Decolorization of triphenyl methane dyes by
Fomitopsis fellei

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ABSTRACT

Six triphenylmethane dyes viz., bromophenol blue, basic fuchsin, methyl violet, methyl green, ethyl violet and malachite green were studied for their decolorization by Fomitopsis fellei. Among, basic fuchsin (98%) was maximum decolorized followed by bromophenol blue (96.8%). However, the rate of bromophenol blue decolorization was very fast. There was no correlation between the lignolytic activity and dye decolorization of the dyes. The highest laccase and lignin peroxidase activities were observed in basic fuchsin (46 U/mL) and methyl green (44 U/mL) respectively after 21 days of incubation, which were poor dye degraders. The triphenylmethane reductase enzyme was the responsible enzyme for the decolorization of these triphenyl methane dyes. The treatment by using fungal organisms was considered to be the cost-effective and ecofriendly method of decolourization of effluents discharged from the dye industries.

Keywords: Triphenyl Methane Dyes; Fomitopsis fellei; Dye Decolorization; Lignolytic Enzymes; Triphenyl Methane Reductase

1. INTRODUCTION

Synthetic dyes are extensively used in the textile industries and significant proportion appears in the form of wastewater and is spilled into the environment. More than 10,000 synthetic dyes include the azo, anthroquinone and triphenylmethane dyes, all of which are generally considered as the xenobiotic compounds, which are very recalcitrant to biodegradation [1]. Triphenylmethane dyes are aromatic xenobiotic compounds that are used extensively in many industrial processes, such as textile dyeing, paper printing, and food and cosmetic manufacture [2]. They are known to be highly toxic to mammalian cells and mutagenic and carcinogenic to humans [3-6]. Based on their potential for adverse human health effects, most countries have nominated triphenylmethane dyes as hazardous material and prohibited the use of them in aquaculture and food industry. However, they are still used in some areas due to their relatively low cost, ready availability and efficacy [7]. Since triphenylmethane dyes occur as contaminants and potential human health hazards, there is concern about the fate of them in aquatic and terrestrial ecosystems [8,9]. Considerable amounts of these toxic and mutagenic dyes are discharged into wastewater treatment facilities and thus impose a selective pressure on the microbial flora residing in wastewater habitats. Several physicochemical methods include adsorption, chemical precipitation and flocculation, photolysis, chemical oxidation and reduction, electrochemical treatment and ion pair extraction have been used to eliminate the colored effluents from wastewater [10] but these methods are expensive, of limited applicability and produce large amounts of sludge. As a better alternative, therefore, the development of biological processes using microorganisms for the treatment of dye-containing wastewater has become increasingly important.

Several triphenylmethane dye-decolorizing microorganisms have been reported and their characteristics were reviewed [11]. Only a few studies have reported the bioremoval potentials of brown rot fungi for different dyes [12-17]. Further research work on such fungal strains is much needed in bioremediation perspective. Fomitopsis fellei is a basidiomycetous fungi belongs to the family Fomitopsidaeae. Brown rot fungus F. fellei has been found to have decolorization capacity but only limited literature is available [18]. The present work was aimed at evaluating basidiomycetous fungi for its ability to de-
colorize several synthetic dyes in shake condition. An attempt has also been made to analyze the concomitant association of ligninolytic enzymes (lignin peroxidase and laccase) production during the decolorization by this fungus.

2. MATERIAL AND METHODS

2.1. Chemicals

Triphenyl methane dyes i.e. Basic fuchsin (BF), Methyl green (Mg), Ethyl violet (Ev), Methyl violet (Mv) were procured from Himedia Laboratories Pvt. Ltd. (Mumbai, India) and Malachite green (MG) and Bromophenol blue (BB) were purchased from Merck Specialities Private Limited (Mumbai, India) and Qualigens Fine Chemicals Ltd. (Mumbai, India) respectively. All media components and chemicals used were of highest purity grade.

2.2. Isolation of Microorganism and Culture Conditions

The wood decaying fungus F. feei was collected from the infected wood logs at Pakhal forest (latitude 18°1’,18.82”N and longitude 80°9’22.16”E) Warangal district, Andhra Pradesh, India. The tentatively identified fungus was phylogenetically confirmed by molecular analysis of D2 region of 28S rDNA as *Fomitopsis feei* (AY 515327.1). All along the duration of the experiment, fungus was maintained on Malt extract agar medium (MEA (g/L): Malt extract 15, Dipotassium hydrogen phosphate 1, Ammonium chloride 1, 1 N Citric acid 15, Agar 20) and stored at 4°C in shaking incubator at 150 rpm speed. After regular intervals include 7, 14 and 21 days of incubation, the fungal culture was harvested by filtering through Whatmann No.1 filter paper and the culture filtrate was centrifuged (12,000 × g; 20 min) at 4°C. Decolorization and ligninolytic enzyme activities was determined from the cell-free supernatant. pH of the supernatant was measured every time using Elico pH meter. The uninoculated dye-containing medium was used as control. The decolorization efficiency was determined using the following equation [19]:

\[
\% \text{decolorization} = 100 \times \frac{A_{ini} - A_{fin}}{A_{ini}}
\]

\(A_{ini} = \) Initial absorbance
\(A_{fin} = \) Final absorbance of dye after incubation time.

