

Review

Color Stains on Paper: Fungal Pigments, Synthetic Dyes and Their Hypothetical Removal by Enzymatic Approaches

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Abstract: Fungi are the main contaminants of books and archival documents. In addition to their degrading power, offered by various types of lignolytic and cellulolytic enzymes, they can also hue the surface of the paper through the production of pigments. The fungi on paper release various types of pigments belonging mostly to two chemical groups (polyketides and carotenoids), which cause unpleasant anaesthetic stains. The paper surface can also be hued with several synthetic colors, which are part, for example, of stamps and inks. These synthetic colors could be degraded by lignin-modifying enzymes (LMEs) and also by dye-decolorizing peroxidases (DyPs). Therefore, the mechanism of action of LEMs and DyPs is illustrated. Moreover, we have examined the potentiality of LEMs and DyPs to remove the synthetic stains and also their hypothetical application in order to clean the fungal hues from the paper surface. Our review article, using the enzymatic removal parallelism between fungal and synthetic pigments, would like to show prospective solutions to this arduous problem.

Keywords: fungal stains; ink stamp; lignin-modifying enzymes; dye-decolorizing peroxidases; bio-cleaning



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1. Introduction

Many books, historical documents, sketches, paintings, music scores, architectural construction documents, postage stamps, and photographs are susceptible to destruction [1–5]. This destruction can be a result of the influence of endogenous factors such as acidity, metal ions, lignin, or paper degradation products, and exogenous factors such as heat, humidity, radiation (light, UV), oxygen, pollutants, or biodeteriogens [6,7].

Filamentous fungi have been considered to be the most important microorganisms responsible for the biodeterioration of collections in museums, archives, galleries, and libraries [6,8,9], causing severe material and information losses in libraries' and archives' collections [8].

Fungi possess the capacity to grow at average relative humidity levels, inducing several chemical and physical decomposing processes through the excretion of metabolites and enzymes that affect the structural organization of the paper [2,10]. These metabolites comprise a wide variety of substances such as colorants, enzymes, acids, chelating agents, etc. [11]. However, one of the major concerns regarding fungal colonization of paper is that it undergoes degradation that affects its mechanical and aesthetic properties [12] since the stains excreted by fungi are colored [13] (Figure 1). Several reviews and research papers can be found describing fungi, their colorants, and the foxing phenomenon causing biodeterioration of historical paper (Figure 1) [2,14–17], but very little data is presented about the acceptable methodology of how to treat and remove the stains from paper. To address these demands, first, it is important to collect available information about the fungal species that stain paper and the chemical nature of those stains. Another important demand is to develop new cleaning methods that would not bleach and destroy the aesthetics of the paper itself.

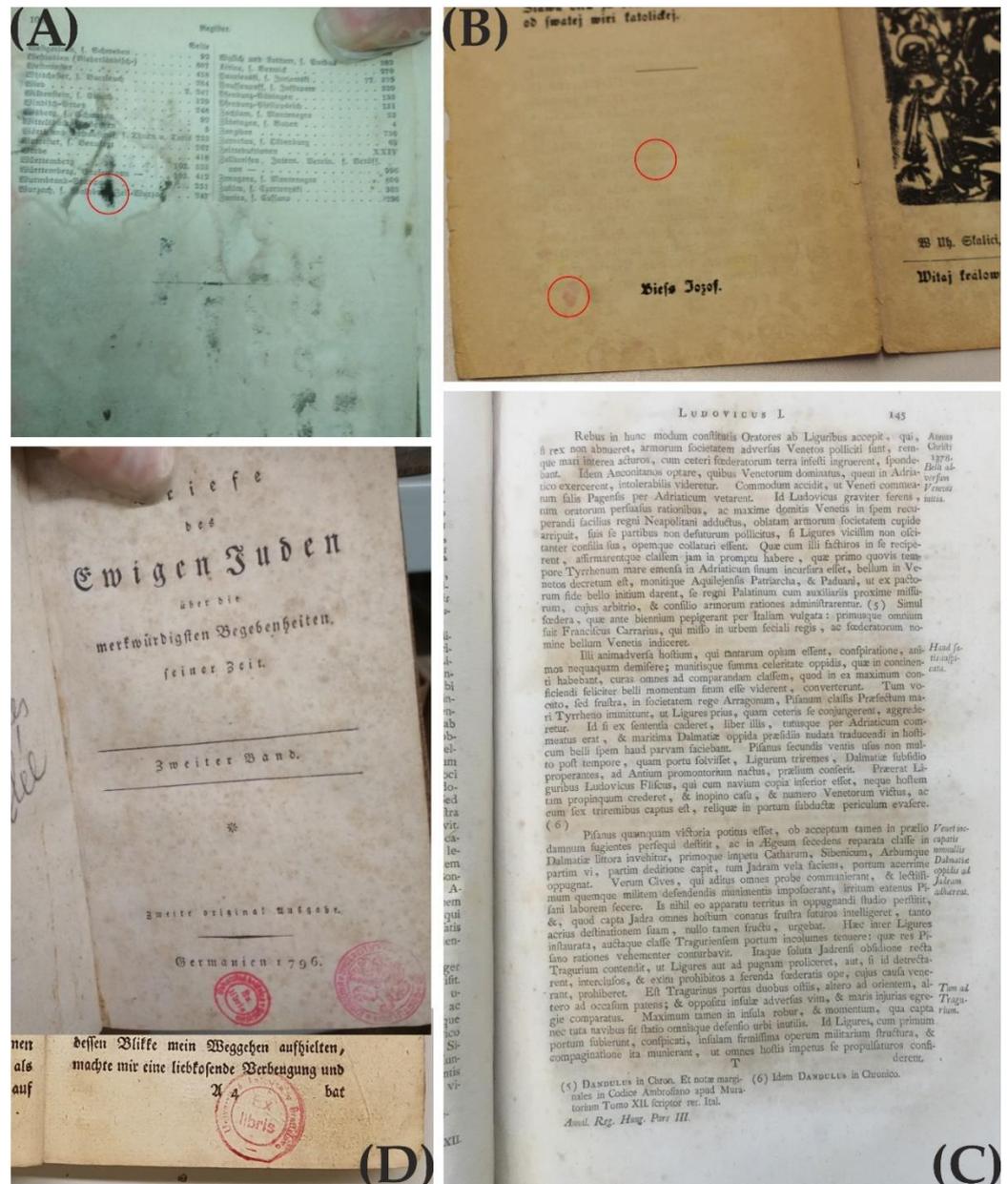


Figure 1. Several types of stains on the paper surface. (A) red circle, probably black fungal stains; many black spots are spread on the book page. (B) red circles showing presumable yellowish and pinkish fungal spots. (C) the phenomenon of foxing. (D) colored stamps on the books' pages.

Apart from fungal stains, another problem for the restorers is the colored stamps placed on a specific sheet (Figure 1), marking the book or the document as a property of the library. These stamps are usually composed of ink containing different kinds of synthetic dyes [18], and when restorers need to restore or digitalize a book or a document, they also need to remove the stamp from the sheet. Therefore, the ink of the stamp can also be considered as a potential colored stain necessary to be removed from the paper surface (personal communication, restorer Radka Benžová).

Generally, microorganisms are considered to be a threat to cultural heritage items in the form of biodeterioration. On the contrary, in the last few decades, these living organisms have been shown to be valuable restoration tools by means of their metabolic properties [19]. Many studies have reported the application of microorganisms and their enzymes in biocleaning, such as the removal of dirt, animal glue, and other organic residues from a variety of artworks [20–26].

