The Use of Algae and Fungi for Removal of Pharmaceuticals by Bioremediation and Biosorption Processes: A Review

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Abstract: The occurrence and fate of pharmaceuticals in the aquatic environment is recognized as one of the emerging issues in environmental chemistry. Conventional wastewater treatment plants (WWTPs) are not designed to remove pharmaceuticals (and their metabolites) from domestic wastewaters. The treatability of pharmaceutical compounds in WWTPs varies considerably depending on the type of compound since their biodegradability can differ significantly. As a consequence, they may reach the aquatic environment, directly or by leaching of the sludge produced by these facilities. Currently, the technologies under research for the removal of pharmaceuticals, namely membrane technologies and advanced oxidation processes, have high operation costs related to energy and chemical consumption. When chemical reactions are involved, other aspects to consider include the formation of harmful reaction by-products and the management of the toxic sludge produced. Research is needed in order to develop economic and sustainable treatment processes, such as bioremediation and biosorption. The use of low-cost materials, such as biological matrices (e.g., algae and fungi), has advantages such as low capital investment, easy operation, low operation costs, and the non-formation of degradation by-products. An extensive review of existing research on this subject is presented.

Keywords: algae; fungi; bioremediation; biosorption; removal of pharmaceuticals; wastewater

1. Introduction

The growth of world population, the promotion of health and better living conditions, and the rise of average life expectancy were accompanied by an increase in the consumption of pharmaceuticals, which are excreted in their original form or as metabolites and collected by the sewage system of urban wastewaters; they are not completely removed by wastewater treatment plants (WWTPs). Their occurrence in the aquatic environment is creating polluting pressure on aquatic ecosystems and is recognized as one of the emerging problems in the last decade [1]. The development of analytical techniques (such as gas chromatography mass spectrometry and liquid chromatography mass spectrometry) enabled the detection and quantification of a broad range of pharmaceuticals (and their metabolites) in environmental matrices [2]. Although they occur at extremely low levels (ranging from \(\mu\text{g}\cdot\text{L}^{-1}\) to \(\text{ng}\cdot\text{L}^{-1}\) or even lower), it is known that their presence in the environment is a potential hazard to public health [3–6].

The release of pharmaceuticals into the aquatic environment is not yet subject to regulation. In Europe, the “Water Framework Directive 2000/60/EC” introduced a strategy for water protection [7]. However, since the water situation in the several European Union (EU) countries was, and still is, different, it was necessary to proceed with adjustments to the original “Water Framework Directive”.

In fact, one year after its publication, Decision 2455/2001/EC of the European Parliament and of the Council of 20 November 2001 established that each European Union member state should create, until 2009, a program of environmental protection measures for each of its hydrographic regions. A list of 33 priority substances in the field of water policy was also established, which includes pharmaceuticals. The revision and update of this list is done at least every four years, this may lead to the inclusion of new priority compounds, or removal, depending on the level of risk they pose [8]. Later, the “Directive on Environmental Quality Standards” (Directive 2008/105/EC) amended some environmental quality standards in the field of water policy and also established 11 new substances to be identified as priority substances or priority hazardous substances [9]. The Directive 2013/39/EU and Implementing Decision 2015/495 incorporated new substances, totaling 45, implementing the first so-called watchlist. A few pharmaceuticals (diclofenac, erythromycin, clarithromycin, and azithromycin), as well as synthetic and natural hormones (estrone, 17β-estradiol, and 17α-ethinylestradiol), were included, and they should be carefully monitored by the Member States in order to set their environmental quality standards [10,11].

In its last review, five substances or groups of substances were removed, among them diclofenac, and three new substances were included, among them amoxicillin and ciprofloxacin. This is consistent with the European One Health Action Plan against Antimicrobial Resistance, which supports the use of the watchlist to improve knowledge and to evaluate the risks to human and animal health posed by the presence of antimicrobials in the environment. This review resulted in the publication of the second watchlist in 2018 (of the Commission Implementing Decision 2018/840 [12]. Also, in the United States, the Environmental Protection Agency (EPA) included 12 pharmaceuticals, personal care products, and endocrine disrupting compounds in a list in order to evaluate their related occurrence and safety risks. Therefore, is expected that, in the near future, legal limits will be established for the concentration of pharmaceuticals in WWTP discharges.

Indeed, pharmaceuticals are introduced into the environment (Figure 1) mainly through the discharge of treated effluents from conventional WWTPs, via domestic and hospital wastewaters, directly or by leaching of the sludge produced by these facilities, or through discharges of the pharmaceutical industry [13–15], as well as veterinary facilities where pharmaceuticals are widely used in livestock production for disease prevention and growth promotion [16–18].

The most frequently detected classes of pharmaceuticals in wastewaters are antibiotics, antiepileptics, antiphlogistics, X-ray contrast media, lipid-regulators, β-blockers, and tranquillizers [19]. Most of these compounds have high solubility, low hydrophobicity, and often negative charge at neutral pH (acidic compounds); these properties add more difficulties to their treatment [20]. WWTPs were not specifically designed to completely remove pharmaceuticals [21,22]. Removal efficiencies can vary from negligible to 100% depending on the compound [23]. The majority of compounds are removed in a percentage of less than 50%, which is related to the structure of the compounds [20]. Many physico-chemical and biological treatments were tested by WWTPs, but none of them are able to efficiently remove them [24]. Tertiary treatments, such as membrane technologies and advanced oxidation processes, usually play a more active role in the removal of these micropollutants than primary and secondary treatments [5,25]. However, they have many drawbacks in terms of energy requirement, large use of chemicals, and formation of undesired and harmful by-products [26]. To overcome these drawbacks, the scientific community focused its attention on the development of eco-friendly, economically viable and comparatively less expensive technologies [27]. Biological tertiary treatments of wastewaters are efficient, less expensive, and more eco-friendly than other technologies. The use of certain microorganisms gained importance in applied environmental microbiology. Fungi and algae-based treatments were pointed out as promising technologies for the remediation of pharmaceuticals. In this article, special attention is given to studies addressing pharmaceutical removal with fungi and algae via bioremediation and/or biosorption mechanisms (Figure 2).
2. Bioremediation

Bioremediation is a biological treatment that involves the use of microorganisms and their enzymes to convert recalcitrant and xenobiotic contaminants to less toxic forms and, therefore, short lifetimes in the environment or even their complete mineralization (the end-products are essentially carbon dioxide and water) [28]. Bioremediation of organic compounds was studied in more detail at the laboratory level, with the metabolic pathways of degradation known for some pharmaceutical compounds [29,30]. The process is influenced by environmental, physical, and chemical factors, namely, the stereochemistry, toxicity, and concentration of the contaminant, efficiency of the microbial strain, conditions during degradation (e.g., pH and temperature), retention time, presence of other compounds, and their concentration [31]. Some of bioremediation’s advantages are as follows: it is accepted as a safe process, transforms pollutants instead of simply moving them from one medium to another [32], and presents lower costs when compared to the other technologies [33]. Although bioremediation proves to be a promising choice, research is needed to overcome some drawbacks of the process, which are the incomplete transformation, the limitation to biodegradable compounds, and the requirement of the selection and use of different microorganisms with specific metabolism for the different pollutants. Some of these disadvantages can be overcome through the use of genetically modified microorganisms [34].
2.1. Mycoremediation

Fungi (“mikes” from Greek) are eukaryotic organisms that include microorganisms such as molds, yeasts, and mushrooms. Some fungi are chemoheterotrophic organisms, being parasitic or saprophytic. Some are unicellular, and many are filamentous and have cell walls. The kingdom Fungi includes phyla Chytridiomycota (the chytrids), Zygomycota (the conjugated fungi), Ascomycota (the sac fungi), Basidiomycota (the club fungi), Deuteromycota (the imperfect fungi), and Glomeromycota. This classification was established according to their mode of sexual reproduction or using molecular data [35].

Fungi were efficiently used to treat water samples contaminated with micropollutants [36,37], pharmaceuticals in particular (Table 1). They are long recognized for their abilities to transform a broad range of recalcitrant compounds using nonspecific intracellular and extracellular oxidative enzymes [38–40]. The physiology and colonization strategy of mycelial fungi allows them to more easily withstand sudden changes in pH or humidity, as well as to degrade more efficiently complex organic compounds [41], although they are limited by a long growth cycle and spore formation [42].

