibility and correct interpretation.

17 Table S1 (Supplementary material) reports a list (necessarily not exhaustive) of the most commonly  
18 adopted toxicity tests, by pointing out the issuing agency and the measured response at the cell, tissue,  
19 organ, organism and ecosystem level. The principle of adverse outcome pathways, AOP, [84] allows to  
20 start from the initiating event, which possibly causes an adverse effect and to explore the whole biological  
21 pathway, up to the ecosystem level, by following a mechanistic approach. Endocrine activity testing is  
22 an example of such an application. The available assays can highlight both the interference with the  
23 hormone receptors, by means of agonistic/anti-agonistic activities, and, more generally, the interference  
24 with hormone synthesis and release.

1 The architecture of an assay involves simple cases, such as the mere formation of a bond between a  
2 ligand (either radio-labeled or bound with a fluorochrome) and an isolated receptor (thus gaining only  
3 analytical information). More complex cases are based on specific endpoints, such as protein activity  
4 (both in terms of protein synthesis and protein interactions with co-factors), cell proliferation and direct  
5 receptor activation linked with a gene reporter. Tests employ different techniques for the quantification  
6 of the biological activity. They range from basic approaches with UV-VIS spectroscopy, to the most  
7 exploited tools like ELISA, radio-immunology, and fluorometry (including flow cytometry) [38], [85],  
8 [86].

9 Reporter gene-assays involve the use of cells (deriving from bacteria, yeasts, fish, humans and other  
10 mammals) to assess gene expression mediated by chemicals. The endpoints consist in cell proliferation  
11 in case of E-SCREEN [86], [87], [88], [89], while, most assays are based on gene expression, often after  
12 specific transfection. The main testing tools are the following: CALUX, CAFLUX, PALM, MELN,  
13 MVLN, T47D-kBluc, HELN, HGELN, MDA-kb2, PR reporter gene assay, YES, YAS, BLYES,  
14 BLYAS, BLYR (luciferase/fluorescent protein gene expression,  $\beta$ -galactosidase synthesis induction);  
15 they measure the binding with estrogen, androgen, progesterone, glucocorticoid, peroxisome proliferator  
16 activator receptors [86], [90], [91], [92], [93], [94], [95], [96], [97]. Other tests are focused on the  
17 quantification of the production of proteins, such as vitellogenin, choriogenin, *zona radiata* protein after  
18 estrogenic stimulation [98], [99], [100], [101]. Steroidogenesis based tests look promising in providing  
19 additional information on disruption mechanisms [101], [102] although *in vivo* compensation of the  
20 effects which occur during *in vitro* assays is far from being defined. Among the *in vivo* assays aimed to  
21 evidence the gene expression induced by pollutants, there is the application of the genetically modified  
22 *Danio rerio* (green fluorescent protein expression, controlled by a thyroid hormone response promoter  
23 of *Xenopus laevis*) already applied to environmental samples [103], [104]. An *in vitro* reporter gene assay  
24 (ER $\alpha$ -luc assay) can be used for estrogen receptor activation to quantify the total estrogenic activity in

1 liquid samples. Extracts from environmental samples (e.g. in petroleum ether) can be used to measure  
2 the estrogenic activity with a reporter gene assay (ER $\alpha$ -luc assay) based on U2OS-ER $\alpha$  cells, with  
3 luciferase as reporter [105]. The method to culture and expose the cells and to assay luciferase activity  
4 has previously been described in [106]. Measurement of the estrogenic activity of nonylphenol during  
5 biological degradation showed a decrease of the estrogenic activity during microbial degradation and can  
6 hence be used to determine the ecotoxicological risk of an environmental sample. An overview of pros  
7 and cons of the main assays applied for ecotoxicological purposes, in case of water ecosystems, together  
8 with basic technical information is reported in the paper recently published by Brack et al. [12].

9 Besides, the study of the effects of EDs on populations and communities requires the setting up of  
10 mesocosm assays or the direct observation of real scenarios. It is worth noting that the pollutants (and  
11 mixtures) are effective, in parallel, at increasing levels, up to the ecological aspects, hence yielding to  
12 significant changes on the trophic web. This is the case, for instance, for *R. rutilus*, a planktivorous fish,  
13 whose grazing capacity is deeply reduced by the exposure to EE2; the population of plankton, as a result,  
14 can undergo a development [107].

15 Among the agencies and organizations which are facing the issue of endocrine disruption, the OECD  
16 approach can be cited, since it prescribes further subsidiary levels of investigations, in order to draw a  
17 complete profile of endocrine disruption [108]. The five levels consist of: 1) acquirement of existing data  
18 about chemical, physical and toxicological properties, 2) execution of *in vitro* assays aimed to highlight  
19 endocrine pathways, 3) execution of *in vivo* assays aimed to highlight endocrine pathways, 4) execution  
20 of *in vivo* assays aimed to highlight adverse effects on endocrine endpoints, 5) execution of *in vivo* assays  
21 aimed to highlight adverse effects on endocrine endpoints throughout the whole life of an organism and  
22 across generations.

23 As far as genetic toxicity is concerned, the assays proposed in the scientific literature, the international  
24 standards (mainly OECD and ISO) and available on the market (automatized, in most cases) allow

1 highlighting and quantifying multiple effects, from early, hence reversible modifications of genetic  
2 material, up to irreversible damages, which can evolve to either apoptosis or neoplastic formations.  
3 Therefore, the assays can be usefully integrated in a multi-layer frame, also due to the option of testing  
4 organisms of growing biological complexity (prokaryotes and eukaryotes) and situated at different levels  
5 of the trophic web (producers, consumers and decomposers). The detection of genetic damage induced  
6 by various mechanisms is made possible by performing *in vitro* and *in vivo* tests. The endpoints are  
7 represented by a) gene mutations; b) chromosomal damage (to parts of the chromosomes); c) genomic  
8 damage (loss/gain of entire chromosomes) c) epigenetics.

9 Among the large number of tests either standardized or just proposed for the evaluation of water matrices,  
10 it is worth mentioning: a) the Ames test, for detecting point mutations in *S. typhimurium* bacterial strains;  
11 it is based on the growth of histidine revertant bacteria over specific culture media, with or without the  
12 addition of rat liver microsomal fractions. It is the most applied in case of environmental evaluations  
13 [109], [110], [111], [112], [113], [114]. It takes 48 hours to obtain a result. b) The micronuclei test, for  
14 detecting chromosomal mutations (generally performed on root cells of *Allium cepa*, throughout 72 h)  
15 [109], [112], [113]; it is a biomarker of chromosomic damage and genome instability. Its exposure  
16 depends on the employed organism. c) The Comet assay (also called SCGE, Single Cell Gel  
17 Electrophoresis), for quantifying the primary DNA damage; it is typically carried out on eukaryotic cells  
18 [109], [110], [113], [114], [115]. d) The reporter gene assays, which detect the SOS response induced by  
19 DNA damage and have a duration of several hours; often automatized, they are less sensitive and robust  
20 than the aforementioned tests [111], [116]. e) The GreenScreen assay (GSA) which employs cells of  
21 *Saccaromices cerevisiae*; it detects a DNA damage, based on the quantification of a green fluorescent  
22 protein linked to the promoter of the RAD54 gene [117], [118]. f) The sister chromatide exchange (SCE  
23 assay) based on mammal cells [115], [119].