For safety reasons, enzymatic cleaning is preferable in many cases over the application of living cells that can leave residues on the surfaces that are about to be restored and which can later serve as nutrients for dangerous microbes [19,27]. In the work of Jeszeová and colleagues [22], hydrolytic enzymes, in particular proteases, nucleases, and phosphatases, have been identified and assayed in biocleaning tests for degradation of animal glue performed on different kinds of surfaces such as wood, glass, and stone. These extracellular enzymes were excreted from *Exiguobacterium undae* and were used for the successful removal of different types of animal glue. Recently, the lipolytic abilities of the yeast *Sporidiobolus metaroseus* were also exploited to remove an oil-based patina from a historical painting from the XVIII Century [24]. Thus, it can be suggested that the isolated extracellular enzymes can be used for the restoration of different artworks and historical items without microbial contamination of the treated surface.

White-rot fungi, considered a physiological group of fungi exhibiting the activity of lignin degradation within lignocellulosic substrates, have been investigated intensively due to their great potential to degrade a wide variety of recalcitrant organic compounds and synthetic dyes [28–32]. These fungi, as well as brown-rot fungi, secrete various extracellular enzymes called lignin-modifying enzymes (LMEs), which include heme-dependent lignin peroxidases (LiP), manganese peroxidases (MnP), dye-decolorizing peroxidases (DyP), and copper-dependent laccases [33–37]. LMEs have found their use in environmental applications, including the transformation and degradation of compounds, such as different synthetic dyes, pharmaceuticals, and pesticides [38–45]. These enzymes are classified as Auxiliary Activity (AA) families in the Carbohydrate-Active enZYmes Database (CaZy database; <http://www.cazy.org/> accessed on 22 September 2022), which provides an overview of cellulose- and lignin-degrading enzymes for biotechnical applications [46,47]. Some other enzymes such as vanillyl-alcohol oxidases, glyoxal oxidases, cellobiose/quinone oxidoreductase, and cellobiose dehydrogenase are correlated with lignin degradation through enhancement of the activity of laccases, MnPs, and LiPs [34,48,49]. Therefore, it could be useful to utilize these enzymes in order to remove the stains on archival documents and books. This review will illustrate several types of fungal pigments and synthetic dyes present in historical books and paper documents and the possible removal tools which can be represented by the ligninolytic enzymes and dye-decolorizing peroxidases, moreover, their function in known and potential decolorization applications.

2. Fungal Pigments Found on Paper

Colorants produced by microorganisms are the result of metabolic processes taking place in their cell structures. These secondary metabolites are complex chemical substances that can be incorporated into spores and mycelium or excreted into a substrate such as paper [50]. Fungi can produce a variety of stains with different colors ranging from brown, black, green, yellow, purple, or pink [2]. Colorants from fungi can be either pigments or dyes. The difference between the two is due to their physical characteristics, meaning that pigments are insoluble in the given medium, whereas dyes dissolve during the application, losing their crystal or particulate structure in the process [51].

Fungal colorants represent an organism's response to light harvesting and processing, photo-protection, absorption, and neutralization of protons that could potentially damage the cell structures [52]. Some colorants are also acknowledged as enzyme inhibitors and antibiotics [53]. Dark pigments, melanins, have had several biological functions such as protection against radiation (e.g., UV), enzymatic lysis, high temperatures, or oxidizing agents; capacity for binding of metals, preventing the entry of toxic metals or concentrating essential metals; action as a virulence factor; or capacity for increasing resistance to fungicides [54].

Most of the fungal colorants are polyketide-based [55], but, in general, they can be classified into different groups, mainly: polyketides (and their derivatives), melanin, and carotenoids [56].

2.1. Polyketide Colorants

The important natural products encompassing a variety of toxins, antibiotics, therapeutic compounds, and colorants are represented by a group of chemical substances called polyketides. Due to their biosynthetic complexity and importance in the pharmaceutical industry, polyketides have drawn great attention among researchers, which has led to extensive molecular genetic studies carried out in actinomycetes and Gram-positive bacteria [57].

Polyketides are one of the largest and most structurally diverse classes of naturally occurring compounds with a variety of biological activities [58]. These compounds are biosynthesized from acyl CoA precursors catalyzed by polyketide synthases (PKSs) [59] that are members of a large enzyme family that are involved in the synthesis of different secondary metabolites such as pigments, toxins, antibiotics, and signaling molecules [60]. Pigments produced by fungi are mostly the result of the paucity of nutrients when mycelium produces secondary metabolites [61]. Fungal colorants are either polyketide- or carotenoid-based. Those that are polyketide-based are composed of tetraketides and octaketides with eight C₂ units, creating a polyketide chain [62]. Polyketides comprise azaphilones and quinones, including anthraquinones, hydroxyanthraquinones, and naphthoquinones [55,62].

Azaphilones can be defined as a small but structurally diverse subset of the polyketide class that are excreted by fungi as second metabolites [55,63]. Due to their affinity for ammonia, the name azaphilones was given to these substances. In addition, azaphilones react with amines such as proteins, amino acids, and nucleic acids to construct red or purple vinylogous γ -pyridones due to the exchange of pyrane oxygen for nitrogen [55]. Fungi possessing these compounds show bright yellow, red, or green colors of mycelia. These polyketide derivatives are produced by a variety of representatives of ascomycetes, including the genera *Aspergillus*, *Penicillium*, *Chaetomium*, *Talaromyces*, *Pestalotiopsis*, *Phomopsis*, *Eremicella*, and *Epicoccum*, as well as *Monascus* and *Hypoxylon* [64], which may colonize paper and cause the appearance of colored spots [2].

Quinones are a class of quinoid compounds widely present in nature [65]. They are derived from aromatic compounds by the conversion of an even number of methylene (-CH₂-) groups into carbonyl (C=O) groups with any necessary rearrangement of double bonds, resulting in a fully conjugated cyclic dione structure [66]. Those biosynthesized by the polyketide pathway have been isolated from fungi showing different colors, from yellow and orange to red. However, the color of the quinone isolated from a fungus does not reflect the color of that fungus. The reason for this may be that the quinone is accompanied by its reduction products, which lead to the formation of the quinhydrone complex [67]. The most produced quinones in fungi are Hydroxyanthraquinone (HAQN) colorants. These colorants are derivatives of quinones, which are widely distributed in microorganisms. In particular, they have been produced mostly by the genera *Penicillium* and *Aspergillus*, with a different color tone [68]. Specifically, *Aspergillus glaucus*, *Aspergillus versicolor*, and *Penicillium purpurogenum* have been conspicuous anthraquinone producers [69]. The shade-like appearance of a specific anthraquinone called chrysophanols has been reported to be a consequence of the metabolic processes of *Trichoderma* with the species *T. harzianum*, *T. polysporum*, *T. viridae*, and *T. aureoviride* [70]. Chrysophanols are peculiar anthraquinones giving the shade-like appearance of different colors due to the presence of chromophore for absorption of electronic spectra at diverse wavelengths [71]. Naphthoquinones, widespread quinones with naphthalene backbones, have been shown to possess biological activities such as antibacterial, antitumoral, antiviral, and antifungal activities [69,72–74]. These compounds have been synthesized in *Fusarium* and *Candida* species in response to stress [75,76]. Recently, novel naphthoquinones of a yellow color called Trypethelamide A and a dark violet-red color called 5'-hydroxytrypethelone isolated from *Trypethelium eluteriae* have been shown to exhibit antioxidant and cytotoxic activities against cancer cell lines [77]. The fungal species *Scytalidium cuboideum* has been characterized as a producer of naturally occurring red/pink pigments that can crystallize [78]. These crystalline pigments have a

naphthoquinone structure and were reported to require an easy methodology for their isolation, which is beneficial for different industries and green energy production [78–80].