2.1.1. Treatment Systems

Fungal reactors, also termed mycoreactors, can be suspended growth (such as slurry reactors) or immobilized systems (such as trickling filters, rotating biological contactors, upflow fixed-film reactors, and fluidized-bed reactors); these latter allow a fast biodegradation. Mycoreactors can be operated in batch, semi-batch, sequencing batch, or continuous mode, like other biological reactors. They can also be operated under aerobic or anaerobic conditions. Mycoreactors for submerged growth include stirred tanks, packed bed, bubble column, and air-lift [43], with stirred tanks as the most common, where the culture medium is agitated mechanically, providing a good fluid mixture and a good oxygenation. However, they have disadvantages such as the stress generated by the agitation, their impracticality for certain microorganisms, and the high energy consumption for high agitation speeds. Air-lifts are pneumatically agitated mycoreactors very similar to the stirred tanks; however, the fluid mixing is done by injecting air or compressed gas into the base of the bioreactor. This bioreactor is usually cylindrical so that the air bubbles remain as long as possible in the fluid. Compared with stirred tank, air-lifts have worse mixing rates; however, they generate less stress to the microorganisms [44,45].

Pellet-forming fungi, which are easier to recover at the end of the treatment, are cultivated in aerated fluidized-bed or suspended air-lift loop reactors [43].

2.1.2. Mechanisms of Removal

White-rot fungi (WRF), in particular, belong to the Basidiomycota phylum, whose potential was explored in several studies about the removal of pharmaceuticals (Table 1) [46–72]. These strains are filamentous wood-degrading fungi, ubiquitous in nature, able to mineralize lignin efficiently. The name white rot derives from the bleached appearance of the wood attacked by these fungi due to the removal of the dark-colored lignin. The same mechanism that gives these fungi the potential to degrade lignin also allows them to degrade a wide variety of recalcitrant pollutants, such as pharmaceuticals, making them promising and attractive microorganisms for wastewater bioremediation. They are able to mineralize a wide variety of pollutants since their enzymatic system is non-specific, non-stereoselective, and based on free-radical levels [73].

Pollutant degradation (Figure 3) seems to involve either an intracellular enzymatic system (i.e., cytochrome P450 system) or an extracellular enzymatic system (mainly lignin peroxidase, manganese peroxidase, laccase, and versatile peroxidase) [73,74]. Peroxidases are secreted during secondary metabolism of WRF in the presence of nitrogen, carbon, or sulfur limitations, while laccases are glycosylated multicopper oxidoreductases that are produced during primary metabolism [75]. Peroxidases generally show higher redox potential than laccases; however, peroxidases suffer deactivation in the presence of hydrogen peroxide while laccases do not [76–78]; laccase performance seems to be affected by the presence of chlorine ions forms [79]. Lignin peroxidase catalyzes the one-electron oxidation of various aromatic
compounds, with subsequent formation of aryl cation radicals which are decomposed spontaneously by various pathways. Manganese peroxidase catalyzes the oxidation of Mn$^{2+}$ to Mn$^{3+}$, which in turn can cause oxidation of several phenolic substrates [80]. Fungal laccases were reported in several studies since these enzymes have wide substrate ranges and use only oxygen as the final electron receptor, producing water as the only by-product [65,81–103]. In many cases, preference was given to the use of isolated enzymes (enzymatic bioremediation) instead of the use of fungi biomass in order to reduce the time of treatment, to avoid the lag phase of fungal growth, to reduce sludge production, and to facilitate process control [39]. It was found that _Phanerochaete chrysosporium_, a major WRF, does not have laccase genes [104]. _Trametes versicolor_ is a WRF frequently cited in the literature that was shown to be effective in the removal of different pharmaceuticals (Table 1) [46,47,50–55,57–62,72].

For example, strains of _Trametes versicolor, Irpex lacteus, Ganoderma lucidum_, and _Phanerochaete chrysosporium_ were simultaneously tested in the removal of ibuprofen, clofibric acid, and carbamazepine. The results suggested that clofibric acid and carbamazepine degradation occurred intracellularly by the cytochrome P450 system of _Trametes versicolor_ [46]. Using the same fungus, Nguyen et al. [54] compared the removal of trace organic contaminants (including pharmaceuticals and steroid hormones) by alive, intracellular enzyme-inhibited and chemically inactivated whole-cell preparations, and a fungal extracellular enzyme extract, predominantly laccase. The low degradation of some hydrophobic compounds by the extracellular extract, and the impact of intracellular cytochrome P450 system inhibition on the degradation of some trace organic contaminants by the whole-cell culture indicated the importance of extracellular enzyme-independent catalytic pathways. Rodriguez-Rodriguez et al. [49] studied the removal of sulfonamides sulfapyridine and sulfathiazole by _Trametes versicolor_. Complete degradation was achieved for both compounds, although a longer period of time was needed to completely remove sulfathiazole when compared to sulfapyridine. In order to determine the effect of cytochrome P450 inhibitors, piperonyl butoxide or 1-aminobenzotriazole was added in the experiments performed. The results showed that sulfathiazole degradation was partially suppressed, while no additional effect was observed for sulfapyridine. In another study, _Trametes versicolor_ was able to degrade carbamazepine in aqueous medium in an air-pulsed fluidized bioreactor in batch and continuous mode. In batch mode, carbamazepine concentration decreased 96%, while, in continuous mode, carbamazepine concentration decreased 54%. In this case, it was not possible to establish a correlation between extracellular laccase activity and carbamazepine degradation, since laccase and manganese peroxidase levels were negligible during the initial period, which may indicate that these enzymes were involved in an early stage of carbamazepine removal [47].

Rodarte-Moralez et al. [48] studied the removal of six pharmaceuticals (citalopram, sulfamethoxazole, diclofenac, ibuprofen, naproxen, and carbamazepine) from an initial mixture by three other WRF strains, _Bjerkandera_ sp. R1, _Bjerkandera adusta_, and _Phanerochaete chrysosporium_, and also confirmed their enzymatic activity, particularly manganese peroxidase. An intense enzymatic activity was also detected during the individual degradation of 17α-ethinylestradiol and carbamazepine by ligninolytic fungi strains _Pleurotus_ sp. P1, _Pleurotus ostreatus_ BS, and a (unidentified) basidiomycete strain BNI. During 17α-ethinylestradiol degradation by _Pleurotus_ sp. P1, laccase and manganese peroxidase activity was detected, while, during carbamazepine degradation by strain BNI, laccase, manganese peroxidase, and lignin peroxidase activity was detected [68].

Nguyen et al. [51,54] revealed in their studies that the enzymatic performance of laccase, in particular, can be enhanced by the addition of mediators. These authors studied the effect of continuous dosing of a mediator (1-hydroxybenzotriazole) in the removal of trace organic contaminants, including pharmaceuticals, from an initial complex mixture by _Trametes versicolor_. Becker et al. [63] studied the removal of 38 antibiotics (majority non-phenolic) from a mixture using immobilized laccase (_Trametes versicolor_), in an enzymatic membrane reactor, with or without the addition of syringaldehyde as a mediator. Thirty-two out of 38 antibiotics were removed by up to 50% when a mediator was used. In contrast, no significant removal was observed in experiments applied without a mediator.
Indeed, mediators can be oxidized by laccase to free radicals, which in turn can oxidize pollutants less specifically, increasing the variety of pollutants potentially degraded by these enzymes [105].

Synthetic and natural laccase mediators were used in enzymatic studies. Synthetic mediators include 2,2’-azino-bis(3-ethylbenzothiazoline-6sulfonate) and 1-hydroxybenzotriazole, while natural phenolic mediators include syringaldehyde and acetosyringone. Natural mediators are more economically feasible and more environmentally friendly than artificial mediators [106]. Each mediator has a specific catalytic mechanism [107,108]. The effect of a mediator depends on the radicals formed, the mediator recyclability, and the laccase stability in the mediator’s presence [106,107,109,110]. Despite the proven increase in efficiency, mediators incur additional costs, and can cause toxicity [63,111] and laccase inactivation [112–114]. For example, Becker et al. [63] observed that, although the addition of syringaldehyde enhanced the removal of antibiotics, unspecific toxicity was also induced.

Figure 3. Fungi and algae strategies to counteract pharmaceuticals.

2.1.3. Factors That Influence the Degradation Capability

Fungi are able to degrade very low or non-detectable concentrations levels [73] and withstand a wide range of pH, further enhancing their degradation capability [115]. Nevertheless, mycoremediation requires specific and controlled conditions in order to maintain a durable and efficient process. Fungi oxidative metabolism can be strongly affected by the presence of nutrients, pH, immobilization on different supports, and agitation/static growth conditions [31]. Zhang and Geißen [116] tested the degradation of carbamazepine and diclofenac by lignin peroxidase produced by WRF Phanerochaete chrysosporium in various conditions. It was found that lignin peroxidase completely degraded diclofenac at pH 3.0–4.5 and 3–24 mg·L⁻¹ H₂O₂, while the degradation efficiency of carbamazepine was mostly below 10%. The addition of veratryl alcohol and the high temperature (30 °C) did not enhance the carbamazepine degradation. In another study, the removal of carbamazepine by Phanerochaete chrysosporium in a plate bioreactor operated in batch and continuous systems showed that carbamazepine removal depends on a sufficient nutrient supply. Carbamazepine concentration decreased about 80% when a diluted synthetic nutrient medium was fed to the reactor and decreased about 60% when a real effluent with additional glucose and nitrogen was fed to the reactor, after 100 days of incubation [49].
By opposition, 17α-ethinylestradiol was completely degraded by Pleurotus sp. P1, Pleurotus ostreatus BS, and a (unidentified) basidiomycete strain BNI, in the presence or absence of another carbon source [68]. Becker et al. [57] showed that the degradation of hormones by laccases is feasible even at very low enzyme concentrations, and that immobilized enzymes displayed better removal performance compared to the free enzyme. Indeed, enzyme immobilization provides a more suitable environment for enzymes and may result in an increasing enzyme stability concerning pH, temperature, and storage time. Efficient use of immobilized cells requires, however, control of physiological/metabolic changes occurring either during the immobilization process or during the biotransformation phase. Enzyme immobilization can be done with irreversible methods (e.g., covalent binding and entrapment) or reversible methods (e.g., adsorption, ionic binding, affinity binding, chelation, and disulfide bonds). The selection of the immobilization method is based on technical–economic criteria. Since cost is a preponderant parameter, the simplest ones are usually used [117].

