

CRITICAL REVIEW

View Article Online
View Journal | View Issue



Cite this: *Environ. Sci.: Adv.*, 2024, **3**, 1500

Laccase-mediated degradation of emerging contaminants: unveiling a sustainable solution

Pooja Thathola,^{ID} *^{ab} Elda M. Melchor-Martínez,^{cd} Priyanka Adhikari,^e Saúl Antonio Hernández Martínez,^d Anita Pandey^f and Roberto Parra-Saldívar^{ID} ^{cdg}

The excessive use of emerging contaminants (ECs) in various applications has led to a global health crisis. ECs are found in groundwater, surface water, soils, and wastewater treatment plants at concentrations ranging from ng L⁻¹ to µg L⁻¹. This review explores the sources of ECs and laccase's role in their degradation. ECs encompass diverse categories with potential implications for human health, animals, and the environment, and their adverse effects are examined. Laccase, a key mediator, can oxidize non-phenolic compounds, broadening its substrate range. The review discusses the intricacies of laccase-mediated degradation and highlights its potential to improve global water resource sustainability. Innovative strategies, like laccase immobilization, are explored for EC removal, benefiting environmental preservation. In summary, the review addresses the issue of excessive EC use, their presence in water sources, and their impact on health, wildlife, and the ecosystem. Laccase offers promise for EC degradation, emphasizing its mechanism and potential for sustainable water resource management. Advanced techniques, including laccase immobilization, further demonstrate the commitment to tackling EC-induced environmental challenges.

Received 28th May 2024
Accepted 7th September 2024

DOI: 10.1039/d4va00173g

rsc.li/esadvances

Environmental significance

The review on Laccase-Mediated Degradation of Emerging Contaminants unveils a promising sustainable solution to the growing concern of environmental pollution caused by novel, unregulated compounds. Emerging contaminants, including pharmaceuticals, personal care products, and industrial chemicals, pose significant risks to ecosystems and human health due to their persistence and bioaccumulative potential. Traditional water treatment methods often fail to effectively remove these pollutants. Laccase, a naturally occurring enzyme, offers an eco-friendly and efficient approach to degrade these contaminants. By harnessing the catalytic power of laccase, this research demonstrates a viable pathway to mitigate the environmental impact of emerging contaminants, promoting cleaner water bodies and healthier ecosystems. The laccase-mediated process not only aligns with green chemistry principles but also supports the advancement of sustainable water treatment technologies, contributing to the broader efforts for environmental protection and public health safety.

1. Introduction

Occurrence of emerging contaminants (ECs) in the environment has become a significant concern due to their potential

adverse effects on both humans and aquatic animals. The concentration of these ECs in soil and water bodies varies from a few nanograms to several micrograms per liter (µg L⁻¹).^{1,2} These ECs (Table 1) are present in trace levels, and they are categorized into different types, *i.e.*, pesticides, steroid hormones, pharmaceuticals, and chemicals from industries. Phytoestrogens are also one of the active ECs, which primarily constrain the sexual and reproductive development of wildlife.^{2,3} Therefore, the remediation of ECs has become the most significant concern in the current age. This portends a potential impact on broader ecology as well as human health (Fig. 1). Due to concerns about widespread human exposure and the potentially harmful health effects, or environmental impact of ECs, research on the treatment of these contaminants has recently gained increasing interest.⁴ Different chemical and physical methods, such as photodegradation,⁵ Fenton treatment,⁶ electrochemical treatment,⁷ ozonation,⁸ thermal degradation, and advanced oxidation processes,⁹ have been implemented for the treatment of ECs. However, several

^aCenter for Land and Water Resource Management, Govind Ballabh Pant National Institute of Himalayan Environment, Kosi-Katarmal 263643, Almora, Uttarakhand, India. E-mail: poojathathola29@gmail.com

^bAnalytical and Environmental Science Division and Centralized Instrument Facility, CSIR-Center Salt and Marine Chemicals Research Institute, Bhavnagar 364002, Gujarat, India

^cInstitute of Advanced Materials for Sustainable Manufacturing, Tecnológico de Monterrey, Monterrey 64849, Mexico

^dSchool of Engineering and Sciences, Tecnológico de Monterrey, Monterrey 64849, Mexico

^eCenter for GMP Extraction Facility (Dept. of Biotechnology), National Institute of Pharmaceutical Education and Research, Changsari 781101, Guwahati, Assam, India

^fDepartment of Biotechnology, Graphic Era Deemed to be University, Dehradun 248002, Uttarakhand, India

^gFacultad de Agronomía, Universidad Autónoma de Nuevo León, Francisco Villa, 66059 Colonia Ex hacienda El Canadá, General Escobedo, Nuevo León, Mexico



Table 1 Frequently detected emerging contaminants in water bodies with their molecular formula and structure

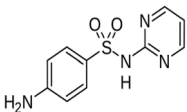
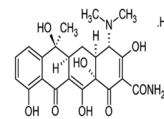
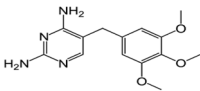
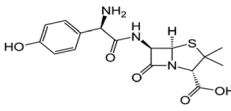
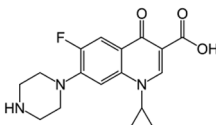
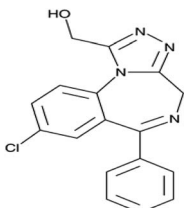
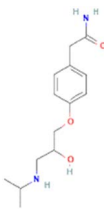
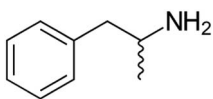
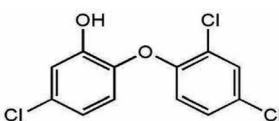
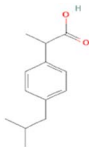
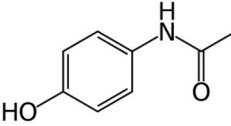
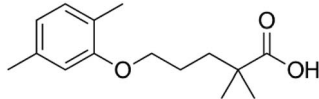
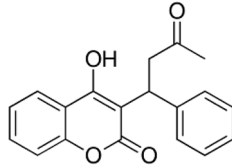
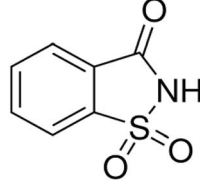
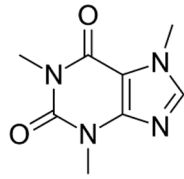
S. no	Emerging contaminants	Molecular formula	Structure
1	Sulfadiazine	$C_{10}H_{10}N_4O_2S$	
2	Tetracycline	$C_{22}H_{24}N_2O_8$	
3	Trimethoprim	$C_{14}H_{18}N_4O_3$	
4	Amoxicillin	$C_{16}H_{19}N_3O_5S$	
5	Ciprofloxacin	$C_{17}H_{18}FN_3O_3$	
6	α -Hydroxyalprazolam	$C_{17}H_{13}ClN_4O$	
7	Atenolol	$C_{14}H_{22}N_2O_3$	
8	Amphetamine	$C_9H_{13}N$	
9	Triclosan	$C_{12}H_7Cl_3O_2$	
10	Ibuprofen	$C_{13}H_{18}O_2$	
11	Paracetamol	$C_8H_9NO_2$	



Table 1 (Contd.)

S. no	Emerging contaminants	Molecular formula	Structure
12	Gemfibrozil	$C_{15}H_{22}O_3$	
13	Warfarin	$C_{19}H_{16}O_4$	
14	Saccharin	$C_7H_5NO_3S$	
15	Caffeine	$C_8H_{10}N_4O_2$	

limitations, including high cost, generation of toxic by-products, and high energy consumption hinder their implementation.¹⁰ Thus, developing novel, efficient, and greener remediation processes has become an important objective for researchers, industrial chemists, and the scientific community. In this context, biodegradation appears to be a greener, sustainable, and cost-effective approach for treating ECs.¹¹

There are many innovations in the degradation of these pollutants, which mainly include filtration technology, advanced oxidation processes (AOPs), membrane bioreactors (MBRs), and adsorption and absorption processes. These methods are mainly based on their physical, chemical, and biological mechanisms. In the filtration process, ECs are effectively separated and studied in both pilot and full-scale systems.^{12,13} But the treatment of these ECs separated by filtration technology is also a main concern as it involves a high concentration of ECs. Adsorption and AOPs have also increased attention due to their efficiency in removing ECs.³ Nonetheless, these methods are normally used after primary or secondary treatments due to the higher content of organic carbon and turbidity. MBR technologies are an alternative to the conventional activated sludge processes for treatment. Many researchers have reported that MBRs effectively remove contaminants, but they are less effective in removing ECs.^{14,15} Recently, the enzymatic transformation and degradation of ECs using microorganisms have been determined as a promising and eco-friendly approach.¹⁶ Enzymes from microorganisms are probable biological agents for the degradation and transformation of these ECs.