2.2. Melanin

Melanin is an ubiquitous, widely distributed dark brown to black pigment in animals, plants, and microorganisms [81]. It is an insoluble and non-digestible, negatively charged polymer formed by oxidative polymerization of phenolic or indolic compounds [82–84]. There are a few forms of melanin existing in living organisms, including fungi: allomelanins, eumelanins, and pheomelanins. Eumelanins are characterized as black-brown insoluble melanin pigments partially derived from the oxidative polymerization of L-dopa via 5,6-dihydroxyindole intermediates. Pheomelanins represent a subgroup of melanins containing sulfur obtained by the oxidation of cysteinyl-dopa precursors via benzothiazine and benzothiazole intermediates. These pigments are alkali-soluble with a yellow to brown color [85]. Allomelanins are defined as melanins containing phenolic, catecholic, and 1,8-dihydroxynaphthalene (1,8-DHN) units [86].

Melanins were previously described as compounds resistant to degradation by hot acids, hot concentrated alkaline solutions, and bleaching by strong oxidizing agents, while nowadays, they stand for melanins, compounds insoluble in aqueous or organic solvents [87]. As a consequence of melanin's recalcitrant nature, it is difficult to achieve a detailed chemical analysis of melanin. Additionally, the use of hazardous chemical methods for the purification of melanin makes it harder to chemically analyze them since the chemicals used in the purification process can modify their structure [87–89]. Microscopic analysis of fungi has suggested that the melanins, in general, have a granular shape and are enclosed in the fungal cell wall [60]. Many filamentous fungi, also including paper deteriorogens [2,18], in the genera *Alternaria*, *Aspergillus*, *Auricularia*, *Cladosporium*, *Epicoccum*, *Eurotium*, *Magnapotha*, *Penicillium*, *Phomopsis*, *Sporothrix*, *Stachybotrys*, and *Wangiella*, have been shown to synthesize melanin [90]. This pigment serves as a protection from disadvantageous conditions for fungal species, even though it is not essential for their development. It has been shown that fungi that produce melanin are more resistant to UV light-induced damage, damage caused by radioactive oxidative species, temperature extremes, different hydrolytic enzymes, antimicrobial drugs, and many other harmful factors [54,91,92]. Recently, it has been discovered that due to the production of melanin fungi, particularly *Cladosporium* and *Alternaria*, species could survive extreme conditions, such as those in industrial and roadside areas. For instance, *Cladosporium* spp., *Alternaria alternata*, and *Aureobasidium pullulans* were found to colonize the walls of the damaged reactor at Chernobyl, where they were exposed to extreme levels of radiation [93]. Melanin protects conidia against hydrolysis and digestion by different enzymes secreted by other microorganisms as well as against bacterial and fungicidal proteins of animal origin [83]. Thus, it can be concluded that melanin contributes to fungal pathogenesis and survival in unfavorable conditions.

2.3. Carotenoids

Despite the fact that fungi are non-photosynthetic organisms and carotenoids are not as widely distributed in them as in plants [2], some carotene hydrocarbons have been found in several fungal species [67,94]. Carotenoids are naturally occurring pigments that are produced in an isoprenoid pathway [95]. Thus, they are comprised of isoprene units that contain light-absorbing conjugated double bonds, forming an aliphatic polyene chain and providing characteristic yellow, orange, or reddish colors [94]. Their essential biological role is in the protection of an organism from oxidative stress and membrane stabilization [94]. It has been suggested that by growing a strain of *Penicillium* sp. under high oxidative stress and in the deficiency of antioxidants, higher carotenoid colorant yields and elevated sclerotial biomass would be obtained [96]. They are also important precursors to essential metabolites such as vitamin A [95].

It has been shown that β -carotene production, as the most abundant carotenoid in nature, seems to be present in species such as *Aspergillus giganteus* [97], *Cercospora nicotianae* [98], *Penicillium* sp. [96], and *Aschersonia aleyroides* [99].

In the review written by Melo et al. [2], it has been reported that of 80 different colorants produced by fungi identified as paper colonizers, approximately 96% were of a polyketide nature and only around 4% were carotenoids. They have found only three different carotenoids in the literature that were mentioned in the context of colorants produced by fungi identified on paper: neurosporaxanthin, β -carotene, and sporopollenins. Moreover, the most common stains produced by fungi on paper have been found to be brownish and black. It has been suggested that these stains can be a mixture of different colorants. Specifically, the hallmark of *A. niger* is the formation of black or dark brown conidia, which appears to be an outcome when dark brown melanins combine with hexahydroxyl pentacyclic quinoid green pigments [100]. Additionally, the color of the fungal stains on paper may be a result of the oxidation process of a colorant in the given medium, which can further lead to the darkening of the stain. This can be seen in the brown pigment sporopollenin, an oxidative polymer of yellow β -carotene [68].

For full insight into the chemical structure of the fungal pigments, refer to Melo et al. [2].

3. Synthetic Dyes

Synthetic dyes have been largely used in the textile, paper printing, food, pharmaceutical, and other industries due to their several effective characteristics, such as cost-effectiveness in synthesis, firmness, and high stability to different factors such as light, temperature, detergent, and microbial attack [101–103]. Nowadays, these dyes represent a big problem for the environment. The color impedes the effect of the sunlight on water and negatively affects photosynthesis, which consequently reduces the food source of aquatic organisms and inhibits their growth [101,103–105]. Additionally, many synthetic dyes have been considered toxic and carcinogenic [106].

Dyes are organic compounds with the property of absorbing light in the visible spectrum (400–700 nm). This property is possible due to the presence of a chromophore or “color-bearing group”, described as a delocalized electron system with conjugated double bonds [107]. The chromophore is composed of groups of atoms, most commonly nitro ($-\text{NO}_2$), azo ($-\text{N}=\text{N}-$), nitroso ($-\text{N}=\text{O}$), thiocarbonyl ($-\text{C}=\text{S}$), carbonyl ($-\text{C}=\text{O}$), and alkenes ($-\text{C}=\text{C}-$). Chemical classification of the dyes is based on these groups present in chromophores that are illustrated in Table 1 [108]. For a chromophore to absorb electromagnetic waves, an excitation of the electrons in a molecule is required. Once this requirement is fulfilled, the molecule becomes chromogenic. In addition, most of the dyes comprise groups called auxochromes or “color helpers” that can be acidic (COOH , SO_3 and OH) or basic (NH_2 , NHR and NHR_2). These groups are mostly responsible for dye solubility and color shifting [109].

Azo dyes have been by far the most utilized class of dyes in the industry. The azo chromophore is composed of at least one azo group ($-\text{N}=\text{N}-$), but sometimes two (diaz) and three (triaz) azo groups can be present in the structure. This azo group is bonded to two groups, of which at least one is aromatic [107].

The main compound of anthraquinone dyes is anthracene, composed of three fused benzene rings with two carbonyl groups on the central ring. After the substitution of the aromatic rings, the whole structure is given the color [110]. In the anthraquinone structure, the carbonyl group acts as an electron acceptor, which is not the case in the structure of azo dyes. In order to break the structure of anthraquinone dye, an electron donor is required [111,112]. Moreover, anthraquinone dyes have a highly stabilized structure due to the resonance effect, which makes it harder to degrade them when compared with azo dyes [112,113].