1 Traditionally a wide number of enzymes, known to be involved in reactions against pollutants, are  
2 employed in toxicity tests. Unfortunately, in several cases enzymes react by means of induction or  
3 inhibition mechanisms, without a direct connection to the chemistry (e.g. leaving groups, electrophilic  
4 or nucleophilic functions) of the specific pollutants. Moreover, it is well-known that the effects of  
5 chemicals can be disguised by the action of several environmental factors, such as the feeding regime,  
6 temperature, water chemistry, as well as biological aspects, including population genetics, reproductive  
7 cycles [120]. Enzymes may rarely induce general stress rather than detoxification.

8 Therefore, it is important to clearly denominate the purpose of the assay in the frame of the toxicity  
9 testing. Enzymes like SOD, CAT, APOX, DHAR, MDHAR, GPOX and GR are members of the  
10 Halliwell-Asada-pathway [48] detoxifying radicals and toxic oxygen species that might build up under  
11 xenobiotic stress [121].

12 Enzymes of the metabolic cascade of xenobiotics, like the P450 and POX, as examples for phase I, would,  
13 on the contrary, act on the xenobiotic directly and activate it by inserting –OH groups into the molecule.  
14 Similarly, in phase II, GST and GT would conjugate glutathione or glucose to the activated xenobiotic,  
15 thereby detoxifying it [122]. However, there are also examples of direct attacks towards the pollutant, as  
16 for P450 and diclofenac or acetaminophen, and GST and lamotrigine.

17 Despite these differences in function, the mentioned enzymes are inducible by xenobiotics, and might  
18 exhibit elevated levels of activity in the respective assays. Table S2 (Supplementary material) lists the  
19 main enzymes employed in bioassays.

20 Among pharmaceuticals, antibiotics give seriously cause of concern, due to their indirect adverse effect  
21 on human health linked to the phenomenon of bacterial resistance. In clinical microbiology standardized  
22 susceptibility tests they clearly dominate among the available methods, aimed to detect possible drug  
23 resistance in common pathogens and to assure susceptibility to drugs for a particular infection [123]. In  
24 these tests, resistance is detected by carrying out growth inhibition tests broth (e.g. the macrobroth

1 dilution test and the miniaturised broth dilution test) or by agar diffusion (*e.g.* the gradient diffusion test  
2 and the disk diffusion test). In most of these tests (except the disk diffusion test) the lowest concentration  
3 of antibiotic that prevents growth, represented by the minimum inhibitory concentration (MIC), is  
4 quantified. A more detailed discussion of advantages and drawbacks of these methods is given by [123]  
5 and [124]. Such culture-based approaches typically require 1-2 days for fast-growing bacteria like  
6 *Escherichia coli* or *Salmonella* spp., and several weeks for slow-growing bacteria, like *Mycobacterium*  
7 *tuberculosis*. However, the main drawback of cultural methods is that the vast majority of strains present  
8 in environmental microbial communities (<1 %; [77]) still cannot grow outside their host environment.  
9 Assessment of antibiotic resistance in such communities based solely on cultivable bacteria will therefore  
10 easily generate unrepresentative and biased results [125].

11 For that reason, tools for molecular detection of antibiotic resistance genes (ARG) have become  
12 increasingly popular [69], [70], [126]. Polymerase chain reaction (PCR) assays such as multiplex PCR  
13 and quantitative real-time PCR (qPCR) have frequently been applied to amplify and detect specific ARGs  
14 in environmental samples [70]. Nevertheless, they only target well-studied pathogens or resistance-  
15 causing genes (as the primers are based on known resistance genes only) and cannot easily be used for  
16 broad-spectrum screening [69]. DNA microarray is a more powerful molecular method than the PCR  
17 assays as it is able to detect the presence or absence of a large range of ARGs simultaneously in a single  
18 assay [126]. However, its use for environmental samples has been limited as it is hampered by low  
19 detection limits (partially overcome if coupled with PCR) and the need for complicated pre-treatment to  
20 reduce the presence of other compounds that inhibit DNA extraction and/or target gene amplification  
21 [70]. Furthermore, both microarray and PCR based technologies are not conclusive regarding the  
22 detection of resistance genes in metagenomes [127].

23 Metagenomic analysis is one of the latest modern approaches for analysing complex microbial  
24 communities and enables to describe the genetic potential of a community and to detect the

1 presence/absence of genes or genetic variations responsible for antibiotic resistance [69]. Metagenomic  
2 analysis usually follows two different approaches, namely sequence-based and functional. In the first  
3 case, a sample of DNA from the studied metagenome is extracted and completely, but randomly,  
4 sequenced in relatively short contiguous sequence read lengths. These sequences are then compared with  
5 known sequences that have accumulated over the years in public databanks (reference sequences; e.g.  
6 [128]) to identify resistance genes and/or mutations that are known to cause resistance [69]. This  
7 approach has the potential to identify all known resistance genes in a metagenome. Though, important  
8 shortcomings are that it can only identify known ARGs and that it gives no information on expression of  
9 the resistance genes [127]. This is, however, overcome by the second approach, functional  
10 metagenomics, in which the extracted DNA is shot-gun cloned into cloning vectors and subsequently  
11 expressed in a cultivable surrogate host (usually *E. coli*) plated onto antibiotic-containing agar. If  
12 bacterial artificial chromosomes (BACs) are used, a larger gene fragment can be inserted, potentially  
13 making it possible to trace the phylogenetic origins of the original host bacteria [127]. These larger gene  
14 fragments are also more likely to include antibiotic resistance that is encoded by multiple genes.  
15 Disadvantages of using BACs is the low copy number (though, they are usually more stable than higher  
16 copy vectors) and the need for the transcription and translation signals to be efficiently recognized by the  
17 host organism. If vectors that only accept small inserts are used, the copy numbers are higher and the  
18 host's transcription and translation systems can be used, hence the drawbacks of using BACs are  
19 circumvented. However, the small size of the insert will not normally allow information about the genetic  
20 background of the resistance gene. However, if coupled with sequence-based metagenomics, this  
21 disadvantage can be overcome to some extent. See [127] and [69] for a more thorough discussion of  
22 advantages and drawbacks in metagenomics analyses.