2.4. Dry Mycelium Weight

Dry mycelium weight (DMW) of the fungal mass was obtained by filtering the contents of each flask through pre-weighed Whatmann No.1 filter paper and then it was left to dry to a constant weight at 70°C. DMW was expressed in terms of grams of biomass per liter of culture.

2.5. Lignolytic Enzyme Assays

Laccase activity was measured [20] by taking the optical density of reaction mixture, prepared by mixing 0.5 mL of distilled water, 1 mL of sodium acetate buffer (pH 4.5), 0.5 mL of substrate solution (46 mM guaiacol) to 0.5 mL of crude enzyme (cell free supernatant) at 440 nm up to 90 sec with 30 sec of time interval. Lignin peroxidase activity was evaluated [21] by following the same procedure of laccase but 0.5 mL of hydrogen peroxide was added in addition to that mixture. For these two enzymes, one activity unit was defined as the amount of enzyme necessary to oxidize one μmol of substrate per minute.

2.6. Triphenylmethane Reductase (TMR) Assay

The standard assay [22] system for TMR comprised 250 μL of 20 mM sodium phosphate buffer (pH 7.0), 250 μL of 20 μM basic fuchsin, 250 μL of 0.1 mM NADH, and 250 μL of the enzyme in a total volume of 1 mL. Each reaction was initiated by the addition of the enzyme, and the initial reaction rate was determined by monitoring the decrease in absorbance at 544 nm in the first 2 min in a temperature-controlled cuvette in a 1.0-cm light path at 40°C. One unit of enzyme activity was defined as the amount that catalyzed the reduction of 1 μmol of basic fuchsin per min per mL. The corresponding wavelengths were used when other triphenylmethane dyes were tested in place of basic fuchsin.

3. RESULT

Table 1 indicates the result showing pH, dry mycelial weight (DMW), percentage of decolorization and lignolytic enzyme activities of *F. feei* in shaking condition. With increasing incubation time, pH of the fungal culture broth in all dyes was decreased, which is inversely proportional to dry mycelial weight and percentage of de-
Table 1. Decolorization of triphenyl methane dyes at 0.001% by *Fomitopsis ffei* and their effect on growth and lignolytic enzyme production.

<table>
<thead>
<tr>
<th>Name of the dye</th>
<th>pH</th>
<th>Dry weight (g/L)</th>
<th>Decolorization (%)</th>
<th>Laccase (U/mL)</th>
<th>Lignin peroxidase (U/mL)</th>
</tr>
</thead>
<tbody>
<tr>
<td>DOI 7 14 21</td>
<td>7</td>
<td>14 21 7 14 21</td>
<td>7 14 21 7 14 21</td>
<td>7 14 21</td>
<td>7 14 21</td>
</tr>
<tr>
<td>BF</td>
<td>2.15 2.12</td>
<td>1.99 7.28 ± 0.01 8.12 ± 0.00 11.16 ± 0.02</td>
<td>73.8 81.7 98.0 12 ± 0.01 14 ± 0.02 46 ± 0.01 24 ± 0.01 30 ± 0.0036 ± 0.02</td>
<td></td>
<td></td>
</tr>
<tr>
<td>MV</td>
<td>6.23 5.18</td>
<td>2.12 4.08 ± 0.02 5.18 ± 0.01 11.28 ± 0.01</td>
<td>85.3 88.7 91.5 - - 8 ± 0.02 - - 6 ± 0.01</td>
<td></td>
<td></td>
</tr>
<tr>
<td>EV</td>
<td>6.36 6.19</td>
<td>5.91 3.32 ± 0.00 4.04 ± 0.02 6.56 ± 0.01</td>
<td>63.1 79.8 83.1 - - - 2 ± 0.02 - -</td>
<td></td>
<td></td>
</tr>
<tr>
<td>MG</td>
<td>4.54 2.18</td>
<td>2.04 3.56 ± 0.01 7.64 ± 0.00 9.32 ± 0.02</td>
<td>81.8 85.1 89.6 - 16 ± 0.01 34 ± 0.00 - 18 ± 0.01 28 ± 0.02</td>
<td></td>
<td></td>
</tr>
<tr>
<td>BB</td>
<td>2.42 2.14</td>
<td>1.98 5.84 ± 0.00 7.16 ± 0.01 9.20 ± 0.03</td>
<td>88.1 92.4 96.8 - 6 ± 0.02 42 ± 0.01 4 ± 0.01 14 ± 0.02 14 ± 0.03</td>
<td></td>
<td></td>
</tr>
<tr>
<td>MetG</td>
<td>5.75 2.34</td>
<td>2.20 3.32 ± 0.02 7.72 ± 0.00 10.1 ± 0.01</td>
<td>34.2 65.1 88.3 - - - - 44 ± 0.01</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