Among all the reported enzymes, laccase is one of the lignin-modifying enzymes responsible for the effective degradation of these ECs. These enzymes are applied in many processes due to their oxidation capacity.¹⁷ Laccase enzymes in their crude, purified, and immobilized forms have been used for the degradation of several types of ECs in all categories of experiments and trials (*i.e.*, continuous and batch experiments).¹⁸ These enzyme technologies are widespread and have promising applications in environmental biotechnology. Based on this, the present review aims to gather current trends and information on ECs, sources, harmful effects, and the role of laccase enzyme in the degradation process.

2. Aim and scope

This review will address emerging pollutants, their sources, and laccase-related studies with an emphasis on bioremediation. Laccases are responsible for breaking down these compounds without producing any harmful metabolites. Laccase enzyme-based bioremediation is highly effective and specific for treating pollutants because these enzymes have a higher efficiency of degradation as compared to the microorganisms as a whole. The quality of enzyme-based bioremediation, *i.e.*, it requires mild conditions and low energy, makes it more suitable for the field of bioremediation.^{19,20} Laccase belongs to the diverse group of enzymes/biocatalysts within the oxidoreductase group which is produced by many microorganisms (*e.g.*, fungi and bacteria). It has several biotechnological applications ranging from lignin degradation to organic pollutant biodegradation.



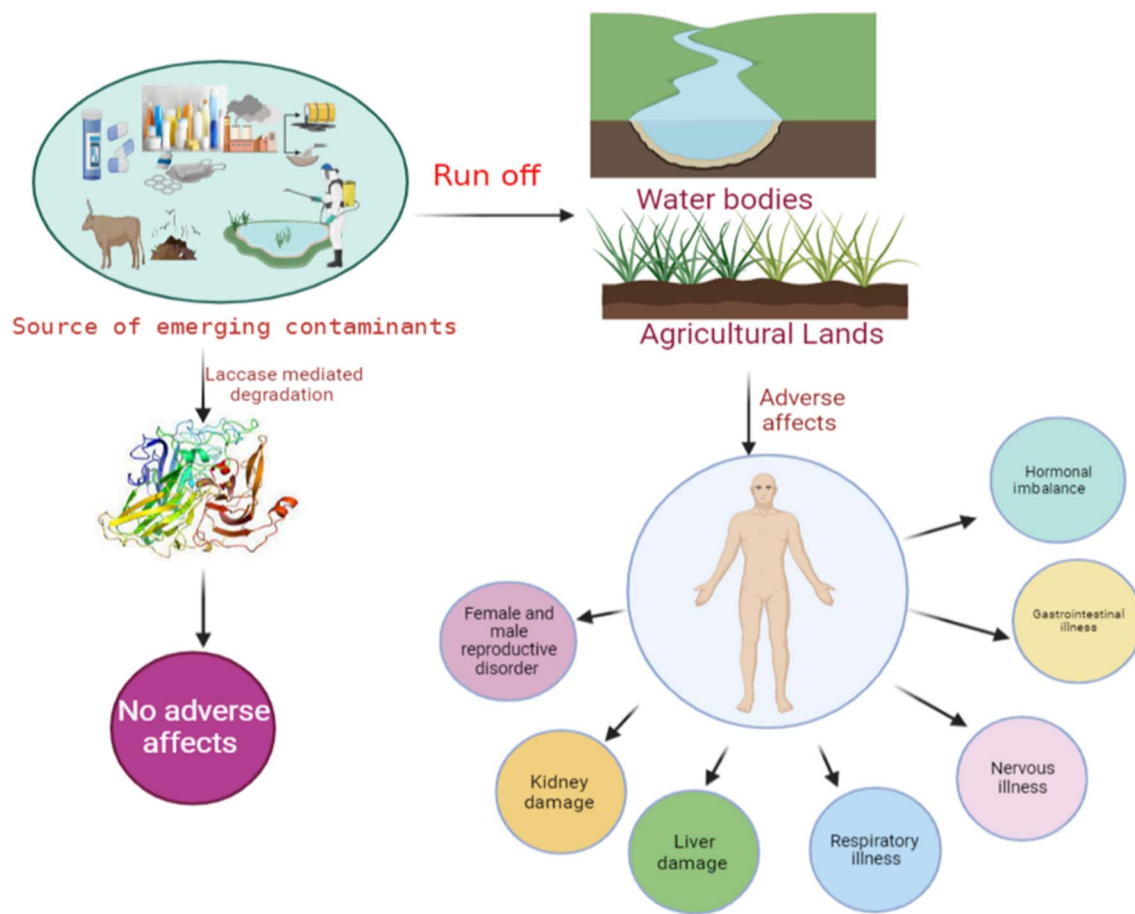


Fig. 1 Harmful effects of emerging contaminants.

Laccases are also involved in the transformation of various groups of ECs which mainly include polyaromatic hydrocarbons (PAHs), pharmaceutical and personal care products (PPCPs), phenols, and dyes.^{21,22} Although the applications of laccase degradation are well proven, most of the studies are limited to the lab scale. So further emphasis on pilot scale studies is necessary for economic feasibility. Therefore, this review expands upon bioremediation studies which will be further contextualized to address future studies. This review also focuses on immobilised laccase for bioremediation with different synthesis methods, hypothesized uptake mechanisms, and trends. Furthermore, the use of laccases as biocatalysts is discussed. Immediate gaps in the laccase bioremediation are highlighted to guide future studies, and potential mechanisms of laccase degradation are also discussed.

3. Emerging contaminants and their sources

The annual production of ECs in both developed and developing countries has been steadily increasing over the past few decades, largely due to their persistent nature. They are commonly used in agriculture, pharmaceuticals, personal care products, and hormones.²³ These compound residues are

incessantly released into the water bodies *via* runoff and wash-off from hospitals and households to the surface, contaminating the agricultural land. Additionally, surface water contaminants enter groundwater bodies through leaching and filtration. Furthermore, the other routes for ECs to enter the diverse water bodies are unused medicine disposal, irrigation methods using wastewater, disposal of animal carcasses, and the treatment plants with wastewater and sewage water.^{16,23} The discharge of these ECs from wastewater plants is a major route for their entry into the aquatic environment.²⁴ Poor management strategies and inefficient treatment technologies for EC pollution may lead to the release of these ECs into the environment (Fig. 1). Disposal of solid waste into the environment is the most common practice nowadays, which mainly increases the EC pollution in both the environment and wastewater treatment plants. However, the concentration is in the range of pg L^{-1} .²⁵ Several studies have indicated the presence of sweeteners. For instance, acesulfame, saccharin, and sucralose have been reported to be present in domestic wastewater *via* human excretion.²⁶ On the other hand, food preservatives, *i.e.*, parabens, have also been detected in groundwater.²⁷ Surface water and groundwater are usually used for drinking in developing countries and some developed countries. These water bodies are a common source of ECs in drinking water. The presence of ECs in water bodies is a common source of contamination in drinking water. The concentration of herbicides detected in drinking water



suggests that agricultural runoff is the primary pathway for transporting these contaminants from agricultural land to water bodies and finally to drinking water sources. These contaminants are detected in higher concentrations. The standard and frequently detected contaminants are atrazine, and terbuthylazine.²⁸

Non-sewered sanitation methods which mainly include urine-diverted dry toilets (UDDTs) are identified as a major source of EC pollution in developed countries. UDDTs, a means to separate human urine toilets, are built on a pilot scale in many developed countries. Still, there are few reports available on ECs from UDDTs. Ref. 29 described the presence of 12 ECs in source-separated urine. Pharmaceuticals are the major ECs detected, including hydrochlorothiazide, clarithromycin, darunavir, atenolol, atazanavir, atenolol acid, acetyl-sulfamethoxazole, diclofenac, emtricitabine, N4-ritonavir, and trimethoprim (TMP).³⁰

The management strategies for these ECs are very poor, so these ECs are also detected in landfill leachate, which increases the possibility of these ECs contaminating freshwater sources. Consequently, the constant monitoring of these landfill leachates should be done, as it is the major factor in detecting these ECs in water bodies.³¹ Moreover, proper disposal of these ECs in the environment and the treatment strategies for these landfill leachates are growing concerns. Untreated disposal of this leachate into the sewer is a common practice (Environmental Affairs 2019), which may lead to loading higher EC concentrations onto municipal wastewater treatment plants (WWTPs). Many studies have reported the presence of ECs in landfill leachates. Daso *et al.* reported that landfill leachates in Cape Town are the main source of polybrominated diphenyl ether (PBDE) in waterbodies and the environment.³² Olukunle *et al.* described the presence of these ECs in the landfill at concentrations in the range of pg L^{-1} .³³