23

#### 24 **4.4 Criteria for selecting a bioassay**

1 Toxicity tests can be prescribed by the law or by voluntary regulations and, consequently, be completely  
2 standardized. Also the results interpretation, in this case, is almost plane. On the contrary, when the final  
3 goal is a deeper investigation of the impact of effluents (or chemicals/mixtures) on specific biological  
4 targets, at different levels (from sub-cellular components to the whole community) several alternative  
5 choices arise. The criteria underpinning the selection of a bioassay (or a battery of complementary assays)  
6 should include the required volume (smaller volumes may favor the miniaturization, hence the  
7 automation of the procedure), the price (consisting in capital and operation costs, as well as in the license  
8 fee), the throughput, the sensitivity (by taking into consideration possible non-monotonic responses), the  
9 specificity, the requirement of trained and skillful operators, the possibility to measure  
10 acute/chronic/transgenerational effects, the capability of evidencing toxicokinetic or, more generally,  
11 specific metabolic pathways of interest.

12 A pivotal role is played by the personnel cost, which differs highly among the countries: the European  
13 example is revealing, varying the minimum wage per month from less than 250 € for Albania, to nearly  
14 2,000 € for Luxembourg [129]. Furthermore, the same test (e.g., Ames on *S. typhimurium*) can be carried  
15 out either by adopting the conventional microbiological approach and cultivating bacteria or using kits  
16 available on the market. Among the options, there is also the possibility to use genetically modified  
17 microorganisms, which allows eliminating the addition of rat liver S9. Therefore, time is reduced (hence  
18 the analyst costs) but CAPEX and OPEX become significant.

19 Besides, also these two items can assume different commercial values: metagenomics assays are getting  
20 progressively less expensive, but the number of clones and libraries increases rapidly.

21 Finally, in any case, before planning a monitoring campaign focused on the assessment of toxicity of a  
22 sewage/effluent/mixture, one has to pay attention to possible species-specific responses and to the  
23 inherent criticism of *in vitro* assays, whose results cannot be translated as a whole to a real situation, due  
24 to the occurrence of compensation effects and the multifactorial nature of toxicity.



1

## 2 **5. Environmental risk assessment: challenges and limitations**

3

### 4 **5.1 Traditional environmental risk assessment**

5 Environmental Risk Assessment (ERA) deals with the interactions of agents or hazards, humans and  
6 ecological resources. It describes human populations, ecological resources and agents, analyses agents  
7 and exposure potential, characterizes the potential for adverse effects, defines uncertainties, generates  
8 options to deal with the risks, and communicates information about the risks to humans and ecosystems.

9 ERA is a process that evaluates the likelihood or probability that adverse effects may occur to  
10 environmental values because of human activities (i.e., a formal procedure for identifying and estimating  
11 the risk of environmental damage). ERA provides information for making reasoned decisions by defining  
12 the range of risks associated with various options, but it does not dictate a specific outcome. ERA also  
13 provides a mechanism for managers to communicate forecasted risks associated with decisions, such that  
14 stakeholders and the public are informed of the implications for environmental values.

15 Based on the toxicological data and measured environment concentrations found in the literature, the risk  
16 for acute toxic effects is unlikely but chronic adverse effects cannot be excluded. Therefore, risk  
17 characterization is one of the important tools to estimate the environmental risk, particularly in view that  
18 co-occurrence of diverse micropollutants in environmental matrices may lead to additive, synergistic,  
19 and antagonistic toxic effects which is difficult to predict if only concentration is available.

20

### 21 **5.2 Calculation of predicted no-effect concentrations and risk assessment**

22 Risk calculation is an essential step for quantitative evaluation of risk from chemical exposure. Risk  
23 characterization involving exposure and effects using effects-related Environmental Quality Standards

1 and Quality Norms is a single-substance concept for dangerous substances and priority hazardous  
2 substances. The environmental risk posed by certain trace pollutants in aquatic ecosystem is  
3 characterized and assessed through calculation of the risk quotients (RQ). Therefore, the hazard quotient  
4 method is commonly used for screening level risk characterization. Namely, under EU legislation  
5 context, guidelines have been developed for new or existing substances as well as compounds, such as  
6 biocides or hazardous substances [130], [131], [132], [133], [134], [135]. Moreover, this approach has  
7 been generally accepted and has widely been used all over the world: Taiwan [136], [137], Japan and  
8 China [64], [138], [139], [140], Germany [141], United Kingdom and the USA [142], Spain [143], [144],  
9 Korean [145] etc. for environmental risk assessment. The RQ is calculated for each micropollutant by  
10 dividing the measured environmental concentration (MEC) by the predicted no effect concentration  
11 (PNEC - the concentration at which no adverse effect is suspected to occur) using the lowest value of all  
12 endpoints, as given in Eq. 1:

$$13 \quad RQ = MEC/PNEC \quad (1)$$

14 PNEC is calculated (Eq. 2) by dividing the lowest chronic no observed effect concentration (NOEC) by  
15 the assessment factor (AF) or uncertainty factor (UF) chosen according to European guidelines [146].

$$16 \quad PNEC = NOEC/AF = EC_{50} \text{ or } LC_{50}/AF \quad (2)$$

17 This method recommends the use of chronic toxicity data for calculation of PNEC. However, for some  
18 compounds, the availability of chronic data referring to NOEC for some compounds is limited. In absence  
19 of NOEC values, acute toxicity data could be used and where no experimental toxicity data are provided,  
20 the toxicity values could be predicted using QSAR (Quantitative Structure Activity Relationship) models.  
21 An assessment factor (AS) is applied to the toxicological benchmark value according to [134], laying  
22 down the principles for the assessment of risks to man and the environment of existing substances in

1 accordance with Council Regulation [133].

2 AF or UF of 1000 is applied when at least one short term L(E)C<sub>50</sub> from each of the three evaluated trophic  
3 levels are provided; an AF of 100 is used when one long term NOEC was available for either cladocerans  
4 or fish in case of pharmaceutically active compounds, and finally AF of 50 is used when two long term  
5 NOEC values are provided for species in two different trophic levels. When the MEC is larger than the  
6 PNEC, giving RQ larger than 1, the investigated MP is risky to human health or the environment, while  
7 a RQ value of 0.01-0.1 and 0.1-1 indicates low or medium risk, respectively. If  $1.0 < RQ < 10$ , some adverse  
8 effects or moderate hazard are probable, and if  $RQ > 10$ , high hazard is anticipated [146].

9 Although risk assessment is a useful tool, the results should be taken with attention, due to several factors.  
10 Single compound exposure scenario, indeed, is unrealistic in the real aquatic environment, since trace  
11 pollutants are present as complex mixtures. They often exhibit toxic effects at concentrations lower than  
12 the NOEC values for each compound if acting alone [147].

