
Enzymatic Degradation of Azo Dyes – A Review

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ABSTRACT

Synthetic organic colourants (e.g., azo dyes) are used commonly in different industries ranging from food, textile production, printing and pharmaceuticals. The majority of these dyes are recalcitrant, so that they can impart colour on various raw materials. Certain dyes, dye precursors and some aromatic amine metabolites produced through biotransformation of dye compounds have been shown to be carcinogenic. The release of dyes into the environment constitutes a small proportion of water pollution, and the coloured wastewaters represent a serious environmental problem and a public health concern. Colour removal, especially from textile wastewaters, has been a big challenge over the last few decades; until now there is no single and economically attractive treatment that can effectively decolourize textile mill effluent. This review narrates the different enzymatic mechanisms in the reduction of azo dyes.

Keywords: Remediation; azo dyes; enzymes; enzymatic degradation pathways

1. Introduction

The textile mills daily discharge millions of liters of untreated effluents in the form of wastewater into public drains that eventually empty into rivers. Most of them are recalcitrant in nature, especially azo dyes. The stability and their xenobiotic nature of reactive azo dyes makes them recalcitrant hence they are not totally degraded by conventional wastewater treatment processes that involve light, chemicals or activated sludge (Chung *et al.*, 1992). The dyes are therefore released into the environment, in the form of coloured wastewater. This can lead to acute effects on exposed organisms due to the toxicity of the dyes, phytoplanktons form abnormal colouration and reduction in photosynthesis because of the absorbance of light that enters the water (Duran and Esposito 2000; Mester and Tien 2000). This also alters the pH, increases the biochemical oxygen demand (BOD) and chemical oxygen demand (COD), and gives the rivers intense colourations and public is greatly concerned about water quality. The presence of unnatural colours is aesthetically unpleasant and tends to be associated with contamination. Without adequate treatment these dyes will remain in the environment for an extended period of time (Olukanni *et al.*, 2006). Several methods were adapted for the reduction of azo dyes to achieve decolourization. These include physiochemical methods (Droste, 2004) such as filtration, specific coagulation, use of activated carbon, chemical flocculation etc. Some of these methods (*viz* reverse osmosis, nanofiltration, multiple effect evaporator (MEE)) are found to be effective but quite expensive (Maier *et al.*, 2004; Do *et al.*, 2002). Table.1 summarizes the advantages and disadvantages of the current methods of dye removal from industrial effluent (Andre *et al.*,

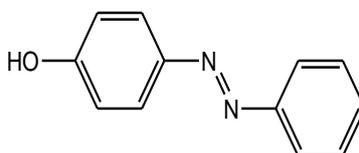
2007). Biotreatment offers a cheaper and environmentally friendlier alternative for colour removal in textile effluents. A number of microorganisms have been found to be able to decolourize textile dyes including bacteria, fungi, and yeasts (Olukanni *et al.*, 2006; Wesenberg *et al.*, 2003; Kirby *et al.*, 2000; Martins *et al.*, 1999; Banat *et al.*, 1996; Paszczynski *et al.*, 1991; Gingell *et al.*, 1971). They have developed enzyme systems for the decolourization and mineralization of azo dyes under certain environmental conditions (Anjali *et al.*, 2006). Although dye molecules display a high structural variety, they are degraded by only few enzymes. These biocatalysts have one common mechanistic feature. They are all redox-active molecules and thus, exhibit relatively wide substrate specificities (Duran and Esposito 2000; Mester and Tien 2000). Preferentially, suitable organisms excrete the active enzymes into the medium. On the other hand dye molecules are transport into the cells; another important requirement for these organisms is its resistance against toxic effects of dyes and other substances present in the effluent. It has been reported after many studies decolourizing rates of dyes by microorganisms' decreases with increasing dye concentrations above certain levels. This may be a limiting factor for bio-elimination. Therefore, in cases where the target molecule or additives inhibit growth, isolated enzyme systems may be preferred.