4. Adverse effects of ECs on the environment

ECs have an enormous destructive impact on the environment; when these environmental pollutants come in contact with the ecosystem, they tend to show diverse effects. These pollutants probably cause many acute and long-term effects (*i.e.*, disruption of endocrine function, immunotoxicity, neurological disorders, cancers, *etc.*) on aquatic life, human health, and the environment, (Fig. 2). Increased release of these ECs in the environment provides many routes for their entry into the aquatic environment.³⁴ ECs are incessantly discharged into the environment but they are efficiently removed only sometimes; subsequently many of these contaminants persist in the environment for a long time.^{35,36} Improper and incomplete degradation of these contaminants on waste treatment plants during the degradation process are one of the main reasons for contaminants to enter into aquatic environment.³⁷ The risks associated with these ECs are chronic for living organisms, and therefore, they may cause lethal effects even at very low concentrations.³⁸ Creatures or living bodies in the aquatic environment are continuously exposed to the mixtures of these ECs, and these contaminants may interact with aquatic organisms and produce synergistic or toxic effects.³⁹ ECs are universally found in the environment primarily in soil and water bodies. These compounds, including organic compounds, brominated flame retardants, phthalates, plastics, microplastics, hormones, and polycyclic siloxanes, are responsible for causing deleterious effects on the health of humans and aquatic animals.⁴⁰ These contaminants are considered one of the foremost reasons for endocrine disorders in humans, which mainly include the modifications of salivary glands, the thyroid, and various genetic disorders in the male body system. These ECs

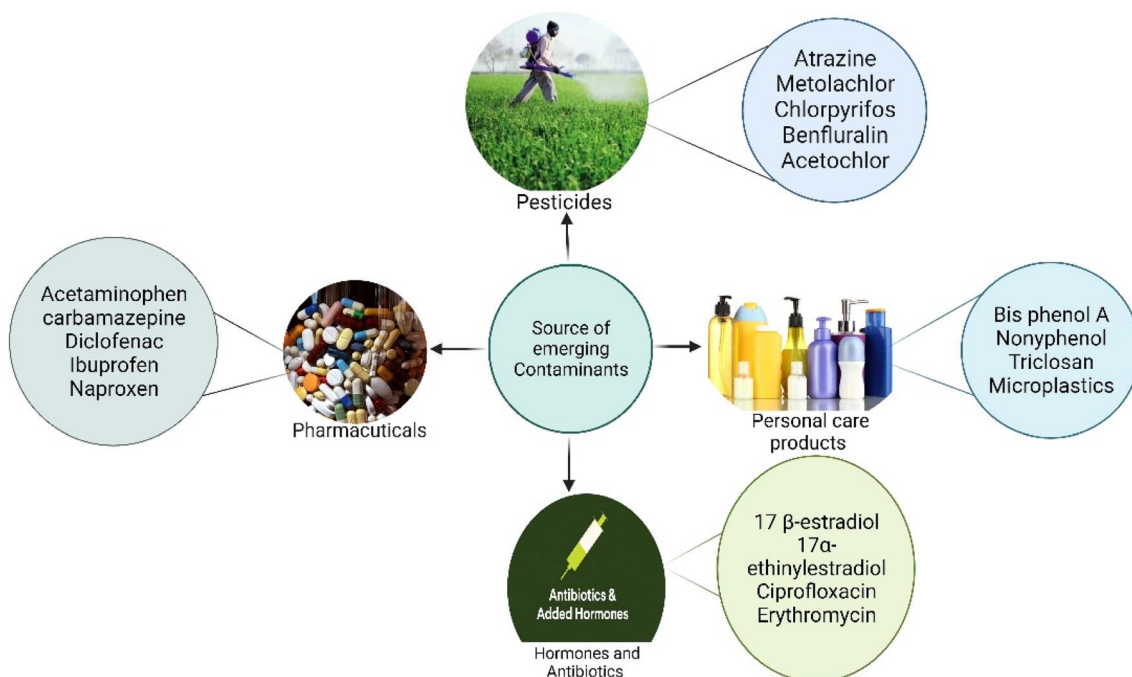


Fig. 2 Source of emerging contaminants in water bodies.



increase the risk of cancer in humans and act as antiandrogens in males, leading to feminizing effects.⁴¹ EC toxicity chiefly affects the endocrine cycle; many groups of these ECs may cause hormonal imbalance, DNA damage, carcinogenic effects, disruptions in thyroid function, impaired brain development, reduced fertility, and overall growth issues. Moreover, the genetic toxic effects of these ECs are abnormality in female implantation, breasts, and the liver, which may lead to increased hormones due to brominated flame retardants.⁴² Their excessive concentrations have been reported to cause abortion and maternity difficulties.⁴³

5. Laccase enzyme

Laccases (EC 1.10.3.2) are monomeric glycoproteins that have great potential in the field of environmental microbiology.^{44–46} Over the last few years, the use of enzyme technology has attracted great interest in the field of green and sustainable technologies. Laccase acts as a biocatalyst in conventional chemical-based processes and pharmaceutical industries.^{39,40} The use of laccases in the industrial and biotechnological sectors is a thriving area of research nowadays.^{47,48} The use of Laccases in industries, *i.e.*, food, textile, paper, *etc.*, as oxidative enzymes due to their ability to act on a wide range of substrates.^{49,50} Laccase enzymes provide an eco-friendly substitute for conventional chemical reactions. Laccase was first discovered in the nineteenth century in the Japanese lacquer tree *Rhus vernicifera*.⁵¹ Although the presence of laccases has been reported in higher plants, microorganisms (fungi, bacteria), and insects, most researchers reported laccases from fungal sps., which mainly include those in genera ascomycetes, deuteromycetes, basidiomycetes, and cellulolytic fungi.^{52–56} Among these, basidiomycetes (white-rot fungi) laccases are most common and frequently reported and these are *Phlebia radiata*, *Trametes (Coriolus) versicolor*, *Coriolopsis polyzona*, *T. hirsuta*, *T. ochracea*, *T. villosa*, *Pycnoporus cinnabarinus*, *T. galli-ca*, *Lentinus edodes*, *Pleurotus ostreatus*, *Coprinus cinereus*, *etc.*⁵² *Monocillium indicum* was the first characterized ascomycete enzyme.^{57,58} Laccases, isolated from fungi, are mainly responsible for fructification, detoxification, phytopathogenic, sporulation, and degradation of lignin (Table 2).

Laccases isolated from bacteria have been reported to be the most stable laccases due to their resistance to high temperatures and wide pH ranges. While fungal laccases can be intra and extra-cellular, bacterial laccases are mostly intracellular, *i.e.*, *Marinomonas mediterranea*, *Azospirillum lipoferum*, and *Bacillus subtilis*.^{59,60} Bacterial laccases normally show the best activity at a higher pH compared to laccases from fungal sources,⁶¹ with the optimum pH being acidic for the latter.⁶² Plant laccases have intracellular physiological properties in the neutral pH range.^{63,64} Plant laccases have a higher isoelectric point compared with fungal laccases. Most laccases have an optimum temperature between 50–70 °C.⁶⁵ The thermal stability mainly depends on the type of microorganism; laccases from bacteria are more stable than those from fungi.⁶⁶ Due to the higher glycosylated content in the plant enzymes, they have higher molecular weight when compared with bacterial and fungal laccases. Laccases are secondary metabolites formed

under conditions of limited growth, mainly nitrogen, which negatively impacts the enzyme yield.⁶⁷ The production of laccases is mainly achieved through the fermentation process (liquid and solid).⁶⁸ Copper,⁶⁹ ethanol,⁷⁰ and aromatic compounds such as xyloidine, anisidine, syringaldazine, gallic acid, veratrine alcohol, ferulic acid, and lignosulfonates⁷¹ are used to induce the production of laccases.

5.1 Biochemical property characterization

Characterization parameters, like optimum temperature and pH, always play a significant role in the degradation process by any enzyme. Previously, many researchers have reported the role of characterization parameters in the laccase enzyme. Laccase (molecular weight (MW) 52 kDa) from *Cyberlindnera fabianii* has been reported for the degradation of 42.7% of Calcium alginate immobilized laccase (AIL) and 39.1% (Cu-AIL) of bisphenol A (BPA) (100 µM) after 24 h. The characterization of the native enzyme has shown an optimum pH and temperature of 5 and 40 °C, respectively.⁷² Likewise, laccase (MW 37 kDa) from *Bacillus subtilis* MTCC 2414 has been reported to degrade 81.72% of yellow GR dye after 120 h at 35 °C and 9 pH.⁷³ *Aspergillus oryzae* derived laccase (MM 57 kDa) with the potential of removing more than 80% of sulfamethoxazole, carbamazepine (CBZ), diclofenac, and bisphenol A at 30 °C and 7 pH,⁷⁴ is also reported to biodegrade 75% of azo-dyes after consecutive cycles at pH 5 at room temperature.⁷⁵ *Trametes versicolor* derived laccase also works as a biosensor for detecting phenolic compound catechol. Additionally, laccase (MW 31 kDa) isolated from *Bacillus atrophaeus* has been stated to clarify 41–58% phenolic compounds by reducing the color turbidity in the juices at 35 °C and 5.5 pH.⁴⁹

Biochemically, the protein is extremely thermostable with a half-life of about 2 h at 80 °C.⁷⁶ In contrast to fungal laccases, bacterial-derived laccases are biochemically more active at high pH values and highly stable at high temperatures. A high alkaline pH optimum of 8.5 to 9 was shown for several *Streptomyces* laccases for 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulfonic acid) (ABTS),^{77,78} whereas an optimum temperature of 85 °C was reported for the multicopper oxidase protein (McoP) laccase derived from *Pyrobaculum aerophilum*.⁷⁹ Furthermore, for *T. thermophilus* laccase, optimum temperature is 92 °C.⁸⁰ Fascinatingly, the laccase retrieved from drained soil with industrial wastewater is biochemically stable in the optimum pH range of 3 to 10.6.⁸¹ Another equally important biochemical parameter is salt tolerance. *Streptomyces ipomoea* derived laccase holds 100% activity in 1 M sodium chloride (NaCl) at an optimum pH of 8.0.⁷ These astonishing biochemical features highlight laccases as a potential basis for robust compounds with valuable biotechnological applications.