An eventual contamination with bacteria can negatively affect the efficiency since it can generate competition for the substrate, damage fungi mycelium, disrupt biomass growth, and destabilize fungal activity [118,119]. To avoid bacterial contamination, some strategies can be implemented. In the biological treatment system, the pH has great influence, since it acts in the development and selection of the microorganisms. Therefore, it is convenient to operate under acidic conditions, since the optimum pH for fungi growth is lower than the optimum pH for most bacteria. Other possible strategies include pretreatment by coagulation–floculation of wastewater, since a reduction of the initial bacterial count usually occurs; coupling the bioreactor with a micro-screen, which allows the retention of fungal biomass and, simultaneously, the washout of bacteria with effluent; the use of disinfectant agents that allow the selection and/or inactivation of bacteria; the immobilization of fungal strains in different carriers; and the periodic replacement of fungal biomass [118]. Sterility conditions do not appear to be a mandatory requirement to apply Trametes versicolor in pharmaceutical removal. The studies performed showed that this microorganism was able to partially or completely remove pharmaceuticals either under sterile or non-sterile conditions [50,52].

2.1.4. Concluding Remarks and Future Challenges for Mycoremediation

In general, studies indicate a significant reduction in wastewater toxicity after treatment. Some fungi strains found in the literature [46,47,50,52,53,61,120] proved to be effective in pharmaceutical removal (Table 1). Final products were reported to be less toxic or more biodegradable than the parent compounds, which emphasizes the potential of fungi as remedial agents.

Genetic tools can represent an essential step to improve fungi performance and overcome some limitations associated with the process. The sequencing of fungi genomes will allow the development of new genetic techniques to improve metabolic and adaptive processes, and consequently achieve efficient bioremediation [121]. To date, efforts met limited success, and there is still a long way to go before the introduction of modified fungi in remediation processes. The first complete eukaryotic genome belongs to the yeast Saccharomyces cerevisiae [122]. Several projects released information about the genome sequences of fungi Aspergillus nidulans, Aspergillus fumigatus, Neurospora crassa, and Coprinus cinereus [123]. The thirty-million-base-pair genome of WRF Phanerochaete chrysosporium strain RP78 was sequenced using a whole-genome shotgun approach [124]. The genome revealed genes encoding oxidases, peroxidases, and hydrolytic enzymes involved in wood decay, which opens new horizons related to the process of biodegradation of organic pollutants and pharmaceuticals in particular [125]. However, the risks involved in the use of genetically modified fungi, their impact on the environment and human health, and existing legal limitations must be considered [126].

2.2. Phycoremediation

Algae (“phyco” from Greek) include prokaryotic organisms (cyanobacteria) and eukaryotic organisms (all the algae species) that contain chlorophyll and carry out oxygenic photosynthesis. Although most algae are of microscopic size and, hence, are clearly microorganisms (microalgae),
several forms are macroscopic (macroalgae) grow to over 30 m in length. Algae are either unicellular or colonial. When the cells are arranged end to end, the alga is said to be in filamentous form [127]. According to Ruggiero et al. [128], algae can be classified according to the pigments they possess. Algae color differences arise due to the proportions of different auxiliary photosynthetic pigments present in addition to the green chlorophylls. Chlorophyta (the green algae) and Euglenophyta (the euglenoids) are green since chlorophyll \( a \) is dominant. If carotenoids are dominant in the algae, they give them a golden-brown color, such as Chrysophyta, whose chloroplasts contain chlorophylls \( a \), \( c_1 \), and \( c_2 \), fucoxanthin, and \( \beta \)-carotene, which are the carotenoids responsible for the golden-brown color. Dinophyta (the dinoflagellates), have a reddish, greenish, or brown appearance due to chlorophylls \( a \) and \( c_2 \) and carotenoids. Rhodophyta (the red algae) have chlorophylls \( a \) and \( d \), phycobiliproteins, and floridean starch as storage products accumulated in the cytoplasm outside the chloroplast. In Phaeophyta (the brown algae), color results from the dominance of fucoxanthin over chlorophylls \( a \), \( c_1 \), and \( c_2 \).

Algae are highly adaptive microorganisms and can grow autotrophically, heterotrophically, or mixotrophically. They can grow in very harsh environmental conditions, such as low nutrient levels, and extreme pH and temperature, which is an advantage over some species of fungi [129]. Unlike strictly heterotrophic microorganisms, the decrease in nutrient concentration does not limit the growth of algae [130]. Microalgae can acclimatize to changes in temperature, salinity, light, and nutrient availability, which allows the improvement of their tolerance and biodegradation capacity. This adaptation mechanism to extreme conditions is explained by genetic changes caused by spontaneous mutation or physiological adaptation [131,132]. Xiong et al. [133] evaluated the biodegradation capability of *Chlorella vulgaris* after acclimation with multiple exposures to levofloxacin and an increase in salinity. Results showed that levofloxacin biodegradation was significantly improved after acclimation.

The characteristics of domestic wastewaters is usually suitable for the cultivation of microalgae, since wastewaters represent a source of nutrients. From an environmental perspective, photosynthetic microalgae are fascinating since they can sequester atmospheric carbon dioxide for their own growth, contributing to the mitigation of this pollutant. In a properly controlled process, the association of microalgae with bacteria may result in a very synergistic relationship; microalgae provide oxygen while bacteria release carbon dioxide, which allows a significant reduction of oxygen needs in the treatment process [134–137]. In addition, microalgae treatment supplies an environment that increases the mortality of pathogenic organisms due to the pH elevation [138]. A win–win situation of using microalgae in wastewater bioremediation offers a tertiary biotreatment of wastewater coupled with the production of potentially valuable biomass as a bioresource for biofuel or high-value by-products [139].

### 2.2.1. Treatment Systems

From the economic point of view, open systems are preferred for wastewater phycoremediation. Operational factors that influence algae growth are essentially mixing, dilution rate, and depth [140]. Open ponds are the most usual for microalgae cultivation since they require lower investment costs and operational capital. Commercially, algae cultivation is mainly performed on open channels stirred with a paddlewheel (raceway), since they are less expensive and easier to construct and operate. There is, however, a high risk of contamination and low productivity due to a poor mixing regime and light penetration, and also due to the difficulty of controlling the operating conditions [141]. Facultative, maturation, and high-rate algal ponds are the most used open systems for wastewater treatment, with the first two being the most used. The major differences between them are the depth and origin of the influent. The major constraints include poor light utilization by the cells, water loss due to evaporation, diffusion of carbon dioxide to the atmosphere, temperature fluctuations, inefficient stirring, and large space requirements for microorganism growth [142]. Open systems usually operate under long hydraulic retention time (between eight and 14 days) in order to consume carbon dioxide
during the day (photosynthesis) and provide oxygen for aerobic biodegradation. Sunlight intensity influences photosynthetic activity, leading to pH and dissolved oxygen variations [143–145].

Closed systems were designed to overcome the problems associated with open systems. Unlike open systems, closed systems allow greater control of the process; however, they are expensive to install and to maintain [146]. Closed systems are most suitable for pure algae strains, and their design must be carefully optimized for each individual strain according to its unique physiological and growth characteristics. These systems avoid losses by evaporation, and reduce the contamination risk and the losses of carbon dioxide to the atmosphere [147]. Closed systems, which are mostly photobioreactors, exist in various design configurations (e.g., horizontal or serpentine tube, flat-plate, bubble column, air-lift column, and stirred tank). Flat-plate photobioreactors have space constraints, and tubular photobioreactors have design limitations. Column photobioreactors are the most efficient, which provide efficient mixing, the highest volumetric mass transfer rates, and the best controllable growth conditions. Some difficulties arise with scale-up caused by inhomogeneous distribution of light inside the culture. The productivity is negatively affected by central, light-deprived zones [148]; therefore, to facilitate penetration of light, photobioreactors are made of glass, transparent plastic material, or sturdy polythene.