Table 1: Advantages and disadvantages of the current methods of dye removal from industrial effluent (Courtesy: Andre *et al.*, 2005)

Physical/chemical methods	Advantages	Disadvantages
Fentons reagent	Effective decolourisation of both soluble and insoluble dyes	Sludge generation
Ozonation	Applied in gaseous state: no alteration of volume	Short half-life (20 min)
Photochemical	No sludge production	Formation of by-products
NaOCl	Initiates and accelerates azo-bond cleavage	Release of aromatic amine
Cucurbituril	Good sorption capacity for various dyes	High cost
Electrochemical destruction	Breakdown compounds are non-hazardous	High cost of electricity
Activated carbon	Good removal of wide variety of dyes	Very expensive
Peat	Good adsorbent due to cellular structure	Specific surface areas for adsorption are lower than activated carbon
Wood chips	Good sorption capacity for acid dyes	Requires long retention times

Silica gel	Effective for basic dye removal	Side reactions prevent commercial application
Membrane filtration	Removes all dye types	Concentrated sludge production
Ion exchange	Regeneration: no adsorbent loss	Not effective for all dyes
Irradiation	Effective oxidation at lab scale	Requires a lot of dissolved O ₂
Electrokinetic coagulation	Economically feasible	High sludge production

The chemical structure of dyes in general is comprised of a conjugated system of double bonds and aromatic rings. The major classes of dyes have anthraquinoid, indigoid, and azo aromatic structures. All of these structures allow strong π - π^* transitions in the UV-visible area, with high extinction coefficients that allow us to consider these structures as dye chromophores. Of all of these structures, the azo aromatic one is the most widespread dye class in the industry. They may have one or more azo (N=N) groups. Azo dyes with one azo group are called mono-azo dyes, with two azo groups, di-azo dyes, followed by tri-azo azo dyes with more than three azo linkages are designated polyazo dyes. The most commercially important are mono-azo dyes and di-azo dyes, tri-azo dyes, whereas polyazo are much less important. The main drawback of this class of dyes is that they are not easily degraded by aerobic bacteria, and with the action of anaerobic or microaerobic reductive bacteria. There is a great environmental concern about the fate of these dyes, and their harmful metabolites especially on reactive dyeing of cellulosic fibers, where large amounts of unbound dye are discharged in the effluent (Pierce, 1994). They can form toxic and/or mutagenic compounds such as aromatic amines like naphthylamines, chloro aniline etc... (Chung *et al.*,1992; Wong *et al.*,1996)



A typical azo compound, 4-hydroxyphenylazobenzene

2. Enzymatic Degradation of Azo Dyes

In the case of enzymatic remediation of azo dyes, azo reductases and laccases seem to be the most promising enzymes. Laccases have been shown to decolourize a wide range of industrial dyes (Rodriguez *et al.*, 1999; Reyes *et al.*, 1999). Low molecular weight compounds like 2, 2'-azino-bis-(3-ethylbenzothiazoline-6-sulfonic acid) (ABTS) may also be necessary to mediate the actual electron transfer steps of laccases (Wong and Yu, 1999). In the presence of redox mediators decolourization events of degradable dyes could be significantly enhanced (Reyes *et al.*, 1999; Abadulla *et al.*, 2000; Soares *et al.*, 2001).

Similarly, in the case of peroxidases, addition of veratryl alcohol was shown to positively influence the decolourization of azo and anthraquinone dyes catalyzed by lignin peroxidase. However, this effect may either be attributed to the protection of the enzyme of being inactivated by hydrogen peroxide or to the completion of the oxidation-reduction cycle of the lignin peroxidase rather than to just redox-mediation (Young and Yu, 1997).