5.2 Mechanism

The biocatalytic processes of laccases have the potential application in the degradation of emerging compounds without any harmful effects; therefore, it increases the interest of the researchers to understand their mechanisms of degradation. Normally, laccases are reported for the oxidation of various ECs



Table 2 Laccase for the degradation of emerging contaminants

Source	Emerging contaminant	Biodegradation rate (%)	Reference
<i>A. densiflorus</i>	Rubin GFL (RGFL) dye	Rubin GFL (RGFL) dye (40 g L ⁻¹) up to 91% within 48 h	59
<i>Tagetes patula</i> , <i>Aster amellus</i> , <i>Portulaca grandiflora</i> and <i>Gaillardia grandiflora</i>	Textile effluent	Reduction in dyes 59, 50, 46 and 73%	60
<i>Glandularia pulchella</i>	Reactive orange HE2R, reactive yellow MEG4, reactive yellow GR, blue 2GL, remazol red, green HE4B, brown 3REL, blue 2RNL, patent blue, and malachite green	Decolorization of all the dyes	61
<i>Brumea malcolmii</i>	Brilliant blue R (BBR), malachite green, reactive red 2, direct red 5B and methyl orange	Compete degradation of dyes	62
<i>Alternanthera philoxeroides</i>	Sulfonated remazol red	Completely decolorize remazol red dye	63
<i>Arabidopsis thaliana</i>	Detoxification of herbicides atrazine (ATR) and isoproturon (IPU)	Detoxification or degradation of ATR or IPU	64
<i>Pycnoporus sanguineus</i> CCT-4518	17-Alpha-ethinylestradiol (EE2)	80% of removal of EE2 after 24 h	65
<i>Pycnoporus sanguineus</i>	Estrogens	96% of estrogens after 8 h of reaction	66
<i>Trametes trogii</i>	Amaranth, carmoisine, cochineal red, sunset yellow, patented blue, blue indigo and alizarin red S	All the dyes were decolorized up to 60% percent after 2 h	67
<i>Myceliophthora thermophila</i>	Anthraquinonic dyes acid blue 25 and acid green	RBBR and the diazo RB-5 were only decolorized with laccase/HBT, 31 and 60%, respectively, after 24 h	68
<i>Trametes versicolor</i>	Triphenylmethane dyes	Complete removal	69
<i>Trametes trogii</i> BAFC 463	Indigo carmine, xylidine, malachite green, gentian violet, bromophenol blue	Decolorized 85% of all dyes	70
<i>Trametes trogii</i>	Azoic, indigoid, triarylmethane, and anthraquinonic with acetosyringone	50–100% decolorization	71
<i>Trametes versicolor</i>	Bromocresol purple, safranin, malachite green, kristal violet, bromothymol blue, nigrosine and phenol red	43%, 54%, 55%, 49%, 56%, 53% and 37% for bromocresol purple, safranin, malachite green, kristal violet, Bromothymol blue, nigrosine and phenol red, respectively	72
<i>P. pastoris</i>	Crystal violet	Decolorization rates 90.7%	8
<i>Oudemansiella canarii</i>	Congo red	80% of 50 mg L ⁻¹ Congo red within 24 h	73
<i>Ganoderma lucidum</i> E47 strain	Decolorizing xanthene, azo and triarylmethane dyes	Decolorization	74
<i>Pycnoporus sanguineus</i> (CS43)	Bisphenol A	100% degradation of bisphenol A (20 mg L ⁻¹) was achieved in less than 24 h	16
<i>Trametes versicolor</i> BAFC 2234	Phenol	84% phenol removal in 4 h	75
<i>Trametes sanguineus</i>	Phenanthrene and benzo[α]pyrene	Removed phenanthrene and benzo[α]pyrene (97 and 99% respectively)	69
<i>Pleurotus ostreatus</i>	Phenol	70% phenol removal	76
<i>T. versicolor</i>	Naphthalene, phenanthrene	100% degradation	77
<i>P. ostreatus</i>	Atrazine	—	78
<i>Pseudomonas putida</i>	Chlorpyrifos	90% of degradation after 8 days of incubation	79
<i>P. pastoris</i>	Atrazine, isoproturon	Complete degradation	64
<i>T. versicolor</i>	Amoxicillin, Ampicillin, cloxacillin, penicillin G, penicillin V, oxacillin	54–100% after 24 h	80
<i>T. versicolor</i>	Sulfapyridine, sulfathiazole	100% after 8 h	81
<i>T. versicolor</i>	Sulfadimethoxine	>90% after 72 h	82
<i>Cerrena</i> sp. HYB07	Oxytetracycline, tetracycline	80% after 12 h	83
<i>A. oryzae</i>	Ciprofloxacin	51% after 5 h	84
<i>Ganoderma lucidum</i>	Triclosan (biocide)	90% after 24 h	85



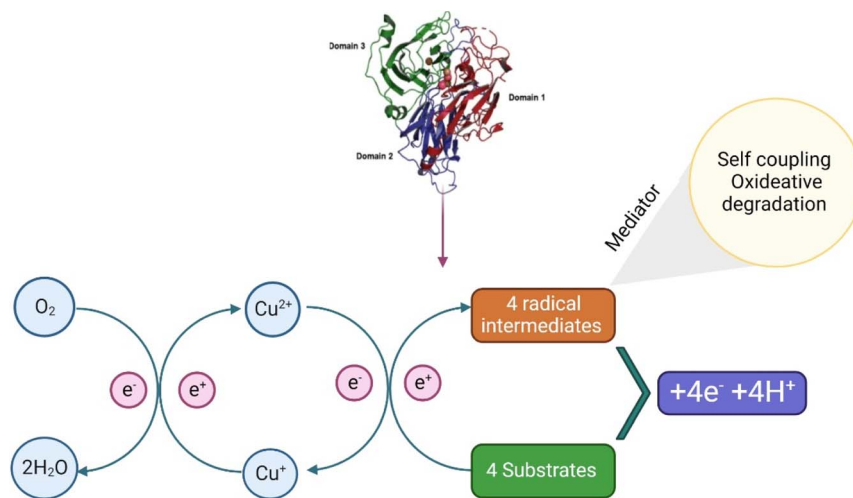


Fig. 3 Pathway of the laccase enzyme.

which mainly include phenols and aromatic amines; after degradation, they are directly transformed into free radicals (Fig. 3).^{82,83} These free radicals are commonly known to initiate domino reactions, primarily driving the complex transformations of ECs through biological processes, such as degradation by laccase and lignin synthesis.⁸⁴ Moreover, the laccase reaction involves one electron ($1e^-$), oxidations consecutive of four molecules of plummeting substrates, concurrently with doubling electron reduction ($2 \times 2e^-$) for oxygen atoms into H_2O molecules of their respective mechanisms. This method is accompanied by a catalytic exchange of $4H^+$ between parallel molecules.^{85,86} The laccase reaction involves two half-reactions associated with an internal electron transfer (IET) step and is supported by catalytic copper ions located at the T_1 copper (Cu) site and T_2 Cu/ T_3 Cu α / T_3 Cu β trinuclear cluster (TNC) sites from the mechanistic, kinetical, and structural points of view.^{85,87–89} The complete conservation of the eleven residues (one Cys and ten His) forming the T_1 copper and TNC laccase sites explains their essential role in the catalytic action. In many studies, comparisons related to sequences and mutagenic approaches have not been clearly demonstrated.^{83–85,90,91} Furthermore, highly conserved residues play important roles in various catalytic steps involved in laccase action.⁹² Despite these advances in the understanding of the action of laccases in terms of structure–function, a complete picture relating their molecular properties and mechanisms with their kinetic performance remains unclear. In brief, how these structural elements are automatically linked to catalytic copper center function and how the kinetic performance of these elements is influenced are listed below.