Among all the drawbacks associated with phycoremediation, harvesting is one of the most challenging processes integrated in the removal treatment. The harvesting process may account for 30% of total microalgae biomass production cost. Several factors affect the harvesting, such as microalgae strain and operational parameters (hydraulic and solid retention times) [149]. Harvesting technologies may involve one or more steps, and different physical (e.g., centrifugation, gravity sedimentation, filtration, and dissolved air flotation), chemical (e.g., chemoflocculation), and biological (e.g., bioflocculation and microalgae immobilization) processes.

Most of these processes are limited by high costs, long processing times, high energy consumption, and low recovery [150,151]. The choice of the harvesting method depends on the microalgae characteristics, such as the density and size of algal cells, as well as the product derived from the algal biomass [152].

The most environmentally friendly method is bioflocculation as it involves formation of extracellular biopolymers that help natural flocculation of small particles [150,151]. Some microalgae species flocculate more readily than others, and these microalgae can be mixed with other species to induce flocculation [153,154]. However, this method may be not efficient due to the small size of microalgae cells and their fast growth. Further research is required to understand the underlying mechanism associated with this process.

New technologies were considered, namely, the addition of other microorganisms, such as bacteria and some fungal species, to the microalgae culture [135]. In the first case, microalgae and bacteria may form flocs that settle more easily than single microalgae. It is necessary to add an extra organic substrate to allow bacteria growth [135], which can be provided by using wastewater. The presence of an organic carbon source in wastewater allows both organisms to thrive together [155,156]. The main disadvantages are the influence of dominant microalgae species and the variability of wastewaters [135]. In the second case, some filamentous fungi can pelletize, entrapping the microalgae cells, which facilitates the harvest by simple sieve filtration or sedimentation in most cases [157]. It is not necessary to add an extra sugar to allow fungi growth [158]. Co-pelletization efficiency seems to depend both on fungi and microalgae strains and on culture conditions (in particular on the pH value). The choice of fungi strains will be the key issue since it determines the overall pelletization efficiency and, therefore, can have a direct impact on the subsequent processes. Since this method does not require the addition of chemicals or inputs of energy, it may offer a solution to two of the major problems of harvesting processes, the high cost and high energy consumption [159]. The competition of nutrients between fungi and microalgae is believed to occur mostly in heterotrophic conditions, since microalgae need external carbon sources to support their growth, or in autotrophic conditions, when other nutrients, such as organic nitrogen, are limited. In this context, neither fungi nor microalgae cells will reach their maximum growth compared with their growth in individual pure cultures [160]. It is not clear if co-culturing fungi affects the growth
and nutrient assimilation of microalgae [161]. It is also unclear how the use of fungi for pelletization is fundamentally different or superior to the use of bacteria for bioflocculation [161,162].

2.2.2. Mechanisms of Removal

Studies about the removal of emerging contaminants (such as pharmaceuticals) with algae are limited; therefore, the mechanisms involved are not yet very clear (Table 1) [163]. The possible mechanisms involved in algal–bacterial systems (Figure 3) seems to be biodegradation, biomineralization (bioprecipitation), biosorption (cell adsorption and/or bioaccumulation), stripping (volatilization), and photodegradation, due to the effect of light [163–165]. However, volatilization can be considered negligible for most pharmaceuticals, because of their low Henry’s constant values [166]. The bioaccumulation of pharmaceuticals in algae cells can induce the generation of reactive oxygen species, free radicals (e.g., \( \text{O}_2^\cdot \))—superoxide radicals, \( \text{OH}^\cdot \)—hydroxyl radical, \( \text{HO}_2^\cdot \)—perhydroxy radical, and \( \text{RO}^\cdot \)—alkoxy radicals) and nonradical forms (e.g., \( \text{H}_2\text{O}_2 \)—hydrogen peroxide and \( \text{1O}_2 \)—singlet oxygen). At normal levels, these species act as essential signaling molecules to control cellular metabolism; however, at excess levels, these species can cause severe damage to cellular components and an increased rate of mutagenesis that ultimately leads to programmed cell death [167]. Such as for fungi, pollutant degradation by algae seems to involve intracellular and extracellular enzymatic systems (Figure 3). Intracellular degradation of pharmaceuticals involves a phase I enzyme (cytochrome P450) [163,168–173] and a phase II enzyme (e.g., glutathione-S-transferases) [171]. Extracellular degradation of pharmaceuticals involves the excretion of various extracellular polymeric substances, such as polysaccharides, protein, enzymes, substituents (polysaccharide-link methyl and acetyl groups), and lipids to their surrounding environment. These extracellular polymeric substances can form a hydrated biofilm matrix that acts as an external digestive system since they keep extracellular enzymes close to the cells [174].

Peng et al. [168] studied the removal of progesterone and norgestrel by the microalgae Scenedesmus obliquus and Chlorella pyrenoidosa. According to the authors, biotransformation was found to be the main mechanism for the removal of progestogens. Hydroxylation, oxidation/reduction, and side-chain breakdown were proposed to be involved in the algal transformation of the target compounds by Chlorella pyrenoidosa. Xiong et al. [169] studied the removal of carbamazepine by the microalgae Chlamydomonas mexicana and Scenedesmus obliquus. The results showed that both species simultaneously promoted biodegradation, adsorption, and bioaccumulation of carbamazepine. De Godos et al. [164] studied the mechanisms of tetracycline removal from a synthetic wastewater by Chlorella vulgaris and identified photodegradation and biosorption as the most important.

2.2.3. Factors That Influence the Degradation Capability

There are several factors that affect the degradation of pharmaceuticals, which are mainly related to the operation conditions, reactor configuration, and the species present. Matamoros et al. [175] studied the removal of emerging organic compounds, including pharmaceuticals, from a real wastewater by the marine algae Lessonia nigrescens Bory and Macrocystis integrifolia Bory in two pilot-scale high-rate algal ponds. Removal efficiency ranged from negligible to up to 90% and was only affected by the hydraulic retention time during the cold season. This effect was not observed in the warm season.

De Godos et al. [164] studied the removal of tetracycline by Chlorella vulgaris in two pilot-scale high-rate algal ponds operated in batch mode. The study demonstrated that the shallow geometry of high-rate algal ponds is advantageous to support tetracycline’s photodegradation.

Díaz-Garduño et al. [176] studied the removal of organic compounds, including pharmaceuticals, from a real wastewater, by the microalga Coelastrum sp., in a pilot-scale photobioreactor and a multibarrier treatment. The multibarrier treatment was the most effective treatment regarding the removal efficiencies. The photobioreactor showed different removal percentages depending on the initial effluent composition.
Hom-Diaz et al. [177] observed that the removal of pharmaceuticals from domestic wastewater in a pilot-scale tubular photobioreactor, in two seasonal periods (September–October; October–December), was highly impacted by temperature and solar irradiation.

The removal of several pharmaceuticals present in wastewater by green algal species, in a real-scale photobioreactor, was positively correlated with light intensity inside the culture, with stronger correlation when the data collected during the night were excluded [178].

Lai et al. [179] studied the removal of natural steroid estrogens (estradiol, estrone, estriol, and hydroxyestrone) and synthetic steroid estrogens (estradiol valerate, estradiol, and ethinylestradiol) by the alga *Chlorella vulgaris*, using batch-shaking experiments in the light and in the dark. The results showed that estradiol and estrone were interconvertible in the presence or absence of light. In the presence of light, 50% of estradiol was further metabolized to an unknown product. Estradiol valerate was hydrolyzed to estradiol and then to estrone. Estrone, hydroxyestrone, estriol and ethinylestradiol were relatively stable, and did not suffer biotransformation. Recently, a combination of photobioreactor and open-pond cultivation was suggested; the first allows a fast algae growth, while the second ensures mass cultivation [141].

In many cases, the complete removal of a compound requires the interaction of several groups of microorganisms (consortium), each of which is responsible for a degradation step. This interaction is extremely positive because pure cultures, isolated from this consortium, may not be able to completely remove this compound as the single carbon source, or the removal rate may be significantly lower than that obtained with the mixed crop that gave rise to it. Shi et al. [180] studied the removal of synthetic hormone $17\alpha$-ethinylestradiol and the natural hormones estrone and $17\beta$-estradiol, from a synthetic wastewater, by an alga (*Anabaena cylindracea, Chlorococcus, Spirulina platensis, Chlorella, Scenedesmus quadricauda*, and *Anabaena*) and duckweed (*Lemna*) pond system. The results showed that all hormones were effectively removed from the continuous-flow algae and duckweed pond even when their concentrations were at ng·L$^{-1}$ level. The simultaneous presence of algae and duckweed accelerated the removal of hormones from the synthetic wastewater since hormones could be quickly sorbed either on duckweed or algae and then degraded by both microorganisms [180].