13 The overall effect is synergistic, additive or subtractive, depending on each specific mode of action; a  
14 compound might also behave completely independent from the others [148]. Also, if the exposure data  
15 used for calculation are based on single sampling events, these data do not accurately represent the long-  
16 term exposure of the organisms present in the studied aquatic matrices. This should be taken with caution  
17 when chronic exposure is estimated for the calculation. Nevertheless, the used exposure data only  
18 represent concentration in water, and, besides, factors like as bioconcentration or bioaccumulation, real  
19 uptake rates should be considered as well. Additionally, available dose-response data in the case of  
20 pharmaceutically active compounds, which are the most studied, are limited, especially regarding chronic  
21 data. Furthermore, the toxicity data provided by QSAR models for fish, algal and *Daphnia* spp. are for  
22 chemicals with non-specific mode of action and hence, should also be applied with caution. On the other  
23 hand, information on compounds for which uncertainties about potential endocrine disrupting effects  
24 exist is still needed. Jones et al. [149] when evaluating chemical mixture toxicity on sites where multiple

1 chemicals of potential ecological concern can simultaneously occur used HQs, which are commonly  
2 referred as Toxic Units (TUs) in order to estimate the relative toxicity of the waters from each site. Thus,  
3 the sum of TUs for each sampling point is calculated by means of the sum of chemical-specific HQ for  
4 each site. As in the case for individual HQ, the sum of TUs should be taken as preliminary for the quotient  
5 addition approach assuming that toxicities are additive or approximately additive. In the case of some  
6 pharmaceuticals similar modes of action of compounds in the same therapeutic group could account for  
7 a certain level of overestimation of the risk. However, at the same time, possible synergic interactions  
8 among compounds with different modes of action could underestimate the risk. Although the risk  
9 assessment scenario and the results obtained using this tool could be accepted with caution due to the  
10 mentioned limitations, its applicability demonstrates the need for water quality monitoring using  
11 composite and periodical samplings, since the discharge of trace pollutants and their metabolites is time  
12 dependent, i.e., fluctuating daily, monthly or seasonally. Nevertheless, the results could show the need  
13 for improving of sewage treatment technologies and could generate concern about the efficacy of the  
14 applied water treatment.

15

### 16 **5.3 Wastewater toxicity assessment and ranking**

17 The problem of wastewater toxicity data management and interpretation is still a current issue, especially  
18 when high toxicity levels are recorded and there are compulsory legislative threshold limits to comply  
19 with [150], [151], [152]. Around the world, countries have developed various toxicity-based methods to  
20 assess the quality of treated wastewater to increase the accessibility to water and sanitation in order to  
21 avoid human health impacts and ecosystem services impairment. Several procedures for discharge hazard  
22 estimation have been proposed generating assessment toolboxes including limit-based threshold  
23 approaches, and toxicity score and index for data integration and interpretation including expert judgment

1 as well [150]. The main goal of wastewater ecotoxicity assessment and ranking should be to minimize  
2 the adverse impact onto the receiving water body as well as treated wastewater recovery and reuse [153].  
3 Apart from the possibility of using toxicity tests to estimate potential hazardous effects on the ecosystem,  
4 they can favour the protection and the optimization of wastewater treatment plant operation, by  
5 discriminating the best available technologies [154], [155]. Consistent wastewater toxicity assessment  
6 can increase the general level of sustainability in the management of water resources pushing ahead both  
7 “zero emission” and “zero discharge” along with the precautionary principle [156].

8 Toxicity is currently used to check effluent quality into various national legislation around the world to  
9 be included in water monitoring and control programs like direct toxicity assessment [157], whole  
10 effluent toxicity, integrating controlling of effluent, whole effluent environmental risk, environmental  
11 effects monitoring [158], and whole effluent assessment [156], [159]. Apart from any program  
12 peculiarities, the main question is still how to use or “interpret” toxicity data keeping in mind that the  
13 objective is to protect the environment and not the “white rat” testing species [160].

14 Generally, legislative requirements tend to refer to a toxicity limit based on a single test or a battery of  
15 toxicity tests considering as final result the worst registered data. This method is quite simple, but not  
16 environmentally realistic, depending on the biological model-endpoint pairs considered and the weight-  
17 of-evidence score attributable to each of them. Sometimes, the classification is attributed just on a  
18 logarithmic [161], [162] or order of magnitude basis [163], [164], [165], [166] or expert judgment and  
19 regression analysis pair [167]. Some authors tried to overcome such drawbacks by identifying tools to  
20 integrate and weight toxicity data on a statistical basis also according to the ecological relevance of the  
21 considered endpoint [150]. For example, Libralato et al. [150], [151] applied the minimum significance  
22 distance (MSD) criterion to support general decisions about the presence or absence of toxicity from  
23 wastewater samples on a database of more than 100 wastewater toxicity data including domestic,  
24 municipal and industrial discharges [168], [169]. This method enabled the consideration on a species-

1 specific basis. Thus, the relative sensitivity of the biological model made the assessment of toxicity  
2 independent to reference wastewater as well. Moreover, expert judgement was reduced to a minimum  
3 just in relation to the choice of the number of ranking classes and their extension in case of more toxic  
4 samples. This kind of approach produced a toxicity score with classes (absent, low, medium, high and  
5 very high toxicity) composed of two sub-scores. The first series of sub-scores (absent or low toxicity)  
6 was partly based on the percentage of effect responses and partly on toxic unit values. The second series  
7 of sub-scores was entirely defined on toxic unit values including a medium, high and extremely high  
8 toxicity threshold. The main limits of this approach are related to the fact that each toxicity score is  
9 species-specific and databases including wastewater toxicity data must be developed *ad hoc* also to  
10 support the data statistical reliability.

11 Further efforts are necessary to identify case-specific toxicity tests (country- or discharge-based),  
12 supporting their round robin and toxicity data integration methods in the perspective of EU legislative  
13 harmonization.

#### 14

#### 15 **5.4 How reliable is our risk assessment in the receiving water bodies?**

16 Within the Water Framework Directive (WFD) the term “ecological status” of a water body primarily  
17 embraces the biological responses caused by other pollutants than micropollutants, but priority  
18 micropollutants are taken into account through an environmental risk assessment (ERA) scheme by  
19 implementing Environmental Quality Standards (EQS) that should not be exceeded in the environment  
20 [170]. The EQS values are set by each member state based on the predicted no effect concentration  
21 (PNEC) for each compound in water, sediment and/or biota. However, available ecotoxicity data are  
22 often limited, especially for metabolites and transformation products. Therefore, traditional ERA, as  
23 described by the European Commission Technical Guidance Document (TGD), allows the use of  
24 assessment factors (AFs) to account for the uncertainty in deriving PNEC values based on acute toxicity