Indigoid dyes are one of the oldest known classes of organic dyes. These dyes, also members of vat dyes (based on their application that takes place in a bucket or a vat), are

insoluble in water and require oxygen for color development. They have an aromatic or heteroaromatic complex with an indigoid carbon skeleton, shown in Table 1 [108,109].

Table 1. Synthetic dyes: classification and examples of dyes according to the chromophore present.

Class	Chromophore	Example
Nitrodyes		C.I. Acid Yellow 24
Nitroso dyes		Fast Green O
Azo dyes		Methyl Orange
Triphenyl methyl dyes		C.I. Basic Violet 3
Phthalein dyes		Phenolphthalein
Indigoid dyes		C.I. Acid Blue 74
Anthraquinone		C.I. Reactive Blue 19

Another important class of dyes widely used in industries producing textiles, paper, leather, plastic, food, and pharmaceuticals is the triphenylmethane class [114–116].

These dyes are composed of a triphenylmethane backbone and are considered inexpensive colorants, but have shown cytotoxic effects on living organisms [117,118].

Stamp inks used in libraries have a complex composition containing dyes or pigments, hydrocarbons, plant or mineral oil as solvent, and many other additives [119]. Jasuja et al. [120] analyzed stamp pad inks by the TLC method and found that blue stamp ink contained Crystal Violet (CV) dye. Red stamp inks have been shown to be composed of alkyl-phenols, castor oil, and polyethylene glycol as volatile organic components [121]. Nowadays, the most commonly used are water-based stamp inks, usually containing colorants such as Acid Red R, Eosin Y, and Pigment Red 112 [18]. Eosin Y is a xanthene dye containing phenolic groups with hydroxyl groups and bromine.

4. Potential Decolorizing Enzymes and Their Characteristics

Many fungal pigments and synthetic dyes are composed of phenolic compounds [2,93], and their structure can be attacked by several oxidoreductive enzymes [122]. In this big group of enzymes, the LMEs and the DyPs are also included. Here, we describe the characteristics of the most promising candidates for decolorizing applications (laccases, LiPs, MnPs, and some DyPs) and their ways of action.

4.1. Laccases

Laccases (EC 1.10.3.2), also known as benzenediol: oxygen oxidoreductase or polyphenol oxidase, are members of the family of blue multicopper oxidases that catalyze single electron oxidation of four reducing-substrate molecules, accompanied by the four-electron reduction of molecular oxygen into the water. At the end of the nineteenth century, this enzyme was extracted from the Japanese lacquer *Rhus vernicifera*, after which it was named laccase [123,124]. Laccases have been considered to be ubiquitous in many white-rot fungi studied so far. The reason for their wide distribution is their function: degradation of the complex polyphenol structure that constitutes lignin in order to break the plant's cell wall and uptake the host's nutrients [125,126]. Fungal laccases also contribute to melanin formation in different fungal species [60,127]. This enzyme showed a specific function in some insects and plants [128]. Plant laccase, together with peroxidases, contributes to the biosynthesis of lignin during protoplast regeneration [129,130]. Laccases have also been found in some bacteria, having a function in melanin synthesis and showing stability at high temperatures and pH [131–134]. Thus, laccases are widely present in nature with a role in either synthetic or degradation processes.

Fungal laccases have shown large-scale applications for industrial and biotechnological processes because they are extracellular and inducible, and they do not require a cofactor and show low substrate specificity [128]. For example, they utilize oxygen as a cofactor [135] and oxidizing agent and showed the degradation of several compounds with a phenolic structure [136]. Thus, laccases have been utilized in many bioremediation processes as well as in the degradation of a wide number of textile dyes [39,128].

Laccase is a monomeric globular glycoprotein comprised of around 500 amino acid residues organized into three consecutive domains (A, B, C) with a Greek key β barrel topology [127].

The overall reaction that is catalyzed by laccase represents the oxidation of four electrons of a reducing substrate with a two-electron reduction of dioxygen to water [$4RH + O_2 \rightarrow 4R + 2H_2O$]. The first step is a reduction of the reducing substrate carried out by the Cu^{2+} (Cu^{2+} to Cu^+) at the T1 site, which is primarily an electron acceptor. The electrons extracted from the reducing substrate are transferred to the T2/T3 site, with three Cu^{2+} ions in total. This results in changing the enzyme state from fully oxidized to fully reduced [124].

Laccases catalyze the degradation of phenolic structures through the subtraction of one electron from phenolic hydroxyl groups to form phenoxy radicals that lead to the cleavage steps, which results in aromatic ring cleavage. Moreover, this enzyme can oxidize non-phenolic compounds but in the presence of a mediator such as 2,2'-

azinobis-(3-ethylbenzthiazoline-6-sulfonate)(ABTS), 1-hydroxybenzotriazole (HBT), and 3-hydroxyanthranilic acid [124].

4.2. Lignin Peroxidases (LiPs)

LiP (E.C. 1.11.1.14) is one of the most significant ligninolytic enzymes that contains heme (protoporphyrin IX) as a prosthetic group. This enzyme was firstly discovered in nitrogen- and carbon-limited cultures of the white rot fungi *Phanerochaete chrysosporium* [137]. The presence of this enzyme was later detected in other white rot fungi such as *Trametes versicolor* [138], *Bjerkandera* sp., *Trametes trogii*, and *Phlebia tremellosa* [139,140]. LiP has been purified in bacteria *Acinetobacter calcoaceticus* [140] and also has shown activity in *Streptomyces viridosporus* [141]. This enzyme is relatively non-specific to its substrate and its main function is the oxidation of different phenolic aromatic compounds as well as a broad range of non-phenolic lignin model compounds [124]. The aforementioned microorganisms secrete LiPs as a family of isozymes with relative composition and isoelectric points (pI), depending on the growth medium and nutrient conditions [142].

4.3. Manganese Peroxidases (MnPs)

Manganese peroxidase (EC 1.11.1.13) is another lignin-modifying enzyme firstly detected in *P. chrysosporium* by Kuwahara et al. [143]. It is secreted by many Basidiomycota species such as *Panus tigrinus*, *Agaricus bisporus*, *Bjerkandera* sp., and *Nematoloma frowardii* [142]. Additionally, this enzyme has been found in bacteria *Bacillus pumilus*, *Paenibacillus* sp. [144], *Azospirillum brasilense* and *Actinobacterium S. psammoticus* [142]. MnPs are glycoproteins with a molecular weight ranging between 38 and 62.5 kDa, comprised of approximately 350 amino acids. The overall structure is shown to be similar to that of LiPs, with two domains, a heme group in the middle, 10 major helices, and a minor helix. MnPs have five disulfide bonds unlike LiPs. Another major characteristic of MgP is the manganese-binding site. The additional disulfide bond participates in the Mn-binding site alongside Aspartic and Glutamic acid residues [145].

4.4. Dye-Decolorizing Peroxidases (DyPs)

Dye-decolorizing peroxidases (EC 1.11.1.19) represent a new family of heme peroxidases with one heme as a cofactor and a role in H₂O₂-dependent oxidation of a wide range of molecules [146,147]. Basidiomycete *Thanatephorus cucumeris* Dec1 was the first species suggested to be excreting these enzymes due to its ability to decolorize 18 types of reactive, acidic, and dispersive dyes, most of which were xenobiotics [148]. In 1999, the same author succeeded in the attempt to purify DyP for the first time. It is a glycoprotein with a molecular mass of 60 kD [149]. Later, it was reported that several bacteria have a role in lignin degradation, suggesting that these bacteria possess a unique class of dye decolorizing peroxidases [150,151]. Due to the low sequence identity shown between bacterial and fungal DyPs, they are divided into four classes, from A to D. The DyP from class D are the ones from Basidiomycota [146].