2.1 Degradation of azo dyes by azo reductases

Azo reductases catalyze the reaction only in presence of reducing equivalents like FADH and NADH. Most of the azo dyes have sulphonate substituent groups and a high molecular weight and they are unlikely to pass through cell membranes. Therefore, the reducing activity referred to the dye is not dependant on the intracellular uptake of the dye (Robinson *et al.*, 2001). Russ *et al.* suggested that bacterial membranes are almost impermeable to flavin-containing cofactors and, therefore, restrict the transfer of reduction equivalents by flavins from the cytoplasm to the sulphonated azo dyes. Thus, a mechanism other than reduction by reduced flavins formed by cytoplasmic flavin-dependent azoreductases must be responsible for sulphonated azo dye reduction in bacterial cells with intact cell membranes (Russ *et al.*, 2000).

One such mechanism involves the electron transport-linked reduction of azo dyes in the extra-cellular environment. To achieve this, the bacteria must establish a link between their intracellular electron transport systems and the high molecular weight, azo dye molecules. For such a link to be established, the electron transport components must be localized in the outer membrane of the bacterial cells (in the case of gram-negative bacteria), where they can make direct contact with either the azo dye substrate or a redox mediator at the cell surface (Myers and Myers, 1992). In addition, Gingell and Walker have shown that low molecular weight redox mediator compounds can act as electron shuttles between the azo dye and an NADH (nicotinamide adenine dinucleotide)-dependent azo reductase that is situated in the outer membrane.

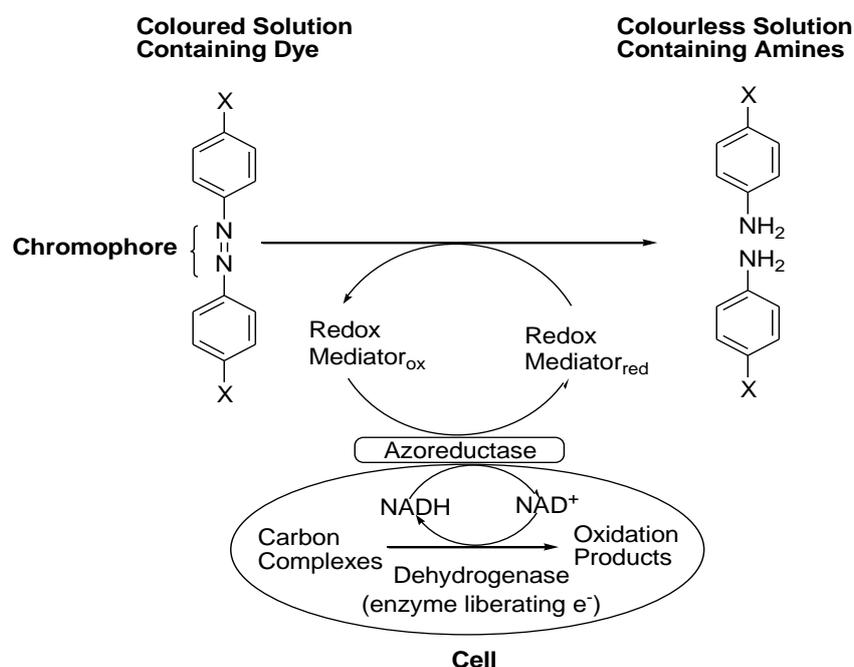


Figure 1: Proposed mechanism for reduction of azo dyes by azo reductase (Courtesy: Keck *et al.*, 1997)

These enzymes are multicopper phenol oxidases that decolourize azo dyes through a highly nonspecific free radical mechanism forming phenolic compounds, thereby avoiding the formation of toxic aromatic amines (Chivukula *et al.*, 1995; Wong and Yu, 1999). Fig. 2 represents the suggested model for the activity of laccase on one of the azo dyes, 3-(2-hydroxy-1-naphthylazo) benzenesulfonic Acid.