In the first half semi-reaction, $1e^-$ substrate oxidation at the T_1 copper site takes place which is located in the substrate binding pocket at the bottom. In *T. versicolor* laccase, this substrate interaction region is delimited by several highly conserved hydrophobic residues, namely, phenol (Phe) 162, leucine (Leu) 164, Phe 265, Phe 332, and proline (Pro) 391, which is useful for the formation of a favourable environment for the docking of hydrophobic molecules and for the

degradation of aromatic compounds substrates by laccase. Moreover, the residue of fully conserved Asp 206 (near His 458 of the T_1 copper site), situated in the binding pocket of the substrate at the bottom, is useful for the substrate orientation and stabilization through interaction with O–H bonds at the catalytic copper site T_1 , by the participation of fully conserved His 458.⁹³ Finally, the last residue should be exposed using a solvent in the interface for the binding cavity of the substrate. Furthermore, Asp 206 acts as a useful mechanistic component using electron subtraction promotion and the transfer of substrate donor molecules from the T_1 copper ion ($Cu^{2+} \rightarrow Cu^{1+}$) with the His 458, direct interaction at T_1 copper site. Moreover, the high E^o' detected on this *T. versicolor* is related directly to the Phe 463 residue at an axial position in the center.⁹⁴

6. Immobilized laccase in biodegradation

In general, biodegradation is the result of advances in molecular biotechnology, with important advantages over traditional degradation or catalytic processes, including higher catalytic behavior, higher specificity, higher reaction rate under mild conditions of pH and temperature, lower energy consumption, and most importantly, biodegradability.⁹⁵ Basically, biodegradation of ECs can be achieved in three different ways: (1) whole-cell biodegradation, which involves using biological molecules as biocatalysts, (2) isolated enzyme biodegradation, which involves using isolated enzymes; and (3) isolated-immobilized enzyme biodegradation, which involves immobilizing enzymes into nanomaterials.⁹⁶ The first two methods have been widely used for the biodegradation of ECs due to important properties, including high reaction rates and relatively low cost and operation time.⁹⁷ For example,⁹⁸ a high biodegradation rate (96%) for sertraline, clomipramine, mianserin, and paroxetine has been reported using *Pleurotus ostreatus* as a biocatalyst. However, both techniques might present problems, such as potential cross-reactivity, side reactions, and poor or null



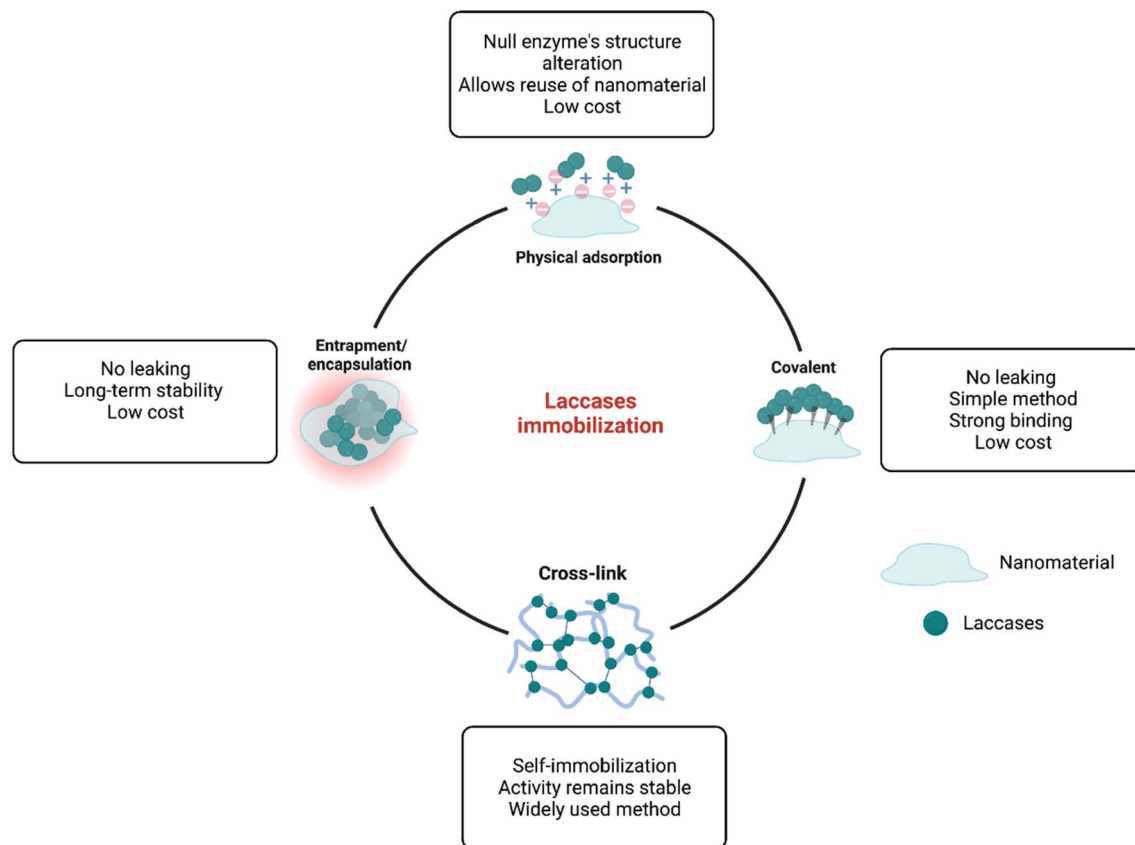


Fig. 4 Routes for laccase immobilization into nanomaterials.

reusability.^{96,99} On the other hand, the immobilization of enzymes into suitable nanomaterials might provide an optimal environment to reach high biodegradation rates. Several important properties are enhanced during this process, *i.e.*, higher thermal and pH stability, and improved storage stability, long-term operation, reusability, and higher enzymatic activity.¹⁰⁰ Different enzymes have been reported as immobilized enzymes for the biodegradation of ECs, including peroxidases, oxidases, catalase, super oxidase dismutase, and laccases.¹⁰¹ Laccases have gained significant interest in developing novel immobilized materials for their application in the biodegradation of emerging pollutants since they can catalyze the biotransformation of a wide range of compounds, such as phenols, dyes, pharmaceuticals, and many others.¹⁰² Different nanomaterials, each one with different properties, have been applied for the immobilization of laccases and its further application in the biodegradation of ECs, including polymers,¹⁰³ metals and metal oxides,⁹⁶ and carbon-based materials.¹⁰⁴ Thus, depending on the nanomaterial used and the procedure of immobilization, the characteristics and properties of the laccase immobilized system change.⁷³ The immobilization of laccases into nanomaterials can be achieved by different methods (Fig. 4), which are briefly described below:

- Physical adsorption immobilization is mainly characterized since it is the simplest immobilization method, which basically involves the interaction of the enzyme with the

nanomaterial by the formation of weak ionic bonds that usually are formed by the dissolution of the enzyme at controlled pH and the subsequent addition of the host matrix.⁹⁶ Important advantages of adsorption immobilization are low operational costs and times, reusability, and increased catalytic stability. However, since the enzyme–nanomaterial interaction is governed by weak binding forces, changes in pH, temperature, or ionic strength, might produce a leaching effect.^{97,105}

- The covalent immobilization technique is widely used due to its important advantages, which include strong bindings between enzymes and supports, high stability across varying pH and temperature conditions, high uniformity, and good control over the immobilized enzyme amount.¹⁰⁶ Covalent immobilization is a two-step process, in which first, the host carrier is activated by the addition of multifunctional reagents (*e.g.*, glutaraldehyde and carbodiimide), which then promotes the formation of covalent bonds with functional groups of the enzyme, most commonly amino, carboxylic, phenolic, and hydroxyl groups.⁹⁶ Even though covalent immobilization offers great advantages, due to the structural modification of the enzyme after the formation of the covalent bond, the active sites of the enzyme might be deactivated, thus affecting the catalytic properties.⁹²

- Cross-linking immobilization involves the self-immobilization of enzymes by the formation of cross-linkages named cross-linked enzyme aggregates (CLEAs), which are



formed by the addition of bifunctional reagents, such as glutaraldehyde, dialdehydes, diiminoesters, diisocyanates and diamines, and the subsequent formation of a three-dimensional complex structure through the interaction of amine groups of the enzyme and the activated bifunctional reagents.¹⁰⁷ This method has the ability to form materials resistant to the effects of pH and temperature, but a major disadvantage is the loss of enzymatic activity after immobilization.¹⁰⁶

• Entrapment and encapsulation are two similar methods involving a two-step process, in which the enzyme and a monomeric solution are mixed, and then polymerization of the monomer confines/entraps the enzyme within the polymeric network without chemical interactions.⁹⁶ These two methods are characterized for providing significant protection from the external environmental conditions to the enzyme, enhancing the stability and decreasing denaturation.⁷⁴ On the other hand, these methods have important mass transfer limitations, which means that substrates might have problems in reaching the active sites of the confined enzyme.^{70,97}

7. Laccase as a biocatalyst for biodegradation

Immobilized laccases have been widely implemented as biocatalysts in the biodegradation of different ECs (Table 3). Even though the application of laccases as biocatalysts might be achieved either through free or immobilized enzymatic systems, it is well known that through the implementation of immobilized laccase biocatalysts, it is possible to overcome the major disadvantages of using free enzymes, which are discussed above.¹¹ Different nanomaterials and immobilization methods

have been applied to design laccase-based biocatalysts for the biodegradation of different ECs. For example, adsorption immobilization was reported to be effective for the biodegradation of CBZ, bisphenol A (BPA), and synthetic dyes, by the immobilization of laccases into polymers,¹⁰⁸ carbon-based nanomaterials,¹⁰⁹ and metal-based nanomaterials.¹¹⁰ Moreover, covalent, entrapment, and cross-link immobilization have been implemented to design novel biocatalysts by using polymers, metal oxide, and carbon-based nanomaterials.¹¹⁵