The catalytic cycle of the DYPs is composed of the resting state and transient intermediates comprising compound I and compound II (Figure 2) [152]. When the enzyme is in a resting state, it contains Fe³⁺. When this resting ferric enzyme is oxidized by H₂O₂, a high-valence compound I, [Fe⁴⁺ = O]^{+•} is produced. This compound I reacts with 1 eq electrons and, from a reducing state, generates a compound II, [Fe⁴⁺ = O]⁺. Compound II is then reduced by a second substrate molecule and the resting state enzyme is produced again [152].

Fernandez-Fueyo and colleagues [153] have shown that DyPs enzymes, *Pleos-DyP1* and *Pleos-DyP4*, from fungi *Pleurotus ostreatus* have the ability to oxidize Mn²⁺ to Mn³⁺, a feature mostly present in MnPs and versatile peroxidases (VPs).

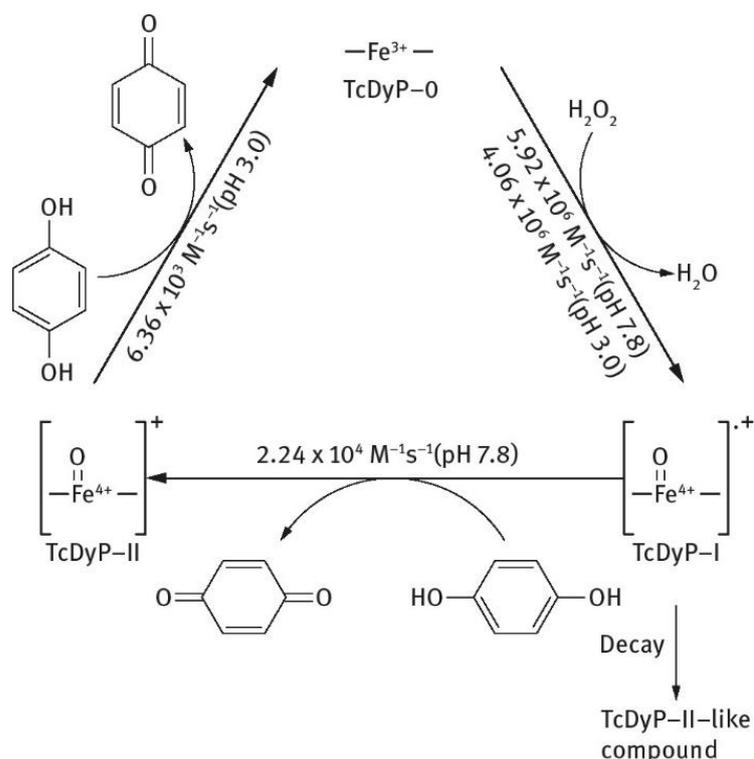


Figure 2. Schematic description of catalytic cycle of *Thermomonospora curvata* DyPs.

To compare the mechanisms of reaction of lignin modifying enzymes, including peroxidases and laccase, a schematic outline is shown (Figure 3) [142].

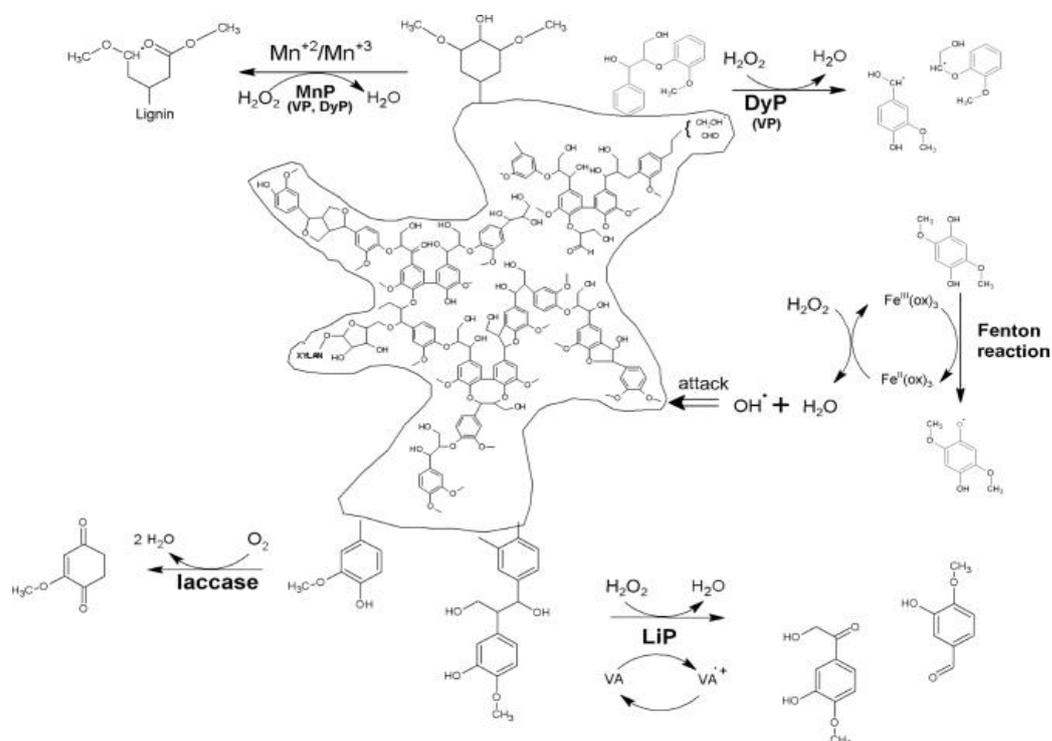


Figure 3. Comparison of lignin degradation by peroxidases and laccase. MnP—manganese peroxidase, VP—versatile peroxidase, LiP—lignin peroxidase, DyP—dye-decolorizing peroxidase, ox—oxalate, VA—veratryl alcohol.

Many stains present on paper can be caused by past fungal contaminations but also by several types of synthetic dyes composing inks and stamp inks. As we saw in the previous paragraphs, there are some similarities between the structures of synthetic dyes and fungal pigments. Therefore, starting with the known properties of LMEs and DyPs to degrade synthetic dyes, we illustrate here the potential of these enzymes to also decolorize fungal pigments.

Osmanova and colleagues [55] were the first to present the transformation pathway of Remazol Brilliant Blue R (RBBR) dye by immobilizing laccase. Laccase was derived from the white-rot fungus *Trametes pubescens* and has shown a degradation activity of 44% in 42 h. The transformation pathway involved steps of deamination and oxidation and resulted in products that were less toxic than RBBR itself [55].

Levin and colleagues have demonstrated that two different strains of *Trametes* species produce different laccase isoforms capable of degrading Acid green 27 (AG27), another anthraquinone dye. Furthermore, they have shown that besides anthraquinone dyes, the culture fluids of the strain *Trametes* sp. LA1 also decolorized the azo dye Reactive Black 5 (RB5) and triphenylmethane CV, suggesting that this strain could be a new laccase source with dye-decolorization potential [154].

Another study showed that due to the presence of laccase, MnP, and LiP, the strain *Aspergillus iizukae* EAN60 was able to decolorize RBBR and two other azo dyes, methyl red (MR) and methyl orange (MO) [155]. After 9 days of monitoring, the maximum removal of RBBR was close to 95%, while the removal of MO and MR observed after 3 days was 84.90% and 47.65%, respectively.