According to this suggested model, laccase oxidizes the phenolic group of the phenolic azo dye, with the participation of one electron generating a phenoxy radical which is sequentially followed by oxidation to a carbonium ion. A nucleophilic attack by water on the phenolic ring carbon bearing the azo linkage to produce 3-diazenyl-benzenesulfonic acid (III) and 1, 2-naphthoquinone then takes place (Susana *et al.*, 2005). The radicals which are formed by the one-electron oxidation of dye by laccase will react with 1, 2-naphthoquinone rather than be oxidized. The formed radicals can undergo oxidation to yield compound VIII, reduction to yield compound X, or further polymerization and again oxidation to form compound IX. According to the reported mechanism, cross-coupling between the reactive species results in the formation of C-C and C-O bonds between phenolic molecules and of C-N and N-N bonds between aromatic amines. By phenolic cross-coupling, an electron is removed from the hydroxyl group, generating an alkoxy radical. Alkoxy free radical forms dimers in the *ortho* and *para* positions with the hydroxyl groups. Phenolic radicals can be further oxidized to yield oligomeric products. Under certain conditions, the C-C-formed dimers can take part in coupling reactions to form extended quinines (Andrea *et al.*, 2005).

2.3 Azo dye reduction by peroxidases

Fungal systems appear to be most appropriate in the treatment of azo dyes (Ezeronye and Okerentugba, 1999). The capacity of fungi to reduce azo dyes is related to the formation of exo enzymes such as peroxidases and phenol oxidases. Peroxidases are hemoproteins that catalyze reactions in the presence of hydrogen peroxide (Duran *et al.*, 2002). Lignin and manganese peroxidases (MnP) have a similar reaction mechanism that starts with the enzyme oxidation by H₂O₂ to an oxidized state during their catalytic cycle. The first ligninolytic peroxidases were isolated from *P. chrysosporium* and called lignin peroxidase (LiP) (Glenn *et al.*, 1983; Tien *et al.*, 1983) and manganese peroxidase (MnP) (Kuwahara *et al.*, 1984; Wariishi *et al.*, 1988). LiP catalyzes the oxidation of non phenolic aromatic compounds like veratryl alcohol. MnP preferentially oxidizes Mn²⁺ to Mn³⁺, and the Mn³⁺ is responsible for the oxidation of many phenolic compounds (Glenn *et al.*, 1986). Mn²⁺ is necessary for completion of the catalytic cycle of MnP. The MnP isoenzymes from *Pleurotus* species and *Bjerkandera adusta* differ from the isoenzymes isolated from *P. chrysosporium* because they are able to oxidize 2, 6-dimethoxyphenol (DMP) and veratryl alcohol in a Mn²⁺-independent reaction (Glenn *et al.*, 1983; Tien *et al.*, 1983; Ezeronye and Okerentugba, 1999). Several reports have shown that LiP or MnP from *P. chrysosporium* is directly involved in the degradation of various xenobiotic compounds and dyes (Paszczynski *et al.*, 1991). In many studies preferential degradation of different sulfonated azo dyes either by MnP and Mn²⁺ or by LiP was demonstrated (Paszczynski *et al.*, 1991; Pasti *et al.*, 1992). Metabolic pathways for azo dye degradation by fungal peroxidases have been postulated in Fig. 3 (Goszczynski *et al.*, 1994; Spadaro *et al.*, 1994).