1 data and a limited number of species [171]. The intention of the use of AFs is to predict a concentration  
2 below which an unacceptable effect will most likely not occur. Data on persistence in the environment  
3 (i.e. lack in biodegradability) and bioaccumulation should also be considered. An AF of 1000 is advised  
4 if only acute toxicity data are available for three trophic levels (algae, daphnids and fish). Only highly  
5 rarely sufficient data on long-term effects at several trophic levels and taxonomic groups exist for a given  
6 compound to be used for statistical extrapolation methods to derive a PNEC value. For biologically active  
7 compounds such as pharmaceuticals, this approach may, however, overlook sub-lethal and subtle  
8 subcellular effects that might occur in some species at much lower concentrations during chronic  
9 exposure. The complexity implied by the cocktail effects of compound mixtures and the large number of  
10 unknown transformation products during degradation in the environment warrants a switch to a more  
11 effects-oriented approach when assessing the environmental risk. Hence, the combined effects from all  
12 compounds in water or sediment samples are assessed using a set of toxicity tests targeting e.g. baseline  
13 toxicity, estrogenic and mutagenic activity and oxidative stress. The main drawback of this effects-  
14 oriented approach is that it is not able to identify the actual compound(s) that are asserting the observed  
15 effects. But if it is combined with the above-described MEC/PNEC (or MEC/EQS), any major  
16 discrepancies between the observed effects and the calculated MEC/PNEC values relevant to the  
17 respective effects may be used to identify “missing” contributing compounds and warrant more detailed  
18 analyses or studies. Still, true food web effects are not covered, leaving the question open whether an  
19 ecosystem hazard may be possible. Discharges from WWTPs are only one of many possible routes for  
20 micropollutants to enter the aquatic environment, and the environmental risk assessment (ERA) of  
21 discharges to a water body should take them all into account. Similar approaches as described above for  
22 the water body may be performed. Instead of measuring the actual environmental concentration (MEC),  
23 the environmental concentration is predicted (PEC) from concentrations in the effluent from the WWTP,  
24 the total discharged volumes and the immediate local dilution in the receiving waters. For compounds

1 that are persistent in the environment and/or bioaccumulate a more long-term and regional assessment  
2 may be needed, including the potential accumulation in sediment.

3 Any industrial agricultural, farming, commercial and recreational activity (including boats and ships), as  
4 well as living units discharging wastewater to water bodies, standing both on freshwater and marine  
5 environments, need to know the nature and the extent of impacts associated with their liquid emissions.  
6 These issues are driving the need for a more detailed assessment of the impact of wastewater discharges  
7 to support decision-making. The integrated assessment of biological effects of discharges in the  
8 ecosystems is relevant and ecotoxicity tests are referred to as extremely useful tools for the identification  
9 of environmental impacts [172]. The use of the ecotoxicology can provide an added value to hazard and  
10 risk assessment of discharges to the receiving water bodies. Environmental management can take  
11 advantage from safe and non-toxic treated wastewater, supporting its recovery and reuse, as in case of  
12 non-potable purposes. Ecotoxicity tests can identify the hazard and be directly used in ecological risk  
13 assessment. Within the WFD, direct toxicity assessment of WWTP discharges can contribute to attain or  
14 keep ecological quality objectives in water masses and finally provide the postulated “good” quality of  
15 all water bodies in the EU.

16

## 17 **5.5 Socio-economic aspects**

18 Monitoring and predicting trace pollutant concentrations in the aquatic environment, together with their  
19 possible subsequent toxicity, are vital to better assess the environmental impact as well as the risks for  
20 human health. Thus, new effective tools for estimating the occurrence of these substances are needed. A  
21 recent method is based on online search queries, though this only applies to those that are widely known  
22 by the public. For example, considering pharmaceuticals, the prescription issuing in the UK of several  
23 substances included in the EU watch list for water monitoring [173] is suggested to be correlated to  
24 online search queries [174]. As the concentrations of antibiotics in wastewater seem to follow the trend



1 of prescriptions [175], search traffic data could be proven a valuable tools in predicting the occurrence  
2 of pollutants in wastewater.

3 The choice of proper removal treatment as well as the overall assessment of its environmental, economic,  
4 and social impacts needs to be assessed with caution [176], and must necessarily take into account  
5 pollutants loads, which, unfortunately, can be affected by extreme variability. Therefore, all the cost  
6 items might be accurately overweighed, to avoid wastes of energy and material resources, land  
7 consumption, and to reduce pollution towards other environmental matrices. Recently, Life Cycle  
8 Assessment (LCA) has been applied to evaluate the economic and environmental viability of processes  
9 aimed to remove trace pollutants from wastewater [177], [178]; this instrument provides standardized  
10 criteria to compare alternative options by taking into account different impact categories.

11 In any case, the effective step towards the reduction of trace pollutants emissions and, consequently, their  
12 effects on the environment is definitely a management at the source. Green chemistry principles [179]  
13 are the essential criteria for designing new production and supply chains, as well as disposal and  
14 treatment. The example of pharmaceuticals is emblematic. Medical professionals and patients should  
15 employ, if possible, products manufactured in accordance with the green pharmacy principles, e.g. using  
16 pharmaceuticals that are designed to be better biodegradable [180], [181]. Disposal of unused medicines  
17 is mostly carried out through household waste [182], toilets and sinks [183], [184], [185]. As many do  
18 not regard this as an environmental issue [182], it is evident, that public awareness is vital, together with  
19 the need for better public information [184]. Over the past decades, attention has also focused on return  
20 policies advertisements [182] and the importance of people information on the correct disposal [182],  
21 [184]. As a consequence, population willingness to pay for a better waste treatment system increases  
22 [183], [186]. Governments should implement the regulatory frameworks for improving the whole water  
23 cycle management [187]. According to the Polluter Pays Principle, environmental damage should be  
24 decreased by introducing advanced treatment technologies, which should be paid by the final users.

1 Therefore, conventional tariff policies aiming to charge all households as a function of wastewater  
2 production are not in accordance with the Polluter Pays Principle. It has been shown, that increased  
3 charge rates and penalties do not contribute to more environmentally friendly practices [188]. Thus, in  
4 order to internalize the externalities of using products, which potentially release micropollutants, the  
5 purchase cost should be increased in order to subsidize the removal/remediation expenses. Revenues  
6 should be allocated to upgrade WWTPs, with the unfailing support of national (and, possibly,  
7 international) policies which consider the global social and environmental costs due to the use of such  
8 substances, together with the costs for water treatment (from drinking water supply, to wastewater  
9 collection and purification).

10

## 11 **6 Interpretation of eco-toxicity data: case studies**

12 In recent years, some authors have applied toxicity tests to diverse applications. In this section, some  
13 case studies are presented, which demonstrate the power and versatility of such investigations. For this  
14 purpose, the examples chosen include a range of different scenarios, in terms of: employed bioassays  
15 (cladocer crustaceans, algae, bacteria, etc.); tested matrices (e.g., municipal and complex wastewater);  
16 adopted treatment systems (conventional activated sludge process, membrane bioreactor, ozonation,  
17 photocatalysis, sonication, anaerobic process). Some of the experiences have been carried out at the full  
18 scale.

19 The pivotal role of bioassays in the integrated assessment of the environmental impact of wastewater is  
20 clearly manifest in all the reported cases.

21

### 22 **6.1 Ecotoxicity removal from complex wastewaters: comparison among conventional and** 23 **advanced technologies**

1 Currently, water quality standards and wastewater discharge limits in the European Union are mostly  
2 based on a limited number of chemical parameters. The aim of The European Water Framework Directive  
3 [189] is to obtain water bodies with a “good” biological quality. The biological or ecological impact of  
4 complex industrial effluent discharges however, cannot be estimated using chemical assays only, but  
5 should be measured using whole effluent toxicity (WET) tests (e.g. [156]).