Although several studies demonstrated the applicability of algae for micropollutant removal from wastewater [137,181], limitations and knowledge gaps still exist to rely on algae biomass production as an effective mean for micropollutant removal. For example, when Wang et al. [182] exposed the freshwater alga *Chlorella Pyrenoidosa* to a triclosan concentration of 800 mg·L$^{-1}$, a reductive dechlorination product of triclosan was formed. Algal cell growth was affected by their toxicity. Algal cell chloroplasts were damaged, decreasing the energy supply for algal growth, which produced an adverse effect on the effectiveness of triclosan biodegradation. Furthermore, most of the studies tested the algae’s ability to remove pollutants, which grew in unpolluted media before the treatment, and ignored the toxic stress caused by the pollutant and its influence on the removal capability. It is possible that the sensitivity or tolerance of algae changes after the first contact with the pollutant and, therefore, it may influence the removal efficiency of subsequent batch treatments.

Chen et al. [183] investigated the removal efficiency of cefradine by *Chlorella pyrenoidosa* in a sequencing batch reactor and identified different results between the first batch treatment and the second, where higher algal growth inhibition rates were observed; however, the alga produced more photosynthetic pigments, enhancing its photosynthetic metabolism as a way of adaptation to more harmful environmental conditions.

2.2.4. Concluding Remarks and Future Challenges

Some algae strains found in the literature [64,133,148,164,166,168–170,176–180,182–194] proved to be effective in pharmaceutical removal (Table 1).
Table 1. Bioremediation studies with fungi and algae for removal of pharmaceuticals. WWTP—wastewater treatment plant.

<table>
<thead>
<tr>
<th>Compounds</th>
<th>Compounds Source</th>
<th>Strains</th>
<th>Removal Mechanisms</th>
<th>Technologies</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ibuprofen, clofibric acid, and carbamazepine</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td></td>
<td>Biodegradation Adsorption</td>
<td>Laboratory-scale batch assays [46]</td>
</tr>
<tr>
<td>Carbamazepine</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td></td>
<td>Biodegradation Adsorption</td>
<td>Laboratory-scale batch assays Pilot-scale glass air-pulsed fluidized bioreactor (continuous and batch feed) [47]</td>
</tr>
<tr>
<td>Citalopram, fluoxetine, sulfamethoxazole, diclofenac, ibuprofen, naproxen, carbamazepine, and diazepam</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td></td>
<td>Biodegradation</td>
<td>Laboratory-scale batch assays [48]</td>
</tr>
<tr>
<td>Carbamazepine</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td></td>
<td>Biodegradation Biosorption</td>
<td>Pilot-scale plate bioreactor (continuous and batch feed) [49]</td>
</tr>
<tr>
<td>Naproxen, ibuprofen, acetaminophen, salicylic acid, ketoprofen, codeine, erythromycin, metronidazole, ciprofloxacin, azithromycin, cefalexine, propranolol, carbamazepine, 10,11-epoxycarbamazepine, 2-hydroxycarbamazepine, acridone, and citalopram</td>
<td>Urban wastewater</td>
<td>Fungi</td>
<td></td>
<td>Biodegradation</td>
<td>Pilot-scale air-fluidized bioreactor (batch feed) [50]</td>
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<tr>
<td>Carbamazepine, ibuprofen, clofibric acid, ketoprofen, metronidazole, trimetazosin, 17-α-ethinylestradiol, 17-β-estradiol-17-acetate, estrone, estriol, 17-β-estradiol, gemfibrozil, amitriptyline, primisodone, salicylic acid, diclofenac, naproxen</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td></td>
<td>Biodegradation Biosorption</td>
<td>Laboratory-scale batch assays Pilot-scale fungus-augmented membrane bioreactor (continuous feed) [51]</td>
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<td>Acetaminophen, ibuprofen, ketoprofen, naproxen, salicylic acid, codeine, phenazone, dexamethasone, diclofenac, piroxicam</td>
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<td>Fungi</td>
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<td>Biodegradation</td>
<td>Pilot-scale glass air-pulsed fluidized bioreactor (batch feed) [52]</td>
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<tr>
<td>X-ray contrast agent iopromide and antibiotic ofloxacin</td>
<td>Hospital wastewater</td>
<td>Fungi</td>
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<td>Biodegradation</td>
<td>Pilot-scale batch assays Pilot-scale glass air-pulsed fluidized bioreactor (batch feed) [53]</td>
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<td>Metronidazole, salicylic acid, primidone, amitriptyline, carbamazepine, ketoprofen, naproxen, ibuprofen, gemfibrozil, diclofenac, triclosan, estriol, estrone, 17-α-ethinylestradiol, 17-β-estradiol, 17-β-estradiol-17-acetate</td>
<td>Synthetic media</td>
<td>Fungi</td>
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<td>Biodegradation Biosorption</td>
<td>Laboratory-scale batch assays [54]</td>
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<td>Cefalexin, ciprofloxacin, metronidazole, trimethoprim, tetracycline, ketoprofen, acridone, carbamazepine, a carbamazepine metabolite, ciprofloxacin, metronidazole and its hydroxilated metabolite, β-blocker carazolol, diazepam, naproxen, cephalaxin, tetracyclin, sertraline, paroxetine, gemfibrozil, amlodipine, furosemide, dimetridazole, azithromycin, romnidazole, olanzapine, piroxicam, β-blockers metoprolol</td>
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<td>Biodegradation</td>
<td>Pilot-scale glass air-pulsed fluidized bioreactor (continuous and batch feed) [55]</td>
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Table 1. Cont.

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<th>Technologies</th>
<th>Reference</th>
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<td>Laboratory-scale batch assays</td>
<td>[56]</td>
</tr>
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<td>Estrone, 17β-estradiol, 17α-ethinyl-estradiol, and estriol</td>
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<td>Fungi Trametes versicolor, Myceliophthora thermophila</td>
<td>Biodegradation Adsorption</td>
<td>Laboratory-scale batch assays</td>
<td>[57]</td>
</tr>
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<td>17β-estradiol and 17α-ethinylestradiol</td>
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<td>Fungi Trametes versicolor</td>
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<td>Pilot-scale air-fluidized bioreactor (continuous feed)</td>
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<td>Sulfapyridine, sulfapyridine, and sulfamethazine</td>
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<td>Laboratory-scale batch assays Pilot-scale air-pulsed fluidized-bed bioreactor (continuous feed)</td>
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<td>Pilot-scale stirred tank reactor and fixed-bed reactor (continuous feed)</td>
<td>[64]</td>
</tr>
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<td>Tetracycline and oxytetracycline</td>
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<td>Pharmaceutical industry wastewater</td>
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</tr>
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<td>Diclofenac, ketoprofen and atenolol</td>
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<td>Fungi Pleurotus ostreatus</td>
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<td>Pilot-scale air-pulsed fluidized-bed bioreactor (continuous and batch feed)</td>
<td>[67]</td>
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<td>Laboratory-scale batch assays</td>
<td>[68]</td>
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<td>[69]</td>
</tr>
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<td>Carbamazepine and clarithromycin</td>
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<td>[70]</td>
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<td>Carbamazepine</td>
<td>Synthetic media</td>
<td>Fungi <em>Pleurotus ostreatus</em></td>
<td>Biodegradation</td>
<td>Laboratory-scale batch assays</td>
<td>[71]</td>
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<td>Clofibric acid, gemfibrozil, ibuprofen, fenoprofen, naproxen, diclofenac, indomethacin, propyphenazone, and carbamazepine</td>
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<td>Fungi <em>Trametes versicolor</em></td>
<td>Biodegradation</td>
<td>Laboratory-scale batch assays</td>
<td>[72]</td>
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<td>Levofloxacin</td>
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<td>Biodegradation, Photodegradation, Biosorption</td>
<td>Laboratory-scale batch assays</td>
<td>[133]</td>
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<td>Acetaminophen, ibuprofen, ketoprofen, naproxen, carbamazepine, diclofenac, and trielosan</td>
<td>WWTP wastewater</td>
<td>Algae <em>Lessonia nigrescens</em> Bory, <em>Macrocystis integrifolia</em> Bory</td>
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<td>Pilot-scale high-rate algal ponds</td>
<td>[148]</td>
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<td>Tetracycline</td>
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<td>Laboratory-scale batch assays, Pilot-scale high rate algal ponds (batch feed)</td>
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<td>Progesterone and norgestrel</td>
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<td>Algae <em>Scenedesmus obliquus</em>, <em>Chlorella pyrenoidosa</em></td>
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<td>Algae <em>Chlamydomonas mexicana</em>, <em>Scenedesmus obliquus</em></td>
<td>Biodegradation, Adsorption, Bioaccumulation</td>
<td>Laboratory-scale batch assays</td>
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Table 1. Cont.