Moreover, Ma *et al.* (2018) also described the biodegradation of dyes, including CV, Congo red (CR), indigo blue (IB), and others by the entrapment of laccases from *Trametes pubescens* into a polymeric chitosan matrix. In this case, biodegradation rates reached about 79%. However, the biodegradation of dyes by implementing this biocatalyst was achieved after 96 h, which might be mainly attributed to the characteristic mass transfer limitations achieved after entrapment immobilization.^{74,114}

The application of immobilized enzyme systems as biocatalysts for the biodegradation of ECs is an important approach to deal with the increasing environmental impact that ECs might have on human well-being and the environment.¹⁰ Thus, designing a novel and highly catalytic biocatalyst is a critical point as a prospect; a suitable biocatalyst might be designed in two main steps: (1) preparation of the host carrier material and (2) choosing a suitable immobilization method.⁹⁶

8. Conclusion and future prospects

In summary, laccases play a crucial role in breaking down ECs. Leveraging modern protein engineering tools makes the use of laccases viable. However, optimizing various factors such as pH, temperature, suspended solids, and mechanical stress is essential to maximize laccase yield. Extensive optimization is

Table 3 Application of isolated-immobilized laccases in the biodegradation of emerging contaminants^a

Laccase source	Support	Immobilization method	Emerging contaminant	Conditions	Biodegradation rate (%)	Reference
<i>Trametes versicolor</i>	Polyimide aerogels	Covalent	CBZ	20 ng mL ⁻¹ of CBZ, 200 rpm, 25 °C, 24 h	74–76	108
<i>Trametes hirsuta</i>	PVDF/MWCNTs	Adsorption	CBZ	5 ppm of CBZ, 223 U mL ⁻¹ of catalysts, pH 5, 25 °C, 4 h	95	109
<i>Trametes versicolor</i>	CNTs	Adsorption	Dyes	10 mg L ⁻¹ of dye, 0.08 g L ⁻¹ of catalyst, 35 °C, 3 h	96	111
<i>Trametes versicolor</i>	Fe ₃ O ₄ @C-Cu ²⁺	Adsorption	Dyes	5–200 mg L ⁻¹ of dye, pH 4.5, 50 °C, 1 h	75–99	112
<i>Trametes versicolor</i>	DAY	Adsorption	BPA	1 mL BPA 2 mM, pH 4.5, 5 mg of catalyst, 45 °C, 1 h	86.7	110
<i>Aspergillus flavus</i>	GO	Covalent	Azo dyes	300 mg L ⁻¹ of catalyst, 10 ppm of dye, pH 5, 45 °C, 0.2 h	48.7–88.7	76
<i>Trametes versicolor</i>	Copper alginate	Entrapment	2,4-DCP	10 mg mL ⁻¹ of 2,4-DCP, pH 4.6, 40 °C, 10 mh	96.4	113
<i>Trametes pubescens</i>	Chitosan	Entrapment	Dyes	50 mg mL ⁻¹ of dye, 10 g of catalyst, pH 5, 55 °C, 96 h	37–79	114
<i>Trametes villosa</i>	CLEAs	Cross-link	RIF, INH	5 mg mL ⁻¹ of drug, 100 U L ⁻¹ of catalyst, pH 5, 40 °C, 24 h	71–95	115

^a Carbamazepine (CBZ), polyvinylidene fluoride (PVDF), multiwalled carbon nanotubes (MWCNTs), bisphenol A (BPA), sodium zeolite Y dealuminated form (DAY), graphene oxide (GO), 2,4-dichlorophenol (2,4-DCP), ultraporous alumina (UPA), rifampicin (RIF), isoniazid (INH), cross-linked.



pivotal for generating substantial laccase biomass, which holds the potential for addressing the pressing challenge of wastewater treatment and purification in the commercial sector. Presently, safeguarding natural water bodies from pollutants stemming from ECs is an urgent concern. These pollutants, encompassing plastics, herbicides, fertilizers, dyes, phthalates, pharmaceuticals, and personal care products, find their way into water bodies through direct runoff and disposal. The degradation of these substances is paramount. Laccases emerge as efficient biocatalytic agents, adept at converting these compounds into less harmful and inert derivatives. It's worth noting that complexities like the intricate composition of contaminated water (high salt concentrations and/or elevated pH levels) pose obstacles to efficient degradation. Nonetheless, contemporary techniques like laccase engineering offer promising solutions. Encouraging reports have highlighted successful EC degradation. Yet, transitioning these processes to pilot-scale bioprocesses hinges on economic feasibility. Immobilizing laccases presents a viable strategy for treatment procedures. Nevertheless, the time-intensive nature of laccase extraction and purification remains a challenge. In this regard, producing recombinant laccases with increased activity and stability holds immense promise. This advancement could streamline the enzyme purification process, significantly enhancing cost-effectiveness.

Ethical approval

All data referenced in this review paper have been obtained from publicly available sources, with no use of confidential or sensitive information.

Data availability

No primary research results, software or code have been included and no new data were generated or analysed as part of this review.

Author contributions

Pooja Thathola: conceptualization, writing of the original draft, review & editing. Elda M. Melchor-Martínez: writing of the original draft, review & editing. Priyanka Adhikari: writing of the original draft, review & editing. Saúl Antonio Hernández Martínez: reviewing. Anita Pandey: conceptualization, reviewing. Roberto Parra-Saldivar: reviewing.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

The authors are grateful to their representative universities and institutes for providing literature facilities.

References

- 1 B. S. Rath, P. S. Kumar and P. L. Show, *J. Hazard. Mater.*, 2021, **409**, 124413.
- 2 V. Uddameri, A. Ghaseminejad and E. A. Hernandez, *Agric. Water Manag.*, 2020, **238**, 106226.
- 3 D. Saidulu, B. Gupta, A. K. Gupta and P. S. Ghosal, *J. Environ. Chem. Eng.*, 2021, **9**, 105282.
- 4 Y. H. Pang, Y. Y. Huang, L. Wang, X. F. Shen and Y. Y. Wang, *Environ. Pollut.*, 2020, **263**, 114616.
- 5 S. Kashefi, S. M. Borghei and N. M. Mahmoodi, *J. Mol. Liq.*, 2019, **276**, 153–162.
- 6 P. Wu, C. Zhou, Y. Li, M. Zhang, P. Tao, Q. Liu and W. Cui, *Appl. Surf. Sci.*, 2021, **540**, 148362.
- 7 V. Cristino, L. Pasti, N. Marchetti, S. Berardi, C. A. Bignozzi, A. Molinari, F. Passabi, S. Caramori, L. Amidani, M. Orlandi, N. Bazzanella, A. Piccioni, J. Kopula Kesavan, F. Boscherini and L. Pasquini, *Photochem. Photobiol. Sci.*, 2019, **18**, 2150–2163.
- 8 J. Wang and H. Chen, *Sci. Total Environ.*, 2020, **704**, 135249.
- 9 R. Bai, W. Yan, Y. Xiao, S. Wang, X. Tian, J. Li, X. Xiao, X. Lu and F. Zhao, *Chem. Eng. J.*, 2020, **397**, 125501.
- 10 R. Morsi, K. A. Al-Maqdi, M. Bilal, H. M. N. Iqbal, A. Khaleel, I. Shah and S. S. Ashraf, *Biomolecules*, 2021, **11**(6), 904.
- 11 R. A. Sheldon and J. M. Woodley, *Chem. Rev.*, 2018, **118**, 801–838.
- 12 D. Wan, H. Wang, I. P. Pozdnyakov, C. Wang, J. Su, Y. Zhang, Y. Zuo, D. D. Dionysiou and Y. Chen, *Water Res.*, 2020, **184**, 116002.
- 13 J. García, M. J. García-Galán, J. W. Day, R. Boopathy, J. R. White, S. Wallace and R. G. Hunter, *Bioresour. Technol.*, 2020, **307**, 123228.
- 14 N. H. Tran and K. Y. H. Gin, *Sci. Total Environ.*, 2017, **599–600**, 1503–1516.
- 15 M. Patel, R. Kumar, K. Kishor, T. Mlsna, C. U. Pittman and D. Mohan, *Chem. Rev.*, 2019, **119**, 3510–3673.
- 16 C. Barrios-Estrada, M. de J. Rostro-Alanis, A. L. Parra, M. P. Belleville, J. Sanchez-Marcano, H. M. N. Iqbal and R. Parra-Saldivar, *Int. J. Biol. Macromol.*, 2018, **108**, 837–844.
- 17 C. A. Lutterbeck, G. S. Colares, N. Dell'Osbelt, F. P. da Silva, L. T. Kist and E. L. Machado, *J. Cleaner Prod.*, 2020, **273**, 122851.
- 18 Y. Chen, J. Vymazal, T. Březinová, M. Koželuh, L. Kule, J. Huang and Z. Chen, *Sci. Total Environ.*, 2016, **566–567**, 1660–1669.
- 19 S. M. Mousavi, S. A. Hashemi, S. M. Iman Moezzi, N. Ravan, A. Gholami, C. W. Lai, W. H. Chiang, N. Omidifar, K. Yousefi and G. Behbudi, *Biochem. Res. Int.*, 2021, 5599204.
- 20 T. D. H. Bugg, J. J. Williamson and G. M. M. Rashid, *Curr. Opin. Chem. Biol.*, 2020, **55**, 26–33.
- 21 M. Bilal, M. Adeel, T. Rasheed, Y. Zhao and H. M. N. Iqbal, *Environ. Int.*, 2019, **124**, 336–353.
- 22 L. Arregui, M. Ayala, X. Gómez-Gil, G. Gutiérrez-Soto, C. E. Hernández-Luna, M. Herrera De Los Santos, L. Levin, A. Rojo-Domínguez, D. Romero-Martínez,