Sing and colleagues [122] investigated the production of ligninolytic enzymes by fungi *Marasmius cladophyllus* during its application in the decolorization of RBBR, as well as two azo dyes, such as Orange G and Congo red (CR). By monitoring specific reactions characteristic of ligninolytic enzymes, it has been found that in the presence and absence of RBBR dye, this strain produced both laccase and LiP. After in vitro RBBR dye decolorization in the culture medium containing unpurified enzymes, it has been shown that laccase activity was in agreement with the dye decolorization. Moreover, the fresh dye has been successively added to the already decolorized culture medium. This step actually sped decolorization up in the subsequent round. After reconcentration of unpurified enzymes from decolorized culture medium, the protein precipitates contained laccase that decolorized 76% of RBBR dye in less than one day. The protein precipitates were also tested for in vitro decolorization of a mono-azo dye, Orange G, and a di-azo dye, CR. It has been shown that laccase decolorized 54% of Orange G and 33% of CR at the same time. On the other hand, the protein precipitates did not show the presence of LiP [122].

Another synthetic dye, Crystal Violet (CV), has been widely utilized in medicine and by the textile industry [44]. Several studies have shown that microorganisms can successfully degrade this dye and have proposed the hypothetical pathway of CV degradation [156,157]. In the first step of degradation, it has been suggested that laccase catalyzed reduction to its leuco derivative before reductively degrading into the next intermediates to the final phenol (Figure 4) [156]. An additional enzyme that appeared to be involved in the process was aminopyrine N-demethylase.

Laccase catalytic activity in the degradation of Bisphenol A and decolorization of synthetic dyes was observed by Chairin et al. (2013) [40]. The purified enzyme from fungi *Trametes polyzona* WR710-1-induced oxidation of Bisphenol A within 3 h with redox mediator 1-hydroxybenzotriazole (HBT) present. Triphenylmethane dye (Bromophenol Blue; BRB), anthraquinone dye (RBBR), and azo dye (MO) were decolorized without the redox mediator present at a rate of 80%.

Another ascomycete, *Leptosphaerulina* sp., has been described as a producer of high amounts of laccase and MnP. These enzymes have shown the activity of decolorization of Remazol Black, Novacron Red, and Turquoise Blue.

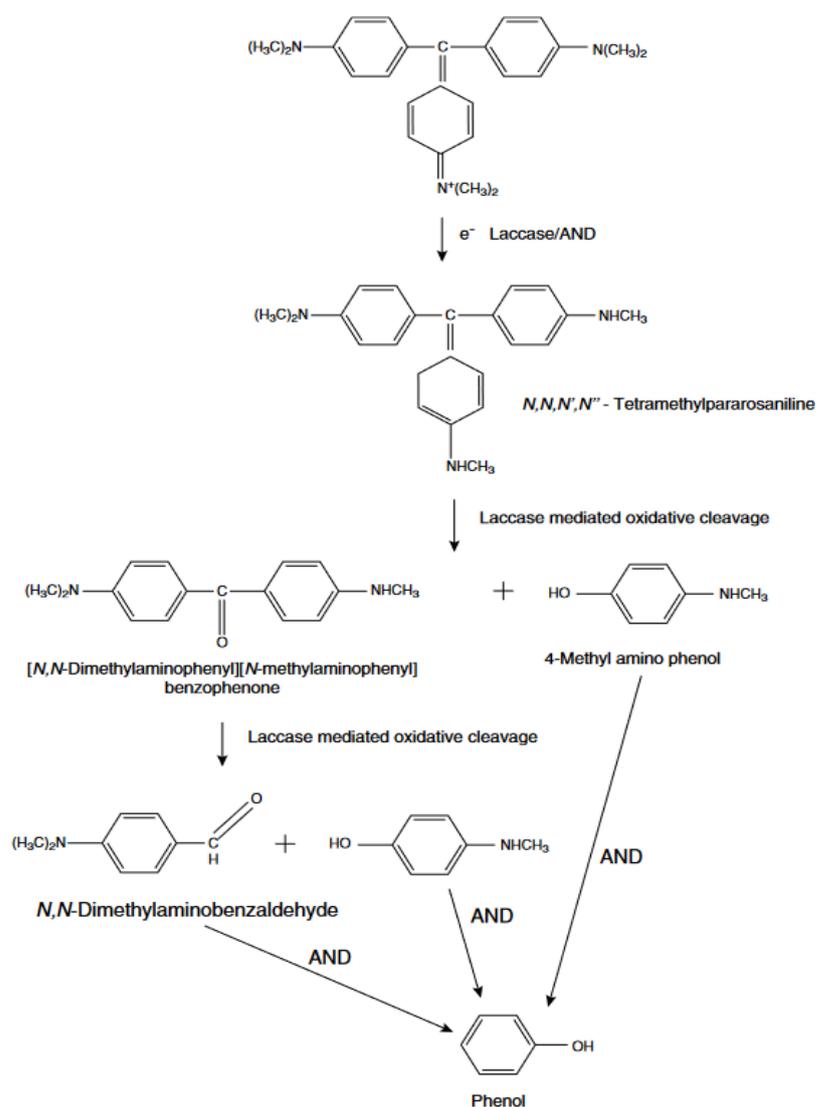


Figure 4. Mechanism of Crystal Violet degradation by laccase.

To improve the production and decolorizing activity of *Leptosphaerulina* sp., Copete-Pertuz et al. [158] co-cultivated this species with other fungal species. Based on response surface methodology, they inferred that co-cultivation with *Trichoderma viride* and *Aspergillus terreus* was the best treatment for the production of ligninolytic enzymes, specifically laccase, MnP, and versatile peroxidase, as well as the degradation of RB5 [158].

Zhang et al. [159] cloned a novel MnP gene (*mnp3*) from the fungal strain *Cerrena unicolor* BBP6 and expressed it in the yeast *Pichia pastoris*. They have demonstrated that this recombinant enzyme, rMnP3-BBP6, is an effective decolorizing agent against different dyes. Additionally, it has been shown that rMnP3-BBP6 has great potential in denim bleaching. The decolorization potential of dyes such as RBBR, MO, BPB, and CV by recombinant MnP was higher than 50%.

Besides laccase and MnPs, LiP was confirmed to decolorize synthetic dyes. This enzyme showed degradative activity in a variety of substrates such as Mg^{2+} , tryptophan, L-Dopa, hydroquinone, and synthetic dyes. Specifically, LiP isolated and purified from bacteria, *Acinetobacter calcoaceticus* decolorized azo-dyes such as MR and MO at a very high rate, 98% and 96% in one day, respectively. Other azo dyes were also decolorized by this enzyme, up to approximately 90% within two days [140]. The reaction mechanism of LiP in the decolorization of azo dyes was demonstrated by Jadhav and colleagues [160,161]. They have proposed that the cleavage of the dye is done asymmetrically, with an active site

available for a LiP to excite the molecule. This asymmetric cleavage by LiP occurs between the nitrogen of the azo group (N=N) and the carbon of the aromatic ring and results in the production of different types of naphthalene-2-sulfonic acid, depending on the type of the dye that was subjected to degradation (Figure 5) [160,161].