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<th>Compounds</th>
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<th>Removal Mechanisms</th>
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<td>17α-Ethinylestradiol</td>
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<td>Algae Desmodesmus subspicatus</td>
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<td>[170]</td>
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<td>Algae Coelastrum sp.</td>
<td>Biodegradation</td>
<td>Pilot-scale photobioreactor microalgae and multi-barrier treatment</td>
<td>[176]</td>
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<td>Acetaminophen, ibuprofen, naproxen, salicylic acid, ketoprofen, codeine, azithromycin, erythromycin, ciprofloxacin, ofloxacin, atenolol, lorazepam, alprazolam, paroxetine, hydrochlorothiazide, furosemide, and diltiazem</td>
<td>Domestic wastewater</td>
<td>Algae Undefined microalgae</td>
<td>Biodegradation</td>
<td>Pilot-scale tubular photobioreactor</td>
<td>[177]</td>
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<td>Alfuzosin, alprazolam, atenolol, atracurium, azelastine, biperiden, bisoprolol, bupropion, carbamazepin, clazapril, ciprofloxacin, citalopram, clarithromycin, clemastine, clindamycin, clonazepam, clotrimazole, codeine, cyproheptadine, desloratadin, dicyclovetin, diltiazem, diphenhydramin, eprosartan, fexofenadine, flecainide, fluconazole, flutepitoxil, haloperidol, hydroxyzine, ibersartan, loperamide, memantin, metoprolol, miconazole, mirtazapine, nefazodon, orphenadrin, pizotifen, ranitidine, risperidone, rosithromycine, sertraline, sotalol, sulfamethoxazol, terbutalin, tramadol, trihexyphenidyl, trimetoprim, venlafaxin, and verapamil</td>
<td>WWTP wastewater</td>
<td>Algae Green algal species (Tetradesmus dimorphus and Dictyosphaerium, between them)</td>
<td>Biodegradation</td>
<td>Real-scale photobioreactor</td>
<td>[176]</td>
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<td>Estradiol, estrene, estriol and hydroxyestrenol) and synthetic steroid estrogens (estradiol valerate, estradiol, and ethinylestradiol</td>
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<td>Algae Chlorella vulgaris</td>
<td>Biotransformation</td>
<td>Laboratory-scale batch assays</td>
<td>[179]</td>
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<td>17α-ethinylestradiol, estrone, and 17β-estradiol</td>
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<td>Algae Anaebena cylindrical Chlorococcus Spirulina platensis Chlorella Scenedesmus quadricauda Anaebena (and duckweed Lemna)</td>
<td>Biodegradation Adsorption</td>
<td>Laboratory-scale batch assays Pilot-scale plug flow reactor (continuous feed)</td>
<td>[180]</td>
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<td>Triclosan</td>
<td>Synthetic media</td>
<td>Algae Chlorella Pyrenoidosa</td>
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<td>[182]</td>
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<td>Cefradinere</td>
<td>Synthetic media</td>
<td>Algae Chlorella pyrenoidosa</td>
<td>Biodegradation</td>
<td>Pilot-scale batch-sequencing reactor algae process (batch feed)</td>
<td>[183]</td>
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<td>Compounds</td>
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<td>Technologies</td>
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<td>Ethinylestradiol</td>
<td>Synthetic media</td>
<td>Algae&lt;br&gt;A. braunii&lt;br&gt;C. ellipsoidea&lt;br&gt;C. pyrenoidosa&lt;br&gt;C. vulgaris&lt;br&gt;S. communis&lt;br&gt;S. obliquus&lt;br&gt;S. quadricauda&lt;br&gt;S. vacuolatus&lt;br&gt;S. capricornutum</td>
<td>Biotransformation</td>
<td>Laboratory-scale batch assays</td>
<td>[184]</td>
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<td>17α-boldenone, 17β-boldenone, 4-hydroxy-androst-4-ene-17-dione, androsta-1,4-diene-3,17-dione, 4-androstene-3,17-dione, carbamazepine, ciprofloxacin, clarithromycin, clindamycin, clotrimazole, diclofenac, enrofloxacin, erythromycin–H2O, estrone, fluconazole, gemfibrozil, ibuprofen, lincomycin, loratadine, norfloxacin, ofloxacin, paracetamol, progesterone, roxithromycin, salicylic acid, salinomycin, sulfadiazine, sulfadimethoxine, sulfamerazine, sulfamethazine, sulfamethoxazole, sulfamonoxazole, sulfapyridine, testosterone, triclosan, trimethoprim, and tylosin</td>
<td>WWTP wastewater&lt;br&gt;Algae&lt;br&gt;C. reinhardtii&lt;br&gt;S. obliquus&lt;br&gt;C. pyrenoidosa&lt;br&gt;C. vulgaris</td>
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<td>Laboratory-scale batch assays</td>
<td>[185]</td>
<td></td>
</tr>
<tr>
<td>Salicylic acid and paracetamol</td>
<td>Synthetic media</td>
<td>Algae&lt;br&gt;C. sorokiniana&lt;br&gt;Chlorella vulgaris</td>
<td>Biodegradation</td>
<td>Pilot-scale reactor (batch and semicontinuous feed)</td>
<td>[186]</td>
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<td>Paracetamol, salicylic acid, and diclofenac</td>
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<td>Algae&lt;br&gt;C. sorokiniana&lt;br&gt;Chlorella vulgaris</td>
<td>Biodegradation</td>
<td>Pilot-scale bubbling column photobioreactor (batch and semicontinuous feed)</td>
<td>[187]</td>
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<td>Biodegradation&lt;br&gt;Adsorption&lt;br&gt;Absorption</td>
<td>Laboratory-scale batch assays</td>
<td>[188]</td>
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<td>[189]</td>
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<td>Strains</td>
<td>Removal Mechanisms</td>
<td>Technologies</td>
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<td>Oxytetracycline, doxycycline, chlortetracycline, and tetracycline</td>
<td>Synthetic media</td>
<td>Algae <em>Haematococcus pluvialis</em> <em>Chlorella</em> sp. <em>Selenastrum capricornutum</em> <em>Pseudokirchneriella subcapitata</em></td>
<td>Biodegradation</td>
<td>Laboratory-scale batch assays</td>
<td>[191]</td>
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<tr>
<td>Salicylic acid or paracetamol</td>
<td>Pharmaceutical industry wastewater</td>
<td>Algae <em>Chlorella sorokiniana</em></td>
<td>Biodegradation</td>
<td>Pilot-scale bubbling column photobioreactor (batch and semicontinuous feed)</td>
<td>[193]</td>
</tr>
<tr>
<td>Ciprofloxacin</td>
<td>Synthetic media</td>
<td>Algae <em>Chlamydomonas mexicana</em></td>
<td>Biodegradation</td>
<td>Laboratory-scale batch assays</td>
<td>[194]</td>
</tr>
<tr>
<td>β-estradiol and 17α-ethinylestradiol</td>
<td>WWTP anaerobic sludge</td>
<td>Algae <em>Selenastrum capricornutum</em> <em>Chlamydomonas reinhardtii</em></td>
<td>Biodegradation Photodegradation Adsorption</td>
<td>Laboratory-scale batch assays</td>
<td>[195]</td>
</tr>
</tbody>
</table>
Chen et al. [182] raised a question worthy for future researches related to the possible improvement of the removal efficiency of pollutants by algae under optimal light conditions and after their acclimation. The final step of the treatment, harvesting, which represents a significant amount of the production cost and energy consumption, is one of the most challenging processes. One of the most environmentally friendly methods is bioflocculation [150,151]; nevertheless, it may not be efficient enough due to the small size of microalgal cells. Other technologies, namely, co-pelletization through the addition of other microorganisms (bacterial and fungal species) to microalgae culture, which do not require the addition of chemicals or inputs of energy, may be promising solutions. However, further research on this topic is recommended.

Such as for mycoremediation, genetic tools can represent an essential step to improve algae performance. However, genetically modified algae for bioremediation are rarely reported. An example is a small laccase from *Streptomyces coelicolor* that was engineered by structure-based design and site-directed mutagenesis to improve its activity on commercially relevant substrates. The variants generated showed up to a 40-fold increased efficiency on 2,6-dimethoxyphenol, as well as the ability to use mediators with considerably higher redox potentials [195].

### 3. Biosorption

Adsorption includes all the processes involving a physical or chemical interaction between the surface of a solid material (adsorbent) and the pollutant (adsorbate); biosorption is a subcategory of adsorption, and may be simply defined as the removal of substances from solution by biological material (live or dead). It is a property of living and dead biomass to bind and abiotically concentrate compounds [196,197]. In addition to being an efficient and low-cost process, biosorption offers advantages over conventional processes, and avoids the use of chemicals, namely, nutrient supply [197].