DOI: days of incubation; The results were reported as the averages of the results on three samples ± SD.

colorization. The growth of *F. ffei* was not inhibited by the dye solution and was increased with the days of incubation, with all dyes. The maximum growth was observed in basic fuchsin after 21 days of incubation (11.16 g/L).

Maximum decolorization with *F. ffei* was observed in basic fuchsin after 21 days of incubation (98%) followed by bromophenol blue (96.8%). The rate of bromophenol blue decolorization was very fast (Figure 1) may be due to the less complex structure of bromophenol blue compared to the other two dyes tested. The rate of decolorization in basic fuchsin (Figure 2) was increased from 73.8% (7 days) to 98% (21 days). 85.3% of decolorization was observed in methyl violet (Figure 3) after 7 days of incubation, which was then increased to 91.5% after 21 days of incubation. Although the initial decolorization of methyl green (Figure 4) was less in 7 days (34.2%), it could reach up to 88.3% after 21 days of incubation but there was no laccase activity. However, it is strange that the decolorization of ethyl violet (Figure 5) was increased from 63% to 83% from 7 to 21 days without showing laccase activity. There was no much difference in the rate of decolorization of malachite green (Figure 6) viz. 81.8%, 85.1% and 89.6% in 7, 14 and 21 days respectively. From these results, it is revealed that the percentage of decolorization was increased with incubation time (Figure 7).

Production of lignolytic enzymes during the decolorization varied depending upon the incubation time (Figure 8). The highest lignolytic activity (46 U/mL) was shown in basic fuchsin followed by bromophenol blue (42 U/mL) after 21 days of incubation. 44 U/mL of lignin peroxidase was observed with methyl green followed by basic fuchsin (36 U/mL) after 21 days of incubation. The presence of both laccase and lignin peroxidase activities were observed in basic fuchsin. Although laccase, lignin peroxidase activities were not observed in methyl violet initially, 8 U/mL and 6 U/mL activities were observed respectively after 21 days of incubation. Although there was no laccase and lignin peroxidase activity in ethyl violet and methyl green, 2 U/mL lignin peroxidase (7
days) was showed in ethyl violet which was not observed thereafter. Laccase, lignin peroxidase were not released up to 7 days in malachite green but afterwards were increased with incubation time. The presence of lignolytic activity was not correlated with the decolorization in all dyes since laccase and lignin peroxidase activities were not resulted in almost all dyes. TMR assay was done using NADH co-factor (Table 2). The presence of this enzyme with decolorization all these dyes showed the specific activity of this enzyme towards decolorization. The highest activity of TMR was observed in malachite green (0.447 µmole/min/mL) followed by methyl violet (0.312 µmole/min/mL).

### 4. DISCUSSION

There are very few reports on the biological decolorization and degradation of textile and dye-stuff industrial wastes containing triphenylmethane dyes. All the reports so far available are on a small scale. Triphenylmethane dyes have been reported to undergo extensive biodegradation by ligninolytic cultures of *P. chrysosporium* [23, 24]. The structural genes encoding lignin peroxidase and laccase have been cloned and characterized [25,26]. The biochemical mechanism underlying the decolorization of triphenylmethane dyes has been elucidated in fungi [11]. Several species of fungi are known to decolorize triphenylmethane dyes by a variety of enzymes as laccase [27] lignin peroxidase [24,28]. After 18 days *Lentinula* (*Lentinus*) *edodes* completely decolorized methyl green, methyl violet, ethyl violet [29].