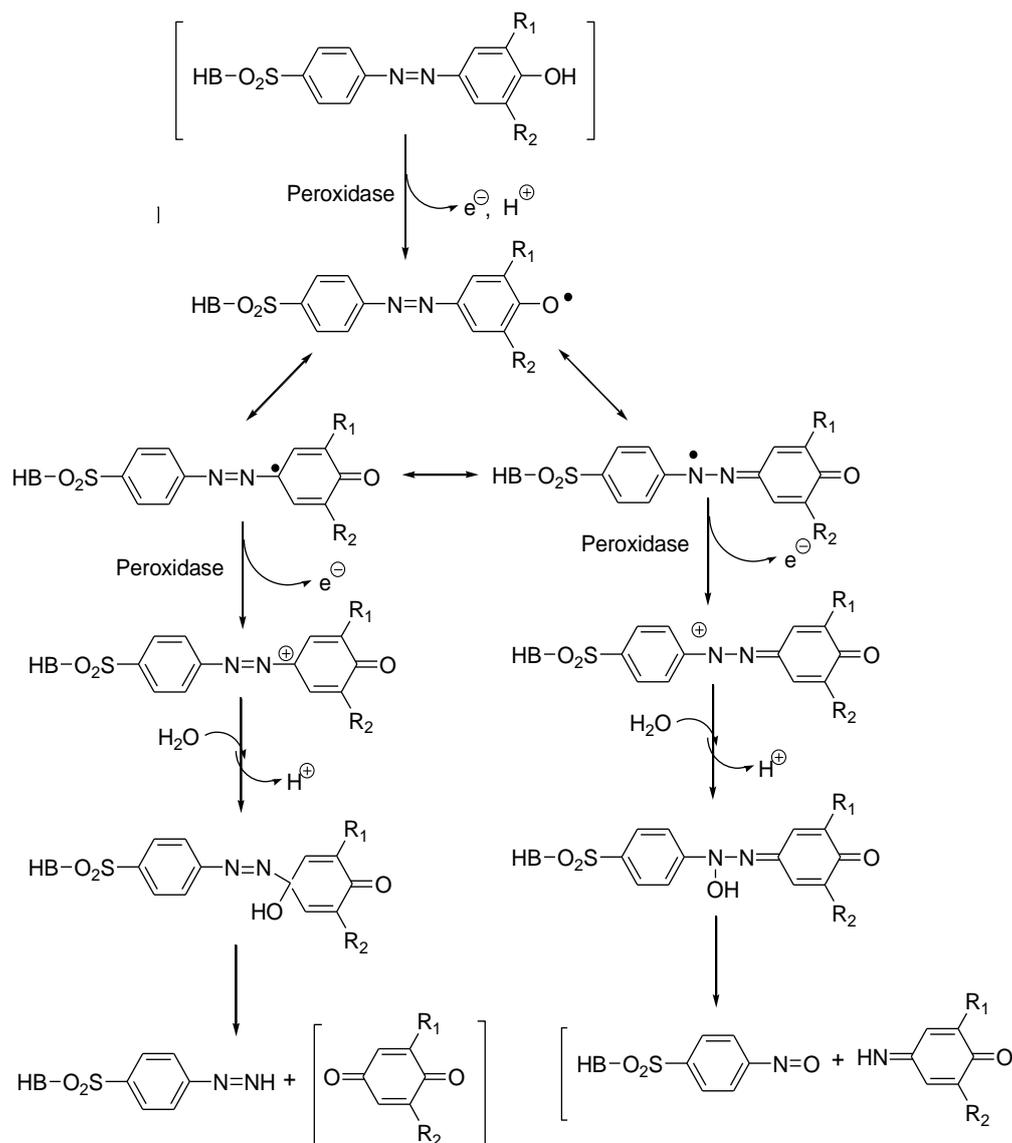


Figure 3: Proposed mechanism for alternative asymmetrical and symmetrical cleavages of sulfonated azo dyes by *P. chrysosporium* and *S. chromofuscus* peroxidases (ligninases). The compounds represented by structures in brackets were found in the reaction mixture. Azo dye 1, $R_1 = R_2 = CH_3$ and $B = O$; azo dye 2, $R_1 = H$, $R_2 = OCH_3$, and $B = NH$.

3. Conclusion

Enzymatic processes are very promising for the decolourization of synthetic azo dyes. Here we have reviewed the initial steps in the pathways for the enzymatic degradation of recalcitrant synthetic azo dyes. To understand the decolourization and degradation mechanism of azo compounds under aerobic conditions, detailed information is needed about the initial enzymatic transformation of azo linkages. The authors in our laboratory are involved in the isolation of potent solo bacterium and bacterial consortium for efficient decolourization of azo dyes from textile mill effluents. Together with this isolation, we too extend our studies to understand of how chemical structure influences the susceptibility of these man-made compounds to degradation, these studies could help to develop a new generation of readily degradable dyes; such research could also exemplify how a new generation of less recalcitrant chemicals might be produced in the future.

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