#### 3.1. Biosorption Materials

Activated carbon is the most widely and effectively used sorbent, since it has a porous structure consisting of a network of interconnected macropores, mesopores, and micropores that provide a good capacity for the adsorption of organic molecules due to its high specific surface area. However, it is quite expensive, United States (US) $20–22/kg [198], and the higher the quality is, the greater the cost is [199,200], which limits its widespread use. According to the Swedish Environmental Protection Agency [201], the costs depend mainly on the size of the treatment plant. For facilities using granular activated carbon, for 2000–10,000 population equivalent, the following costs are associated: 0.72 million US $ for installation, 0.04 million US $/year of capital expenditure, 0.1 million US $/year of operating expenditure, and less than 0.01 kWh/m³ of operational electricity consumption, corresponding to a total cost of 0.09–0.11 US $/m³ of treated wastewater. It is verified that the annual operating expenditure has a major contribution to the total cost and that the major fraction of the operating costs is due to activated carbon, which demonstrates the need to look for low-cost adsorbents. This led to the search for low-cost sorbents, which require little processing and that are abundant in nature [202]. In this search, biosorbents from microorganisms were shown to be a promising alternative since, due to their reduced size, they present high specific surface area and they were shown to be able to sorb different organic and inorganic pollutants from solutions [203]. A wide range of microorganisms were studied in biosorption processes; these include microalgae and fungi. The complexity of the microorganism structure implies that there are many ways for the contaminants to be captured by the cells; in some cases, these are still not very well understood. The cell wall is the first component that meets the pollutants, where the solutes can be deposited on the surface or within the cell-wall structure.

Fungal cell walls are structurally complex with several functional groups such as carboxyl, hydroxyl, amino, sulfonate, and phosphonate, which bring about the excellent adsorption properties of fungi [204–208].

Algal cell walls exhibit some variations in their structure. The main groups present are amino, amine, hydroxyl, imidazole, phosphate, and sulfate [209]. This availability of functional groups
minimizes the selectivity problems and allows the high efficiency of removal to be more easily achieved [210]. For example, Navarro et al. [211] proposed the formation of hydrogen bonds as the main mechanism for the removal of sulfamethoxazole and sulfacetamide by marine algae Lessonia nigrescens Bory and Macrocystis integrifolia Bory.

3.2. Treatment Systems

Biosorption can proceed in batch or continuous mode. Both modes are frequently employed to conduct laboratory-scale experiments, while continuous mode is the most employed for industrial applications. Packed-bed, fluidized-bed, and continuous-stirred-bed reactors are three types of design used in biosorption experiments. Several studies showed that packed-bed columns are the most suitable for liquid–solid separation, and scaling was found to be minimal [212,213]. Fluidized-bed and continuous-stirred-bed reactors are only occasionally used, since fluidized-bed reactors require a high flow rate, which is sometimes difficult to achieve, and stirred-bed reactors require that biomass be in powdered form; beyond that, they have high associated costs and high requirements of operation and maintenance [207,214].

In the last few decades, many patents were developed, focused on improving the sorption capacity of biosorbents through their modification or immobilization, mainly for metal removal [197]. However, further attention should be given to the removal of pharmaceuticals.

3.3. Mechanisms of Removal

Biosorption is a physico-chemical process that involves several mechanisms (e.g., adsorption, ion exchange, surface complexation, and microprecipitation), as shown in Figure 4 [215]. Biosorbent surfaces are characterized by active, energy-rich sites that are able to interact with compounds in the adjacent aqueous phase due to their specific electronic and spatial properties. Sorption occurs because, from the thermodynamic point of view, the molecules prefer to be in a low-energy state. A molecule sorbed onto a surface has a lower energetic state on a surface than in the aqueous phase. Therefore, the molecule is attracted to the surface and to a lower-energy state. The attraction of a molecule to a surface can be caused by physical and/or chemical forces. Electrostatic forces govern the interactions between most sorbates and biosorbents. These forces include dipole–dipole interactions or London/van der Waals forces, and hydrogen bonds [216].

![Figure 4. Biosorption mechanisms.](image)

Biosorption equilibrium is not established instantaneously. The mass transfer from the solution to the sorption sites within the adsorbent particles is constrained by mass transfer resistances that determine the time required to reach the state of equilibrium. The rate of adsorption is usually limited by diffusion processes toward the external adsorbent surface and within the porous adsorbent particles.

The progress of the biosorption process includes four consecutive steps: transport of the sorbate from the bulk liquid phase to the hydrodynamic boundary layer localized around the biosorbent particle; transport through the boundary layer to the external surface of the biosorbent, termed film
diffusion or external diffusion; pore diffusion and/or surface diffusion toward the interior of the biosorbent particle (termed intraparticle diffusion or internal diffusion); and energetic interaction between the sorbate molecules and the sorption sites.

Kinetic and equilibrium data are modeled using different approaches in order to explain the biosorption mechanism of the pollutants’ removal [217]. The equilibrium distribution on the sorbed pollutant between the biosorbent and the aqueous phase is required to determine the maximum biosorbent’s uptake capacity. The sorption kinetics provides additional important information about the sorption mechanism, especially the rate of pollutant removal. In wastewater treatment, kinetics information is important for setting an optimum residence time of the wastewater at the biosolid phase interface.

Typically, the dependence of the sorbed amount on the equilibrium concentration is determined experimentally at constant temperature, and the measured data are subsequently described by an appropriate isotherm equation.

For fixed-bed columns, the dynamic behavior and efficiency are described in terms of the effluent/affluent concentration as a function of time (or volume of treated liquid), i.e., the breakthrough curve. Experimental “breakthrough curve” determination allows verifying the applicability of a chosen adsorption model for a given biosorbent/sorbate system and estimating the related mass transfer coefficients [217].

3.4. Factors That Affect the Process

Several factors affect biosorption; most of them are common to adsorption processes, like pH, temperature, contaminant concentration, nature of sorbent and sorbate, contact time, ionic strength, the presence of other compounds, the use of dead or alive biomass [196,214,218], and the presence/absence of metabolic processes, in the case of living cells [215]. In living organisms, metabolic processes may affect physico-chemical biosorption mechanisms, pollutant bioavailability, chemical speciation, and accumulation/transformation [196].

Organic pollutants have a wide diversity of chemical structures; therefore, factors such as the molecular size, charge, solubility, hydrophobicity, and reactivity are important factors in biosorption. Human pharmaceutical metabolites, for example, are usually more polar and hydrophilic than the parent compounds; therefore, it is expected that they will not be significantly removed by sorption [219].

By changing the properties of the liquid phase (e.g., concentration, temperature, pH), sorbed species may be released from the surface and transferred back into the liquid phase. This reverse process is referred to as desorption.

In the study of Aksu and Tunc [220], the removal of penicillin G potassium salt by dried fungus *Rhizopus arrhizus* was strongly dependent on the pH and temperature. Maximum sorption was observed at initial pH values of 6.0 and at 35 °C. They also [220] explored the possibility of using the dried fungus for the batch removal of penicillin G potassium salt from aqueous solution and compared the experimental uptake with the results obtained with powdered activated carbon, in the same conditions. The uptake capacity determined was 459.0 mg·g⁻¹ and 375.0 mg·g⁻¹ for dried *Rhizopus arrhizus* and for an activated carbon (commercial powdered activated carbon, SIGMA, C-4386, washed with hydrochloric acid), respectively.

The use of dead biomass has advantages compared to the use of live biomass, such as the absence of toxicity limitations and the absence of nutrient requirement for microorganism’s growth; however, live biomass has the advantage of using metabolic processes that complement the overall removal process [220].

Tam et al. [188] investigated tributyltin removal from artificial wastewater by dead and live cells of four microalgal species, *Chlorella miniata, Chlorella sorokiniana, Scenedesmus dimorphus,* and *Scenedesmus platydiscus.* In general, dead cells were more efficient than live cells during the three days of exposure; however, at the end of 14 days, removal efficiencies were identical to those that were reached by live cells. More than 90% and 85% of tributyltin was removed by dead cells of *Scenedesmus*
and Chlorella, respectively, in the first three days. After three days of treatment, total amount of sorbed tributyltin per unit of biomass for dead cells of Chlorella miniata, Chlorella sorokiniana, Scenedesmus dimorphus, and Scenedesmus platydiscus was, respectively, 11.3 ± 0.6, 4.0 ± 0.3, 1.31 ± 0.09, and 0.35 ± 0.04 mg·g⁻¹, while, for live cells, it was, respectively, 0.24 ± 0.03, 0.58 ± 0.12, 0.083 ± 0.003, and 0.0603 ± 0.0005 mg·g⁻¹. Many other studies allowed observing that the fast concentration decrease in the beginning of the assays was caused by the sorption process, followed by a much slower concentration decrease caused by other biological removal mechanisms. This fact is probably due to the large surface area of the individual cells, which provides a large interface between the aqueous and cell phases. Nevertheless, unlike the other removal mechanisms, sorption is an equilibrium process that ends when equilibrium is reached, which may lead to an incomplete removal of pollutants [164,180,182,189]. Combining biosorption with other removal mechanisms seems to be a successful strategy for the removal of pharmaceuticals.