- M. C. N. Saparrat, M. A. Trujillo-Roldán and N. A. Valdez-Cruz, *Microb. Cell Factories*, 2019, **18**(1), 1–33.
- 23 K. Balakrishna, A. Rath, Y. Praveenkumarreddy, K. S. Guruge and B. Subedi, *Ecotoxicol. Environ. Saf.*, 2017, **137**, 113–120.
- 24 J. M. Brausch and G. M. Rand, *Chemosphere*, 2011, **82**, 1518–1532.
- 25 H. Wang, H. Xi, L. Xu, M. Jin, W. Zhao and H. Liu, *Sci. Total Environ.*, 2021, **788**, 147819.
- 26 F. Freeling, N. A. Alygizakis, P. C. von der Ohe, J. Slobodnik, P. Oswald, R. Aalizadeh, L. Cirka, N. S. Thomaidis and M. Scheurer, *Sci. Total Environ.*, 2019, **681**, 475–487.
- 27 M. E. Scheurer, M. L. Bondy, K. D. Aldape, T. Albrecht and R. El-Zein, *Acta Neuropathol.*, 2008, **116**, 79.
- 28 J. M. Dabrowski, *S. Afr. J. Sci.*, 2015, **111**, 1–7.
- 29 H. N. Bischel, B. D. Özel Duygan, L. Strande, C. S. Mc Ardell, K. M. Udert and T. Kohn, *Water Res.*, 2015, **85**, 57–65.
- 30 S. A. Madhi, C. Cutland, K. Ismail, C. O'Reilly, A. Mancha and K. P. Klugman, *Clin. Infect. Dis.*, 2002, **35**, 1120–1126.
- 31 K. Narasimhulu, *Innovations in Microbiology and Biotechnology*, 2022, vol. 5, pp. 70–81.
- 32 A. P. Daso, O. S. Fatoki, J. P. Odendaal and O. O. Olujimi, *Environ. Monit. Assess.*, 2013, **185**, 431–439.
- 33 O. I. Olukunle, I. v. Sibiya, O. J. Okonkwo and A. O. Odusanya, *Environ. Sci. Pollut. Res. Int.*, 2015, **22**, 2145–2154.
- 34 M. H. Wu, C. C. Lee, A. S. Hsiao, S. M. Yu, A. H. J. Wang and T. H. D. Ho, *FEBS Open Bio*, 2018, **8**, 1230–1246.
- 35 A. M. Deegan, B. Shaik, K. Nolan, K. Urell, M. Oelgemöller, J. Tobin and A. Morrissey, *Int. J. Environ. Sci. Technol.*, 2011, **8**(3), 649–666.
- 36 S. Mohapatra, C. H. Huang, S. Mukherji and L. P. Padhye, *Chemosphere*, 2016, **159**, 526–535.
- 37 A. Agüera, M. J. Martínez Bueno and A. R. Fernández-Alba, *Environ. Sci. Pollut. Res.*, 2013, **20**(6), 3496–3515.
- 38 M. C. V. M. Starling, C. C. Amorim and M. M. D. Leão, *J. Hazard. Mater.*, 2019, **372**, 17–36.
- 39 T. K. Kasonga, M. A. A. Coetzee, I. Kamika, V. M. Ngole-Jeme and M. N. Benteke Momba, *J. Environ. Manage.*, 2021, **277**, 111485.
- 40 P. Chaturvedi, P. Shukla, B. S. Giri, P. Chowdhary, R. Chandra, P. Gupta and A. Pandey, *Environ. Res.*, 2021, **194**, 110664.
- 41 Y. Atih, M. AlBashtawy, A. Alkhawaldeh, B. Albashtawy, S. Albashtawy, A. Ibnian, Z. Albashtawy, A. Kazaleh, A. Ayed, M. Suliman, K. al Dameery, E. Rn and F. Jordanian, *J. Hospital*, 2017, **1**, 7.
- 42 H. Ramírez-Malule, D. H. Quiñones-Murillo and D. Manotas-Duque, *Emerging Contam.*, 2020, **6**, 179–193.
- 43 S. Zahmatkesh, A. Bokhari, M. Karimian, M. M. A. Zahra, M. Sillanpää, H. Panchal, A. J. Alrubaie and Y. Rezakhani, *Environ. Monit. Assess.*, 2022, **194**(12), 1–15.
- 44 R. R. Nair, P. Demarche and S. N. Agathos, *New Biotechnol.*, 2013, **30**, 814–823.
- 45 K. Agrawal, V. Chaturvedi and P. Verma, *Bioresour. Bioprocess*, 2018, **5**, 1–12.
- 46 J. Zdarta, K. Anteck, R. Frankowski, A. Zgoła-Grześkowiak, H. Ehrlich and T. Jesionowski, *Sci. Total Environ.*, 2018, **615**, 784–795.
- 47 M. Bilal, T. Rasheed, F. Nabeel, H. M. N. Iqbal and Y. Zhao, *J. Environ. Manage.*, 2019, **234**, 253–264.
- 48 D. M. Mate and M. Alcalde, *Microb. Biotechnol.*, 2017, **10**, 1457–1467.
- 49 L. K. Narnoliya, N. Agarwal, S. N. Patel and S. P. Singh, *J. Microbiol.*, 2019, **57**(10), 900–909.
- 50 F. Lassouane, H. Ait-Amar, S. Amrani and S. Rodriguez-Couto, *Bioresour. Technol.*, 2019, **271**, 360–367.
- 51 H. Yoshida, *J. Chem. Soc. Trans.*, 1883, (43), 472–486.
- 52 P. Baldrian, *FEMS Microbiol. Rev.*, 2006, **30**, 215–242.
- 53 P. Schneider, M. B. Caspersen, K. Mondorf, T. Halkier, L. K. Skov, P. R. Østergaard, K. M. Brown, S. H. Brown and F. Xu, *Enzyme Microb. Technol.*, 1999, **25**, 502–508.
- 54 P. Sharma, R. Goel and N. Capalash, *World J. Microbiol. Biotechnol.*, 2006, **23**(6), 823–832.
- 55 A. Pandey, G. Szakacs, C. R. Soccol, J. A. Rodriguez-Leon and V. T. Soccol, *Bioresour. Technol.*, 2001, **77**, 203–214.
- 56 A. Hatakka, *FEMS Microbiol. Rev.*, 1994, **13**, 125–135.
- 57 H. Patel, V. K. Yadav, K. K. Yadav, N. Choudhary, H. Kalasariya, M. M. Alam, A. Gacem, M. Amanullah, H. A. Ibrahim, J. W. Park, S. Park and B. H. Jeon, *Water*, 2022, **14**(19), 3163.
- 58 J. M. Khaled, S. A. Alyahya, R. Govindan, C. K. Chelliah, M. Maruthupandy, N. S. Alharbi, S. Kadaikunnan, R. Issac, S. Murugan and W. J. Li, *Environ. Res.*, 2022, **207**, 112211.
- 59 O. v. Morozova, G. P. Shumakovich, S. V. Shleev and Y. I. Yaropolov, *Appl. Biochem. Microbiol.*, 2007, **43**(5), 523–535.
- 60 F. Rosconi, L. F. Fraguas, G. Martínez-Drets and S. Castro-Sowinski, *Enzyme Microb. Technol.*, 2005, **36**, 800–807.
- 61 J. Margot, C. Bennati-Granier, J. Maillard, P. Blánquez, D. A. Barry and C. Holliger, *AMB Express*, 2013, **3**, 1–14.
- 62 K. Dhakar and A. Pandey, *Enzym. Res.*, 2013, 869062.
- 63 C. Pezzella, V. Lettera, A. Piscitelli, P. Giardina and G. Sannia, *Appl. Microbiol. Biotechnol.*, 2012, **97**(2), 705–717.
- 64 U. N. Dwivedi, P. Singh, V. P. Pandey and A. Kumar, *J. Mol. Catal. B Enzym.*, 2011, **68**, 117–128.
- 65 B. Chefetz, Y. Chen and Y. Hadar, *Appl. Environ. Microbiol.*, 1998, **64**, 3175–3179.
- 66 K. Hildén, T. K. Hakala and T. Lundell, *Biotechnol. Lett.*, 2009, **31**(8), 1117–1128.
- 67 L. Gianfreda, F. Xu and J. M. Bollag, *Bioremediat. J.*, 2010, **3**, 1–26.
- 68 S. Mazumder, S. K. Basu and M. Mukherjee, *Eng. Life Sci.*, 2009, **9**, 45–52.
- 69 G. Palmieri, P. Giardina, C. Bianco, B. Fontanella and G. Sannia, *Appl. Environ. Microbiol.*, 2000, **66**, 920–924.
- 70 B. Bertrand, F. Martínez-Morales, R. Tinoco, S. Rojas-Trejo, L. Serrano-Carreón and M. R. Trejo-Hernández, *World J. Microbiol. Biotechnol.*, 2013, **30**(1), 135–142.
- 71 J. A. Saraiva, A. P. M. Tavares and A. M. R. B. Xavier, *Appl. Biochem. Biotechnol.*, 2012, **167**(4), 685–693.