6 A typical example of a complex industrial effluent is the water originating from tank truck cleaning  
7 (TTC) activities. The TTC process mainly involves the cleaning of tank truck interiors. The wide  
8 spectrum of transported cargo, ranging from food products to hazardous chemicals, results in wastewater  
9 with a highly variable composition. De Schepper *et al.* [190] reported that a significant residual toxicity  
10 was still present in biologically treated TTC effluent. A battery of acute ecotoxicity assays, with  
11 *Raphidocelis subcapitata* (primary production), *Vibrio fischeri* (decomposition) and *D. magna* (primary  
12 consumption) was applied to assess the whole effluent toxicity. It was found that the effluent of the full-  
13 scale treatment plant was extremely toxic to *R. subcapitata* with toxicity values ranging from 800 to 3260  
14 TU (toxic units).

15 The aim of a subsequent study was to investigate the removal of acute toxicity from TTC wastewater by  
16 a series of key unit operations applied during the treatment of industrial wastewater, i.e. chemical  
17 coagulation, activated sludge treatment and sorption by activated carbon [191]. The treatments steps were  
18 performed on a laboratory scale, to assess the full toxicity removal potential of these technologies. The  
19 rapid *V. fischeri* bioluminescence inhibition test (applying a 30 min contact time) was used to assess  
20 toxicity removal. Chemical pretreatment of the wastewater by coagulation with FeCl<sub>3</sub> removed approx.  
21 38% of the influent chemical oxygen demand (COD) and reduced the bioluminescence inhibition by 8%.  
22 Biological treatment with activated sludge subsequently removed another 77% of the remaining COD.  
23 This treatment step also reduced the bioluminescence inhibition, but the removal efficiency varied  
24 strongly from 5 to 92% for the different samples.

1 The ecotoxicity of the biotreated samples was also analyzed with the 72 h algal growth inhibition assay  
2 using *R. subcapitata*. The TU values ranged from 610 to 5,470, confirming the very high algal growth  
3 inhibition reported for the same type of wastewater by De Schepper *et al.* [190].

4 Powdered activated carbon (PAC) almost completely removed the remaining COD and inhibition in all  
5 samples. The algal growth inhibition after PAC addition ranged from 23 to 82 TU, corresponding to a  
6 reduction of more than 95%.

7 These results suggest that conventional technologies did not suffice for complete removal of toxicity  
8 from TTC wastewater, and that advanced wastewater treatment technologies are required for a  
9 satisfactory detoxification.

10

## 11 **6.2 Removal of estrogenicity from municipal wastewater: comparison between MBR and CAS** 12 **systems**

13 A monitoring campaign was conducted on a full scale municipal WWTP, consisting of 2 CAS and 1  
14 MBR (ultrafiltration) parallel lines. The design size is 250,000 p.e. and the influent load is split about  
15 50% on the MBR train and 25% on each CAS line. The plant is operated according to the modified  
16 Ludzak-Ettinger process scheme, with chemical phosphorus removal (aluminium sulphate dosage into  
17 the biological reactors).

18 Both chemical and biological analyses were carried out all along a 19 days period, in order to compare  
19 the CAS and MBR processes in terms of EDCs removal. The following target substances were measured:  
20 4-nonylphenol (NP), its parent compounds 4-nonylphenol monoethoxylate (NP1EO) and 4-nonylphenol  
21 diethoxylate (NP2EO), and bisphenol A (BPA). The same wastewater samples used for chemical  
22 analyses were submitted to the measurement of hormonal activity by means of human breast cancer  
23 MCF-7 based reporter gene assay, using  $17\beta$ -estradiol (E2) as a standard.

1 Removal efficiency and residual effluent concentration of target compounds were quite similar for both  
2 CAS and MBR lines, ranging between 70% (BPA) and 95% (NP1EO) and from 0,3 mg/L (NP1EO) to  
3 0,8 mg/L (NP), respectively. The CAS and MBR lines were operated at a sludge age of 9 and 15 d,  
4 respectively, the sewage temperature being around 23°C. The reason for the different plants to have  
5 similar performances can be explained based on the well-known relevance of these operating parameters:  
6 Clara et al. [192], [193] demonstrated that any increase of sludge age and temperature above 10 d and  
7 10°C does not lead to noticeable improvements, regardless of the type of process (either CAS or MBR).  
8 Moreover, several Authors (e.g. [194], [195], [196], [197]) evidenced the positive effect of an efficient  
9 nitrification on EDCs removal.

10 Nevertheless, even if no appreciable difference in the EDCs effluent concentration was detected,  
11 biological measurements showed that the MBR effluent exerted a lower estrogenic activity  
12 (estrogenicity, expressed as Relative Light Units, and normalized towards protein concentration, was up  
13 to 50% lower in MBR effluent samples, ranging from 1.0 to  $3.5 \times 10^7$  RLU/mg<sub>protein</sub>). The higher  
14 performance of the MBR system is likely attributable to the more efficient retention of suspended solid,  
15 and, consequently, of specialized slow-growing bacteria and of the organics to be degraded (in case they  
16 are adsorbed onto the suspended solids).

17 The findings confirm the irreplaceability of bio-assays in the monitoring of any impact on the ecosystems  
18 (in this case, the biological reactor of a WWTP). Detailed results are reported in [92].

19

### 20 **6.3 Removal of antibiotics and their effects of anaerobic and aerobic systems**

21 As the working principle of antibiotics inhibits biological activities directly, their adverse/inhibitory  
22 effects on the biodegradation of organic compounds in the wastewater treatment plants are one of the  
23 main concerns. In order to evaluate the inhibitory impact of these compounds in biological systems, two

1 different experimental approaches are commonly applied: short-term (acute) and long-term (chronic)  
2 tests. The short-term, acute tests usually involve a non-acclimated microbial population to the inhibitor.  
3 In long-term experiments with continuous feeding of the antibiotics, the test may reflect, aside from  
4 changes in substrate removal and utilization, adaptation and/or resistance of the microbial community  
5 or even shifts in microbial composition in response to continuous exposure [198], [199]. While  
6 Kümmerer and his colleagues [200] argue that short-term assays would not be sufficient to investigate  
7 the effect of antibiotics on complex microbial systems because of different mechanisms associated with  
8 acute and chronic inhibition, Alighardashi *et al.* [201] propose that the microbial community becomes  
9 well adapted to a synthetic substrate, which is a significantly different scenario from biomass in a full-  
10 scale plant under long-term exposure. Despite different opinions expressed in the literature, these two  
11 inhibition tests complement one another and reflect real-life inhibition schemes encountered in  
12 wastewater treatment.

13 In the light of this knowledge, acute and chronic tests were applied to aerobic and anaerobic biological  
14 treatment systems with three selected antibiotics: sulfamethoxazole (SMX), tetracycline (TET) and  
15 erythromycin (ERY).