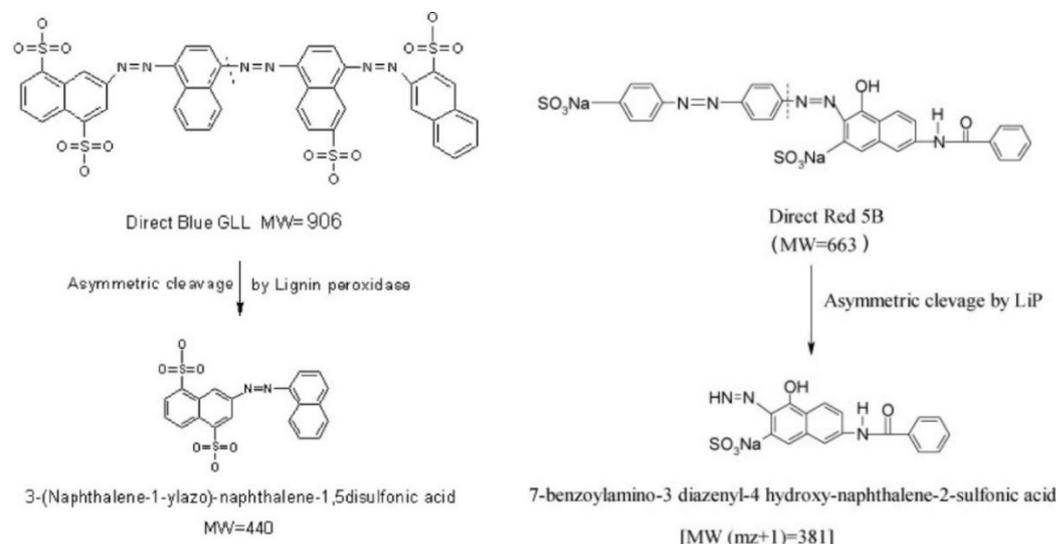


Figure 5. Reaction mechanism of LiP in a degradation pathway of azo dyes.

The anthraquinone dye degradation pathway has been reported by Sugano et al. (2009) [162]. The DyP from the basidiomycete *Thanatephorus cucumeris* was utilized in the process of degradation of RB5 dye. The pathway has the following intermediates: phthalic acid, Product 2, and Product 3 (Figure 6) [162]. These Product 2 and Product 3 were generated by the peroxidase reaction, while in the case of phthalic acid, it has been suggested that DyP acted as a hydrolase or oxygenase rather than a peroxidase [162].

Only a few attempts have been published, describing the degradation of uncharacterized fungal and melanin stains from various surfaces. Two different approaches were applied, a laccase-based method able to decolorize the fungal stains on paper and parchment [163] and a biomimetic copper-pyridine oxidation system able to remove fungal melanin from paper [164] and decolorize fungal stains from textile [165].

However, it has been shown that many quinone colorants are also synthesized by fungi are isolated from paper [2]. Quinones are a class of organic quinoid compounds widely distributed in nature [65]. A common basic structural pattern is an ortho or a para substituted dione. If this dione is conjugated to an aromatic nucleus, it is called benzene quinone, or to a condensed polycyclic aromatic system, then it can be naphthoquinones, anthraquinones, and anthracynones [65]. As was shown above, anthraquinone dyes used in textiles and in other industries can be degraded by LMEs synthesized by microorganisms, thus, we hypothesize that similar action can be applied to the quinoid colorants synthesized by different fungal species on historical paper.

When one fungal pigment from the quinone class is compared with RBBR (Figure 7) [158,159] we can see this common cyclic dione with different functional groups. In the degradation of RBBR dye, laccase appears to be the first enzyme that cleaves hydrogen bonds between N-H groups due to the reduction of Cu^{2+} to Cu^{+} , generating two compounds. In the case of fungal colorants, this cleavage would take place between the OH groups.

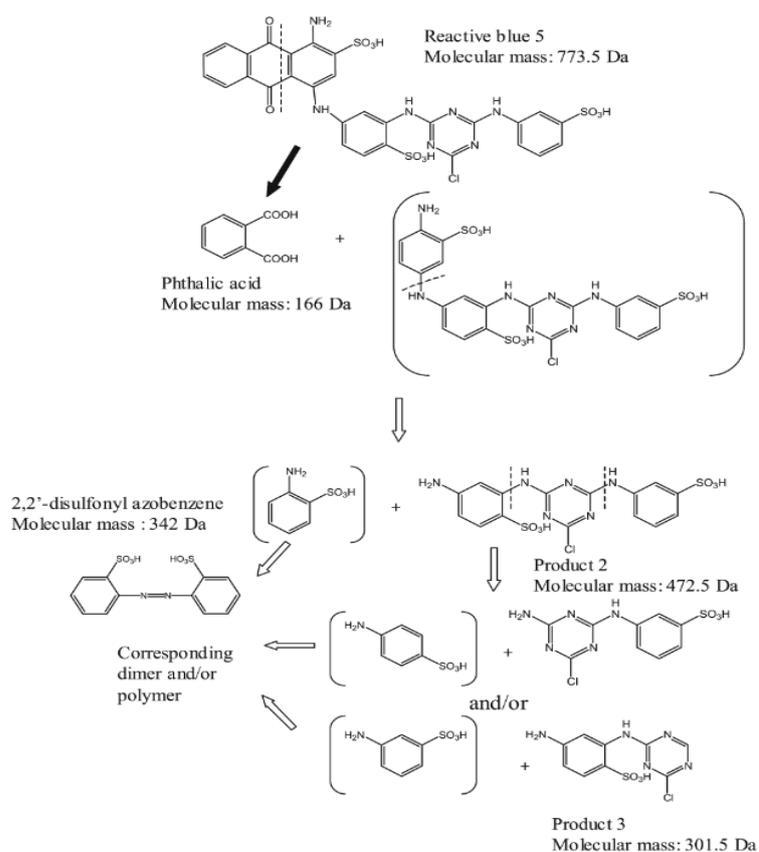


Figure 6. Degradation pathway of RB5 by DyP.

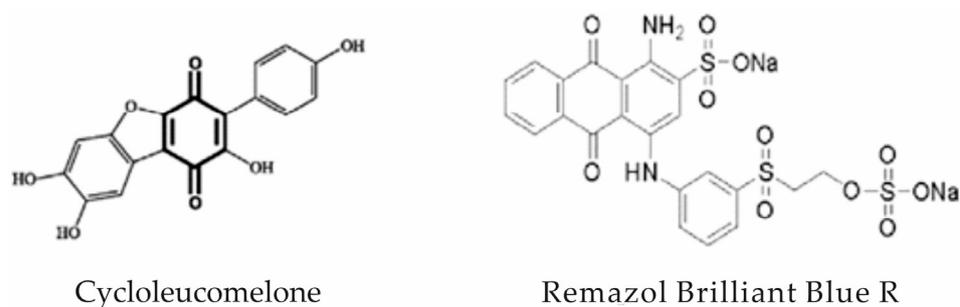


Figure 7. Fungal pigment cycloleucomelone and anthraquinone dye RBBR.

If we take a look at the structures of pigments produced by fungi and synthetic dyes, we can see that most of them contain a phenolic ring bound to the hydroxy group (for a closer look, refer to [2]). For example, simple flavipin, a yellow pigment produced by *Epicoccum purpureescens* and *Chaematomium globosum* [2,166], has a phenyl group bonded to three hydroxyl groups (Figure 8b). Under the catalysis of MnPs, one of those hydroxyl groups can be oxidized, and Mn^{3+} is reduced to Mn^{2+} . As the reaction goes further, with the series of oxidoreduction reactions, the phenolic ring would break down, causing the degradation of the pigment. Bilal et al. [39] have demonstrated a hypothetical degradation pathway of a phenol containing an organic compound (Figure 8a, [39]) as many fungal pigments and synthetic dyes are [39]. Based on the reactions shown by Bilal and colleagues [39], we proposed a similar degradation pathway of the fungal pigment flavipin (Figure 8b).