- 72 F. M. Olajuyigbe, O. Y. Adetuyi and C. O. Fatokun, *Int. J. Biol. Macromol.*, 2019, **125**, 856–864.
- 73 L. N. Nguyen, F. I. Hai, A. Dosseto, C. Richardson, W. E. Price and L. D. Nghiem, *Bioresour. Technol.*, 2016, **210**, 108–116.
- 74 H. H. Nguyen and M. Kim, *Appl. Sci. Conver. Technol.*, 2017, **26**, 157–163.
- 75 S. Kashefi, S. M. Borghei and N. M. Mahmoodi, *J. Mol. Liq.*, 2019, **276**, 153–162.
- 76 M. F. Hullo, I. Moszer, A. Danchin and I. Martin-Verstraete, *J. Bacteriol.*, 2001, **183**, 5426.
- 77 M. C. Machczynski, E. Vijgenboom, B. Samyn and G. W. Canters, *Protein Sci.*, 2004, **13**, 2388–2397.
- 78 K. N. Niladevi, P. S. Sheejadevi and P. Prema, *Appl. Biochem. Biotechnol.*, 2008, **151**, 9–19.
- 79 A. T. Fernandes, J. M. Damas, S. Todorovic, R. Huber, M. C. Baratto, R. Pogni, C. M. Soares and L. O. Martins, *FEBS J.*, 2010, **277**, 3176–3189.
- 80 K. Miyazaki, *Extremophiles*, 2005, **9**, 415–425.
- 81 J. Bains, N. Capalash and P. Sharma, *Biotechnol. Lett.*, 2003, **25**, 1155–1159.
- 82 T. Kupper, C. Plagellat, R. C. Brändli, L. F. de Alencastro, D. Grandjean and J. Tarradellas, *Water Res.*, 2006, **40**, 2603–2612.
- 83 A. Wick, O. Marincas, Z. Moldovan and T. A. Ternes, *Water Res.*, 2011, **45**, 3638–3652.
- 84 S. Comtois-Marotte, T. Chappuis, S. Vo Duy, N. Gilbert, A. Lajeunesse, S. Taktek, M. Desrosiers, É. Veilleux and S. Sauvé, *Chemosphere*, 2017, **166**, 400–411.
- 85 A. Palardy, J. P. Gagné and L. Tremblay, *J. Xenobiot*, 2016, **5**, 37–39.
- 86 S. N. Grove and C. E. Bracker, *J. Bacteriol.*, 1970, **104**, 989–1009.
- 87 H. Kojima, F. Sata, S. Takeuchi, T. Sueyoshi and T. Nagai, *Toxicology*, 2011, **280**, 77–87.
- 88 S. Foteinis, A. G. L. Borthwick, Z. Frontistis, D. Mantzavinos and E. Chatzisyneon, *J. Cleaner Prod.*, 2018, **182**, 8–15.
- 89 J. A. Rivera-Jaimes, C. Postigo, R. M. Melgoza-Alemán, J. Aceña, D. Barceló and M. López de Alda, *Sci. Total Environ.*, 2018, **613–614**, 1263–1274.
- 90 L. You, V. T. Nguyen, A. Pal, H. Chen, Y. He, M. Reinhard and K. Y. H. Gin, *Sci. Total Environ.*, 2015, **536**, 955–963.
- 91 L. You, V. T. Nguyen, A. Pal, H. Chen, Y. He, M. Reinhard and K. Y. H. Gin, *Sci. Total Environ.*, 2015, **536**, 955–963.
- 92 S. Zhou, C. Di Paolo, X. Wu, Y. Shao, T. B. Seiler and H. Hollert, *Environ. Int.*, 2019, **128**, 1–10.
- 93 T. Tsuda, T. Igawa, K. Tanaka and D. Hirota, *Bull. Environ. Contam. Toxicol.*, 2011, **87**(3), 307–311.
- 94 A. S. Mansano, R. A. Moreira, M. Pierozzi, T. M. A. Oliveira, E. M. Vieira, O. Rocha and M. H. Regali-Selegim, *Environ. Pollut.*, 2016, **213**, 160–172.
- 95 E. L. Bell, W. Finnigan, S. P. France, A. P. Green, M. A. Hayes, L. J. Hepworth, S. L. Lovelock, H. Niikura, S. Osuna, E. Romero, K. S. Ryan, N. J. Turner and S. L. Flitsch, *Nat. Rev. Methods Primers*, 2021, **1**(1), 1–21.
- 96 S. A. H. Martínez, E. M. Melchor-Martínez, J. A. R. Hernández, R. Parra-Saldívar and H. M. N. Iqbal, *Fuel*, 2022, **312**, 122927.
- 97 D. M. Liu, J. Chen and Y. P. Shi, *TrAC, Trends Anal. Chem.*, 2018, **102**, 332–342.
- 98 Y. Chen, J. Vymazal, T. Březinová, M. Koželuh, L. Kule, J. Huang and Z. Chen, *Sci. Total Environ.*, 2016, **566–567**, 1660–1669.
- 99 M. Bilal and H. M. N. Iqbal, *Coord. Chem. Rev.*, 2019, **388**, 1–23.
- 100 F. Darvishi, M. Moradi, C. Jolivald and C. Madzak, *Ecotoxicol. Environ. Saf.*, 2018, **165**, 278–283.
- 101 D. O. Lopez-Cantu, R. B. González-González, E. M. Melchor-Martínez, S. A. H. Martínez, R. G. Araújo, L. Parra-Arroyo, J. E. Sosa-Hernández, R. Parra-Saldívar and H. M. N. Iqbal, *Int. J. Biol. Macromol.*, 2022, **194**, 676–687.
- 102 L. Alvarado-Ramírez, M. Rostro-Alanis, J. Rodríguez-Rodríguez, C. Castillo-Zacarias, J. E. Sosa-Hernández, D. Barceló, H. M. N. Iqbal and R. Parra-Saldívar, *Int. J. Biol. Macromol.*, 2021, **181**, 683–696.
- 103 S. Aslam, M. Asgher, N. A. Khan and M. Bilal, *J. Water Proc. Eng.*, 2021, **40**, 101971.
- 104 H. M. Xu, X. F. Sun, S. Y. Wang, C. Song and S. G. Wang, *Sep. Purif. Technol.*, 2018, **204**, 255–260.
- 105 W. Zhou, W. Zhang and Y. Cai, *Chem. Eng. J.*, 2021, **403**, 126272.
- 106 X. Lyu, R. Gonzalez, A. Horton and T. Li, *Catalysts*, 2021, **11**, 1211.
- 107 D. M. Liu, J. Chen and Y. P. Shi, *Trends Anal. Chem.*, 2018, **102**, 332–342.
- 108 C. Simón-Herrero, M. Naghdi, M. Taheran, S. Kaur Brar, A. Romero, J. L. Valverde, A. Avalos Ramirez and L. Sánchez-Silva, *J. Hazard. Mater.*, 2019, **376**, 83–90.
- 109 M. Masjoudi, M. Golgoli, Z. Ghobadi Nejad, S. Sadeghzadeh and S. M. Borghei, *Chemosphere*, 2021, **263**, 128043.
- 110 T. Taghizadeh, A. Talebian-Kiakalaieh, H. Jahandar, M. Amin, S. Tarighi and M. A. Faramarzi, *J. Hazard. Mater.*, 2020, **386**, 121950.
- 111 S. Zhang, Z. Wu, G. Chen and Z. Wang, *Catalysts*, 2018, **8**, 286.
- 112 H. F. Ma, G. Meng, B. K. Cui, J. Si and Y. C. Dai, *Chem. Eng. Res. Des.*, 2018, **132**, 664–676.
- 113 S. Zhang, Z. Wu, G. Chen and Z. Wang, *Catalysts*, 2018, **8**(7), 286.
- 114 Z. Li, Z. Chen, Q. Zhu, J. Song, S. Li and X. Liu, *J. Hazard. Mater.*, 2020, **399**, 123088.
- 115 H. de P. Riedi, M. V. de Liz, D. M. Braga, A. B. Ianoski, T. de Freitas Pereira, T. Brugnari, C. W. I. Haminiuk and G. M. Maciel, *Int. J. Environ. Res.*, 2022, **16**(3), 1–12.