16 For the aerobic acute tests; laboratory-scale fill-and-draw reactors with hydraulic retention time of one  
17 day were established and sustained at sludge ages of 10 and 2 days at steady state under aerobic  
18 conditions [202], and a series of fully aerated batch reactors for kinetic investigations of peptone-meat  
19 extract mixture biodegradation and acute/chronic inhibition of the selected antibiotics ([203], [204],  
20 [205]). Fill-and-draw reactors were fed with peptone-meat extract mixture at concentrations  
21 characterizing domestic wastewaters. To determine the acute and chronic inhibition effects of the  
22 selected antibiotics, batch experiments were conducted with 50 mg/L antibiotic additions [202].  
23 Respirometric tests were performed to determine the effect of antibiotics on unacclimated (acute effect)  
24 and acclimated (chronic) biomass, which yielded oxygen uptake rate (OUR) profiles. Obtained OUR

1 profiles were used for simulation to determine the kinetic properties of each activated sludge biomass  
2 ([204], [205]). Reactors were monitored for COD, suspended solids (SS), volatile suspended solids  
3 (VSS) and polyhydroxyalkanoates (PHA) [206]. The inhibitory impact of selected antibiotics was  
4 observed as a decrease in the amount of oxygen consumed in the OUR tests, which led to the conclusion  
5 that antibiotics have the property to block the microbial substrate consumption [202], [203]. The kinetic  
6 evaluation revealed that antibiotic substances mainly increase endogenous decay levels, the half-  
7 saturation constant of the substrate and inhibit hydrolysis of different COD fractions [202], [203], [204],  
8 [205].

9 For the aerobic acute tests; laboratory-scale fill-and-draw reactors with hydraulic retention time of one  
10 day were established and sustained at sludge ages of 10 and 2 days at steady state under aerobic  
11 conditions and a series of fully aerated batch reactors for kinetic investigations of peptone-meat extract  
12 mixture biodegradation and acute/chronic inhibition of the selected antibiotics [202]. Fill-and-draw  
13 reactors were fed with peptone-meat extract mixture at concentrations characterizing domestic  
14 wastewaters. To determine the acute and chronic inhibition effects of the selected antibiotics, batch  
15 experiments were conducted with 50 mg/L antibiotic additions. Respirometric tests were performed to  
16 determine the effect of antibiotics on unacclimated (acute effect) and acclimated (chronic) biomass,  
17 which yielded oxygen uptake rate (OUR) profiles. Obtained OUR profiles were used for simulation to  
18 determine the kinetic properties of each activated sludge biomass. The inhibitory impact of selected  
19 antibiotics was observed as a decrease in the amount of oxygen consumed in the OUR tests, which led  
20 to the conclusion that antibiotics have the property to block the microbial substrate consumption [203].  
21 The kinetic evaluation revealed that antibiotic substances mainly increase endogenous decay levels, the  
22 half-saturation constant of the substrate and inhibit hydrolysis of different COD fractions [202].

23 For the determination of short-term inhibition effects of the selected antibiotics under anaerobic  
24 conditions, a series of batch reactors seeded with acclimated microbial culture were run and fed with

1 volatile fatty acids (VFAs) in terms of acetate, butyrate, and propionate. Each reactor was also inoculated  
2 with a different concentration (1–1000 mg/L) of the selected antibiotics [207], [208]. The batch reactors  
3 were kept running for six days. Soluble COD and VFAs concentrations were monitored both at the  
4 beginning and at the end of the observation period. Total COD with soluble and particulate fractions  
5 were measured at the completion of the test in selected reactors for mass balance. Biogas production and  
6 methane generation were measured daily through- out the experiment. Organic substrate removal was  
7 monitored by both soluble COD and acetate measurements, together with daily measurements of biogas  
8 and methane generation. Sole acetate fed test showed that acetate was almost fully removed in all  
9 experiments, while methane generation exhibited a significant drop with increasing antibiotics doses  
10 [207]. Almost complete methane inhibition was observed for antibiotics doses above 500 mg/L. The  
11 monitored effect was found coherent with uncompetitive inhibition, which similarly exerts a binding  
12 impact on substrate–enzyme complex. For VFA mixture (acetate, propionate, and butyrate fed system),  
13 at lower doses, the VFA mixture was completely removed but partially used, leading to reduced biogas  
14 and methane generation, suggesting the resemblance of uncompetitive inhibition [208], [209].

15 Anaerobic chronic inhibition tests represented different results from acute tests. The experiments  
16 involved anaerobic sequencing batch reactors fed with a synthetic substrate mixture including glucose,  
17 starch, and volatile fatty acids, and operated in a sequence of different phases with gradually increasing  
18 antibiotics, for more than five months. TET exerted a terminal/lethal effect at 8.5 mg/L on the microbial  
19 community, which caused the inhibition of substrate/COD utilization and biogas production and leading  
20 to a total collapse of the reactor [210]. The microbial activity could not be retrieved and re-started within  
21 a period of more than 10 days, even after stopping TET dosing. During the experiments, TET was  
22 partially removed either through biodegradation or conversion into its by-products. The adverse long-  
23 term effect was quite variable for fermenting heterotrophic and methanogenic fractions of the microbial  
24 community based on changes generating on the composition of remaining/residual organic substrate.



1 The results revealed that anaerobic treatment was suitable for pharmaceutical industry wastewater with  
2 concentrations of up to 40 mg/L of SMX. Higher levels exerted toxic effects on the microbial  
3 community under anaerobic conditions, inducing the inhibition of substrate/COD utilization and biogas  
4 production and leading to a total collapse of the reactor. The adverse long-term impact was quite variable  
5 for fermentative bacteria and methanogenic archaeal fractions of the microbial community depend on  
6 changes inflicted on the composition of the residual organic substrate and mRNA expression of the key  
7 enzymes [211]. ERY fed reactors showed that methane production and VFA recovery are simultaneously  
8 possible up to 2 mg/L of ERY. ERY exerted a terminal effect at 3 mg/L on the biomass, and the activity  
9 could not be recovered after stopping ERY dosing [209].

10 Also, another study was performed to reveal if anaerobic-aerobic biological treatment strategy is proper  
11 for antibiotic production waste streams. Although activated sludge treatment systems are inhibited by the  
12 low concentration of antibiotic mixture, the same aerobic system can tolerate higher concentrations of  
13 the same mixtures after an anaerobic pre-treatment [212].

14

#### 15 **6.4 Removal of estrogenicity from textile wastewater by means of ozonation**

16 A pilot scale ozonation plant was installed at the outlet flow of a CAS plant (design size 370,000 p.e.,  
17 located in Northern Italy) treating mainly domestic wastewater. The CAS process scheme includes  
18 primary settling, pre-denitrification and oxidation-nitrification, secondary settling. Main CAS effluent  
19 characteristics are: 30 mgCOD/L, 5 mgBOD<sub>5</sub>/L, 12 mgTSS/L, 6.5 mgTKN/L; 4 mgNH<sub>4</sub><sup>+</sup>-N/L, 4 mgNO<sub>3</sub><sup>-</sup>  
20 -N/L, <0.1 mgNO<sub>2</sub><sup>-</sup>-N/L, 1.3 mgP<sub>TOT</sub>/L.