3.5. Biosorption Potential as a Wastewater Treatment Technology

The potential of biosorption as a wastewater treatment technology was frequently cited in the literature, as shown in Table 2. Usually, the performance of commercial adsorbents is very good due to the high specific surface area and the high porosity, but they present high costs which prevents their wide use, especially when high flowrates are involved. The use of biosorbents, which are usually easily available, presents advantages such as their low cost, and a more eco-friendly and sustainable nature. Like bioremediation, several aspects make biosorption a promising choice, since it is a low-cost treatment, has low operating costs, does not produce chemical sludge to handle, and is highly efficient [221]. Some disadvantages associated with the process are the low specificity and low robustness of the systems [222], the impossibility of long-term use of the suspended biomass, and the difficult separation of the biomass and the treated effluent [223]. The use of packed-bed columns is considered the most suitable biosorption, allowing an easy liquid–solid separation. The use of biosorbents in granulated form instead of powdered form presents the advantage of avoiding column clogging. Furthermore, the immobilization of fungi and algae can increase the efficiency of the process as it allows their use for several cycles [224].
Table 2. Biosorption studies with fungi and algae for removal of pharmaceuticals.

<table>
<thead>
<tr>
<th>Compounds</th>
<th>Compounds Source</th>
<th>Strains</th>
<th>Removal Mechanisms</th>
<th>Technologies</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ibuprofen, clofibric acid, and carbamazepine</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td>Biodegradation and Biosorption</td>
<td>Laboratory-scale batch assays</td>
<td>[46]</td>
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<tr>
<td>Carbamazepine</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td>Biodegradation and Biosorption</td>
<td>Pilot-scale glass air-pulsed fluidized bioreactor (continuous and batch feed)</td>
<td>[47]</td>
</tr>
<tr>
<td>Carbamazepine</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td>Biodegradation and Biosorption</td>
<td>Pilot-scale plate bioreactor (continuous and batch feed)</td>
<td>[49]</td>
</tr>
<tr>
<td>Carbamazepine, ibuprofen, clofibric acid, ketoprofen, metronidazole,</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td>Biodegradation and Biosorption</td>
<td>Pilot-scale batch assays</td>
<td>[51]</td>
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<tr>
<td>triclosan, 17-α-ethinylestradiol, 17-β-estradiol-17-acetate, estrone,</td>
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<td>estriol, 17-β-estradiol, gemfibrozil, amitriptyline, primidone, salicylic acid, diclofenac, naproxen</td>
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<td>Estrone, 17β-estradiol, 17α-ethinylestradiol, and estriol</td>
<td>WWTP wastewater</td>
<td>Fungi</td>
<td>Biodegradation and Biosorption</td>
<td>Laboratory-scale batch assays</td>
<td>[57]</td>
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<tr>
<td>Sulfaipyridine, sulfapyridine, and sulfamethazine</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td>Biodegradation and Biosorption</td>
<td>Pilot-scale air-pulsed fluidized bed bioreactor (continuous feed)</td>
<td>[59]</td>
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<tr>
<td>Diclofenac, ibuprofen, naproxen, carbamazepine, and diazepam</td>
<td>Synthetic media</td>
<td>Fungi</td>
<td>Biodegradation and Biosorption</td>
<td>Pilot-scale continuous stirred tank reactor and fixed-bed reactor</td>
<td>[64]</td>
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<tr>
<td>Diclofenac, ketoprofen, and atenolol</td>
<td>Hospital wastewater</td>
<td>Fungi</td>
<td>Biodegradation and Biosorption</td>
<td>Pilot-scale air-pulsed fluidized bed bioreactor (continuous and batch feed)</td>
<td>[67]</td>
</tr>
<tr>
<td>Compounds</td>
<td>Compounds Source</td>
<td>Strains</td>
<td>Removal Mechanisms</td>
<td>Technologies</td>
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<td>17α-ethinylestradiol and carbamazepine</td>
<td>Synthetic media</td>
<td>Fungi, Pleurotus sp. P1, Pleurotus ostreatus BS (unidentified) basidiomycete strain BNI</td>
<td>Biodegradation, Adsorption</td>
<td>Laboratory-scale batch assays</td>
<td>[68]</td>
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<td>Tetracycline</td>
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<td>Algae, Chlorella vulgaris</td>
<td>Photodegradation, Biosorption</td>
<td>Laboratory-scale batch assays, Pilot-scale high-rate algal ponds (batch feed)</td>
<td>[164]</td>
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<tr>
<td>Progesterone and norgestrel</td>
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<td>Algae, Scenedesmus obliquus, Chlorella pyrenoidosa</td>
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<td>Laboratory-scale batch assays</td>
<td>[168]</td>
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<td>Biodegradation, Adsorption, Biaccumulation</td>
<td>Laboratory-scale batch assays</td>
<td>[169]</td>
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<td>Acetaminophen, ibuprofen, ketoprofen, naproxen, carbamazepine, diclofenac, and triclosan</td>
<td>WWTP wastewater</td>
<td>Algae, Lessonia nigrescens Bory, Macroptila integrifolia Bory</td>
<td>Biodegradation, Photodegradation, Biosorption</td>
<td>Pilot-scale high-rate algal ponds</td>
<td>[175]</td>
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<td>17α-ethinylestradiol, estrone, and 17β-estradiol</td>
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<td>Algae, Anabaena cylindrica, Chlorococcus, Spirulina platensis, Chlorella, Scenedesmus quadricauda, Anabaena (and duckweed Lemna)</td>
<td>Biodegradation, Adsorption</td>
<td>Laboratory-scale batch assays, Pilot-scale plug flow reactor (continuous feed)</td>
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<td>Synthetic media</td>
<td>Algae, Chlorella Pyrenoidosa</td>
<td>Biodegradation, Biosorption</td>
<td>Laboratory-scale batch assays</td>
<td>[182]</td>
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<td>Tributyltin</td>
<td>Synthetic media</td>
<td>Algae, Chlorella miniata, Chlorella sorokiniana, Scenedesmus dimorphus, Scenedesmus platydiscus</td>
<td>Biodegradation, Adsorption, Absorption</td>
<td>Laboratory-scale batch assays</td>
<td>[188]</td>
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<td>Trimethoprim, sulfamethoxazole, and triclosan</td>
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<td>Algae, Nannochloris Sp.</td>
<td>Biodegradation, Adsorption</td>
<td>Laboratory-scale batch assays</td>
<td>[189]</td>
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<tr>
<td>Compounds</td>
<td>Compounds Source</td>
<td>Strains</td>
<td>Removal Mechanisms</td>
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<td>Diclofenac, ibuprofen, paracetamol, metoprolol, carbamazepine and</td>
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<td>Algae Chlorella sorokiniana</td>
<td>Biodegradation</td>
<td>Laboratory-scale batch</td>
<td>[190]</td>
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<td>trimethoprim, estrone, 17β-estradiol, and ethinylestradiol</td>
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<td>Chlamydomonas reinhardtii</td>
<td>Photodegradation</td>
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<td>Adsorption</td>
<td>Laboratory-scale batch</td>
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<td>Fungi Rhizopus arrhizus</td>
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<td>Paracetamol</td>
<td>Synthetic media</td>
<td>Algae Alga Synechocystis sp.</td>
<td>Biosorption</td>
<td>Laboratory-scale batch</td>
<td>[226]</td>
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<td>assays Pilot-scale</td>
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<td>photobioreactor</td>
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</table>
4. Conclusions and Future Perspectives

The urban wastewater treatment industry is facing challenges, such as the fate of pharmaceuticals, which induce the development of wastewater treatment alternatives. In this review, an overview was drawn of fungi and algae potential to remove pharmaceuticals by bioremediation and biosorption processes from aquatic matrices. Algae present interesting advantages when compared with fungi, as they have a very fast growth and are able to remove both pharmaceuticals and nutrients, which is quite important since tertiary treatments are often needed for this purpose in domestic wastewater treatment. Moreover, this biomass is a valuable bioresource for the production of biofuel or high-value by-products. Bioremediation and biosorption studies proved to be eco-friendly and low-cost promising alternatives for pharmaceutical removal when compared to conventional methods applied for the same purpose. However, most of these studies remain at a laboratory-scale and are performed with synthetic media when it is well known that wastewaters are complex matrices that differ between WWTPs and can vary per geographical region. Only a few researchers are working on real field applications, since extensive research is still needed in order to understand the complexity of the processes, their dependence on physico-chemical and biological factors, and the mechanisms involved, in order to ensure high efficiencies for pharmaceutical removal. New knowledge in genetic engineering should be introduced in order to select and amplify the most effective algae or fungi strains for pharmaceutical removal. Several microorganisms contain key metabolic genes that could be introduced into other organisms. Genetically modified fungi and algae, armed with new or increased capacities for degrading various compounds, will likely have an important future in this field since it will allow these microorganisms to effectively remove pharmaceuticals from wastewaters and also allow their continuous use, considering legal limitations, as well as a pragmatic market and cost rationale.

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