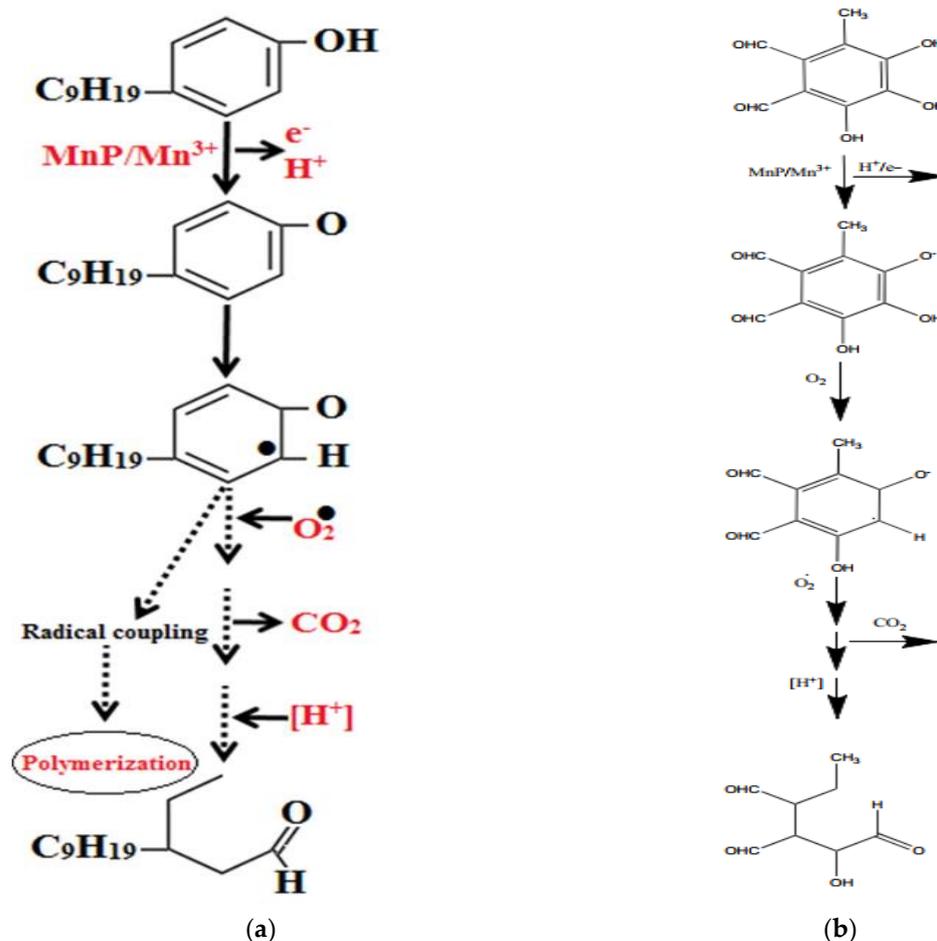


Figure 8. (a) Hypothetical degradation of phenolic compounds (b) Hypothetical degradation of fungal pigment flavipin.

Due to the characteristic of DyP to act as a hydrolase/oxygenase, this enzyme can also be employed to degrade quinone structures that have been found abundantly in filamentous fungi, leaving different color hues on the paper. In Figure 9, we have demonstrated a hypothetical degradation pathway of a randomly chosen pigment, 3-acetyl-2,8-dihydroxy-6-methoxy-anthraquinone, found in *Fusarium oxysporum*. The chemical structure of the pigment itself is not as complicated as those of synthetic dyes; thus, it can be degraded by the cleavage of the $-C(=O)-$ group. The same reaction is shown in Figure 6 [162] as the initial step of RB5 degradation. When it is broken down into simple phenolic compounds, these compounds can further be degraded by the MnP, as has previously been shown in Figure 8b.

Additional problems that libraries and book restorers are dealing with are stamp inks. Those inks are water-based and usually contain colorants such as Acid Red R, Eosin Y, and Pigment Red 112 [18]. Eosin Y is a xanthene dye containing phenolic groups with hydroxyl groups and bromine. Thus, this structure could also be degraded by MnPs. Acid Red R and Pigment Red 112 are typical azo dyes, so they should easily be degraded by laccase or LiP in a similar way as is shown in Figure 5 [160,161].

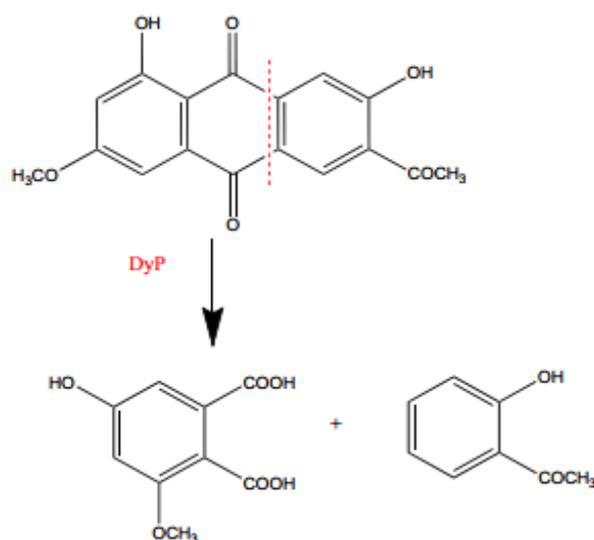


Figure 9. Degradation of pigment 3-acetyl-2,8-dihydroxy-6-methoxy-anthraquinone.

5. Concluding Remarks and Future Outlook

In summary, the data reviewed above illustrated information about the pigments produced by fungi that can colonize paper cultural heritage, synthetic dyes, ligninolytic and decolorizing enzymes, and their application. During the past few years, enzyme-based catalytic engineering has been used in the bioremediation of industrial pollutants. In this regard, the development of enzyme-based immobilized systems suggests a great potential for biotechnological applications for cultural heritage bioremediation purposes. Ligninolytic and dye-decolorizing enzymes have demonstrated features to degrade, decolorize, or detoxify several types of synthetic dyes and pollutants. When these dyes are compared with the fungal pigments found in the books, we hypothesized that the described enzymes have the ability to also degrade these structures. Depending on the side groups of the pigments' structure, with the gathered action of laccase that can catalyze oxidation or demethylation, DyP that can cleave the $-C(=O)-$ group and MnP that catalyzes oxidation-reduction reactions, specific fungal pigments could be degraded, which further leads to the cleaning of the stained paper. Similarly, water-based stamp inks can also be removed by the same principle since these inks are composed of azo dyes, previously shown to be removed by ligninolytic enzymes secreted by different microorganisms.

Even though the costs of biocleaning methods are comparable to those of chemical-physical techniques, the biological approach is more favorable in terms of both environmental and human health safety. Thus, future research should focus on the development of new protocols for the large-scale production of decolorizing enzymes and their application on paper and perhaps also on architectural cultural heritage since the direct application of living microorganisms can leave residues on the surfaces that are about to be restored and later serve as nutrients to other dangerous microbes. Additionally, the discovery of new chemical reactions catalyzed by ligninolytic and dye decolorizing enzymes represents a great rationale for their further utilization in bioremediation processes.

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