21 The O<sub>3</sub> pilot plant consisted of a stainless-steel tubular reactor (volume = 1,460 L) equipped with a pure  
22 oxygen supply system (capacity = 400 gO/h). The reactor was fed with a flow-rate up to 6 m<sup>3</sup>/h in a  
23 continuous mode of operation. Two different dosages were tested, namely 8 and 11 mgO<sub>3</sub>/L, with an  
24 HRT of 20 min.

1 The estrogenicity of wastewater was reduced from 7.35 down to  $3.25 \times 10^7$  RLU (Relative Light  
2 Units)/mg<sub>protein</sub> (about 55% removal efficiency) by means of ozonation, under the lower dosage  
3 conditions. Nevertheless, while the higher O<sub>3</sub> dosage led to an appreciable improvement of EDCs  
4 removal (data not shown: see full data in [92]), only a slight additional reduction of hormonal activity  
5 was achieved (measured value =  $2.90 \times 10^7$  RLU/mg<sub>protein</sub>; removal efficiency = 60%). The difference  
6 between the chemical and the biological answer may be due to the formation of active by-products,  
7 metabolites and/or conjugates, able to exert an estrogenic activity comparable to those of parent  
8 compounds, and to the synergistic effect among the different compounds.

9 In summary, the information gathered from chemical analyses was somehow misleading: the power of  
10 ozonation was overestimated; on the contrary, the bioassay gave a more realistic evaluation of the results  
11 obtainable.

12

### 13 **6.5 Removal of emerging pollutants from municipal wastewater by means of photocatalysis and** 14 **ultrasound treatments**

15 Photocatalysis and ultrasound treatments have been widely investigated for the treatment of emerging  
16 pollutants in urban wastewaters, including EDCs, pharmaceuticals, personal care products, drugs [213],  
17 [214], [215], [216]. Since during the oxidation process some by-products (intermediates) are formed and  
18 the effluent may become more toxic than the untreated solutions or the parent compounds, respectively,  
19 the overall efficiency of the treatment process for this class of chemical pollutants strictly depends on the  
20 toxicity and estrogenic potency of treated effluents.

21 The toxicity of photocatalytic degradation of caffeine, the number one drug worldwide, has been  
22 investigated in aqueous suspensions of titanium dioxide (TiO<sub>2</sub>) (29.3 – 170.7 mg/L) and initial drug  
23 concentrations (0.76 - 9.24 mg/L) by Carotenuto *et al.* [215]. Caffeine was quickly degraded, but not  
24 mineralized as quickly, and it was found that persistent toxic organic intermediates resist further

1 oxidation producing toxicity on *D. magna* at 24h and 48 h. *R. subcapitata* showed to be more sensitive  
2 to by-products than *L. sativa*.

3 A set of bioassays (*D. magna*, *R. subcapitata* and *Ceriodaphnia dubia*) was performed to evaluate the  
4 potential detoxification of the antibiotic vancomycin B hydrochloride (VAN-B, 50 mg/L) and its  
5 oxidation by-products under acute and chronic conditions. The toxicity of the photo-catalytically treated  
6 VAN-B samples varied during the oxidation, due to the formation of some intermediate by-products that  
7 are more toxic than VAN-B. Despite almost total removal of VAN-B that was achieved within 120 min  
8 of irradiation with 0.2 gTiO<sub>2</sub>/L, a significant increase in toxicity was observed in chronic tests proving  
9 that the chronic assays are more sensitive than acute ones to detect the impact of by-products formed  
10 during the photocatalytic degradation of antibiotics [217]. The residual toxicity of photo-catalytically  
11 treated solutions of chloramphenicol sodium succinate (CAP, 25 mg L<sup>-1</sup>), which is a broad-spectrum  
12 antibiotic, evidenced a decreasing trend in toxicity at increasing concentrations of TiO<sub>2</sub> and photo-  
13 oxidation times. After 120 min of photo-oxidation the most significant effect on *V. fischeri* (p<0.05) was  
14 obtained at 1.6 g/L of TiO<sub>2</sub> with a residual toxicity of 8 ± 6% (5min) and 10 ± 4% (15min). Lower TiO<sub>2</sub>  
15 concentrations showed toxicities ranging between 45–62% (5min) and 53–76% (15min) [216].

16 The toxicity of the mixtures of three pharmaceuticals (2.5 mg/L, diclofenac, DCF, 2.5, 5 and 10 mg L<sup>-1</sup>,  
17 amoxicillin, AMX, 2.5, 5 mg/L carbamazepine, CBZ) at different concentrations in contaminated urban  
18 wastewater treated by ultrasound has been evaluated by Naddeo *et al.* [218]. Sonication decreased  
19 toxicity of contaminated WW sample to *R. subcapitata* and no significant effect on this decrease by either  
20 the sonication time or the applied power density was observed. *R. subcapitata* was found more sensitive  
21 than *D. magna*.

22 Toxicity data about photocatalysis and ultrasound treatments are still in their infancy, especially for  
23 sonolysis where just few studies have been performed. From the available results it can be stated that  
24 photocatalysis can be suitable to fully remove toxicity at the discharge but focused research must be

1 oriented specifically, not only on target compound removal but also on effluent toxicity goal. Moreover,  
2 toxicity investigation must comply with the international recognized approach, considering the  
3 integration of at least three species belonging to different phylogenetic levels [150], [151].  
4

## 5 **7. Conclusions**

6

7 This paper reports the shared opinions of the participants to COST Action ES1202 Conceiving  
8 Wastewater Treatment in 2020-Energetic, environmental and economic challenges (Water\_2020) about  
9 the topic of toxicity of wastewater organic trace pollutants.

10 Notwithstanding the valuable literature production, which, up to now, includes also hundreds of reviews,  
11 the choice to write another work about the topic of toxicity of wastewater organic trace pollutants arose  
12 from the awareness that there are still gaps between the different scientific sectors involved in this  
13 research.

14 The debated subjects, indeed, pertain to several disciplines and have been connected based on the final  
15 goal to propose criteria for choosing the proper tools to assess and reduce the possible environmental  
16 impact of such pollutants on the human health and the aquatic ecosystems.

17 Some aspects appear particularly worth noting:

- 18 - bioassays must be chosen considering the meaning of the biological response;
- 19 - laboratory and in situ bioassays must be integrated, by considering their specific reliability and  
20 applicability;
- 21 - trace pollutants can cause unpredictable and non-linear responses by biological systems;
- 22 - wastewater variability in terms of pollutants concentrations and flowrates deeply affects any  
23 toxicity assessment;

1 - the choice of the optimal WWTPs process schemes for the abatement of trace pollutants must be  
2 based on the global assessment of environmental and socio-economic impacts.

3

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10

#### 11 **Disclaimer**

12 The content of this article is the authors' responsibility and neither COST nor any person acting on its  
13 behalf is responsible for the use, which might be made of the information contained in it.

14

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