Ollika et al. [30] purified iso-enzymes of lignin peroxidases from *P. chrysosporium* and studied the decolorization efficiencies for several dyes including azo and triphenylmethane dyes. Synthetic dyes crystal violet and

### Table 2. Details of triphenyl methane dyes and TM reductase assay of *Fomitopsis feei*.

<table>
<thead>
<tr>
<th>Dye</th>
<th>λ&lt;sub&gt;max&lt;/sub&gt;</th>
<th>Empirical formula</th>
<th>M.W.</th>
<th>TMR assay (µmole/min/mL)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Basic fuchsin</td>
<td>544</td>
<td>C&lt;sub&gt;20&lt;/sub&gt;H&lt;sub&gt;20&lt;/sub&gt;CIN&lt;sub&gt;3&lt;/sub&gt;</td>
<td>337.85</td>
<td>0.032 ± 0.01</td>
</tr>
<tr>
<td>Methyl violet</td>
<td>585</td>
<td>C&lt;sub&gt;25&lt;/sub&gt;H&lt;sub&gt;30&lt;/sub&gt;CIN&lt;sub&gt;3&lt;/sub&gt;</td>
<td>407.99</td>
<td>0.312 ± 0.00</td>
</tr>
<tr>
<td>Ethyl violet</td>
<td>596</td>
<td>C&lt;sub&gt;31&lt;/sub&gt;H&lt;sub&gt;42&lt;/sub&gt;CIN&lt;sub&gt;3&lt;/sub&gt;</td>
<td>492.14</td>
<td>0.258 ± 0.03</td>
</tr>
<tr>
<td>Malachite green</td>
<td>620</td>
<td>C&lt;sub&gt;23&lt;/sub&gt;H&lt;sub&gt;25&lt;/sub&gt;N&lt;sub&gt;2&lt;/sub&gt;Cl</td>
<td>364.92</td>
<td>0.447 ± 0.02</td>
</tr>
<tr>
<td>Bromophenol blue</td>
<td>423</td>
<td>C&lt;sub&gt;19&lt;/sub&gt;H&lt;sub&gt;10&lt;/sub&gt;Br&lt;sub&gt;4&lt;/sub&gt;O&lt;sub&gt;5&lt;/sub&gt;S</td>
<td>669.96</td>
<td>0.228 ± 0.00</td>
</tr>
<tr>
<td>Methyl green</td>
<td>653</td>
<td>C&lt;sub&gt;27&lt;/sub&gt;H&lt;sub&gt;35&lt;/sub&gt;N&lt;sub&gt;3&lt;/sub&gt;ClBr.ZnCl&lt;sub&gt;2&lt;/sub&gt;</td>
<td>653.2</td>
<td>0.070 ± 0.01</td>
</tr>
</tbody>
</table>

malachite green showed 100% and 90% of decolorization on the PDA media [31]. In liquid culture, *P. calyp- tratus* was able to decolorize crystal violet and malachite green during 14 days of cultivation in Kirk medium after 14 days the extents of decolorization were only 5 and 27% of the dye, respectively [32]. Decolorization of triphenylmethane dyes (crystal violet, bromophenol blue and malachite green) by three birds nest fungi—*Cyathus buleri*, *C. stercoraeus*, *C. striatus* was reported [27]. Among the three organisms, *C. buleri* was found to be the most efficient in decolorization.

Particularly, the usefulness of the brown rot fungi for decolorization of wastewater has not been probed at all. *Polyporus ostreiformis* produced lignin peroxidase along with manganese peroxidase, acid protease, a-amylase in nitrogen-limited liquid media for decolorization [33]. Studies with the brown-rot fungi *Gloeophyllum trabeum* and *Fomitopsis pinicola* on liquid media revealed no lignin peroxidase or manganese-dependent peroxidase activity, although nonspecific peroxidase activity was detected [34]. No induction in laccase, lignin peroxidase and tyrosinase activity was observed during decolorization process and significant induction in malachite green reductase (73%) activities were observed during decolorization of malachite green by *K. rosea* [35]. An enzyme that decolorizes triphenylmethane dyes is designated triphenylmethane reductase (TMR) purified from *Citrobacter* sp. strain KCTC 18061P and characterized, cloned of the gene encoding this enzyme and its heterologous expression in *Escherichia coli* was reported by [22].

From our results also it is confirmed that triphenyl methane reductase enzyme plays specific role in the de- colorization process compared to lignolytic enzymes. But lignolytic enzyme production was also supported that they might be definitely useful in the degradation process. In contrast to the previous research [18] in which methyl green was not decolorized by *F. feei*, our result showed the decolorization of methyl green upto 88.3%. In their research they reported that bromophenol blue was decolorized upto 64.8% but in our present research it was decolorized upto 96.8% after 21 days of incubation period. This difference might be in days of incubation and the concentration of dye used. Our findings could contribute to a better knowledge of the properties of *F. feei*, which has as yet not been studied in detail and its decolorization abilities could be promising for further biotechno- logical applications.

5. CONCLUSION

It may be concluded from the result that it is possible to decolorize or degrade triphenylmethane dyes by biological means. Biological treatment is the only way for ultimate controlling of pollution generated by textile and dye-stuff industries. Fungal decolorization is a promising alternative to replace or supplement present treatment processes. In conclusion, our results showed that *F. feei* is able to decolorize efficiently several synthetic dyes belonging to triphenylmethane group.

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