

Original Research Article

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## Biodegradation and Decolourization of Orange G by using *Bacillus* Sp. Isolated from Textile Effluents

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### ABSTRACT

#### Keywords

Dye, *Bacillus* Species, Orange G, Textile effluents, Biodegradation

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In many commercial industries the azo dyes are the frequently used dyes. 25 bacterial isolates were taken from the textile waste, and from this 3 bacterial sample (AK 1, AK 2, AK 3) shows the best degradation capacity to degrade Orange G. By observing all the standard morphology and biochemical tests it was observed that these bacteria were the *Bacillus* Species, and it is found that isolate AK 3 showed the maximum decolourization i.e. 96% within 48 hours. The temperature at which the maximum degradation was observed is at 30°C, the pH was set up to be normal and glucose was used as a carbon source and for the nitrogen source yeast extract was used. The enzymatic activity such as Azo Reductase, Tyrosinase, Lignin Peroxidase (LiP), Manganese Peroxidase (MnP) was observed as their role in biodegradation. This research paper is an analysis of illuminated collection of the benefit of enzymes for decolourization of azo dye and also their utilization for analysis of azo compounds present in waste water.

### Introduction

Dyes play a very important role in almost every industry. Among the various dyes used in the industry Azo dyes are the most frequently and most widely used dye. Its structure is composed of the aromatic rings that are normally bound by two or more azo groups (-N=N-). It is observed from the study that the annually world azo dye production is around 1 million tons (Pandey *et al.*, 2007) and there are almost 2500 different structure azo dyes are used today (Kumar *et al.*, 2007). It has been reported that about 10-12% of dye

which are used during the dyeing activity goes into waste water (Asad *et al.*, 2007). These dyes are directly affecting the organism present in the water and cause various problems among them. (Ozturk & Abdullah, 2006). Because of the complex aromatic molecular structure and synthetic origin these azo dyes are very difficult to degrade and form as a stable compound in the atmosphere.

Among these dyes reactive dyes form the covalent bonds as in binds to textile fibres mainly cotton by covalent bonds. (O' Mahony *et al.*, 2002). For dyeing the cellulosic fibres

the most effective method is reactive dyeing. The two main component of azo dye is the presence of azo bands and the Sulfonated group which we called the xenobiotic compound (Rieger *et al.*, 2002). The treatment of this azo dye is difficult. But now a days there are various methods for treatment of these dyes. The biological method for the treatment of dye is most effective and eco-friendly as compared to other physico-chemical method.

Various microbial stains are reported by the different researchers which are able to degrade the dye. (Rajaguru *et al.*, 2000; Stolz, 2001). The breakage of azo dyes forms the aromatic amines which are known as mutagens. The bacterial degradation of dyes is much faster than the fungal degradation of textiledye stuffs (Pourbabee *et al.*, 2005). Decolourization of dye solution by bacteria can be due to biosorption or biodegradation. The bacterial activity is mostly biodegradation. However, in some cases, bacterial decolorization under aerobic conditions usually results in adsorption of dye stuffs on bacteria rather than their oxidation (Pourbabee *et al.*, 2005). The adsorption of dye stuffs is rare in bacteria than in fungi as fungal cell wall is rich in chitin, in which hydroxyl and amino groups are present which makes it an efficient adsorbent for dye effluents. (Chen *et al.*, 2003) found that dye biosorption or biodegradation can be judged by the colour of cell mat. If the cell mat gets deeply coloured by the dye then the mechanism of dye removal is biosorption but if cell mat retains its original color then phenomenon of biodegradation can be attributed for dye removal.

The biodegradation of azo dyes involve 2 steps, first being the cleavage of azo dye and second being mineralization of intermediates. It is currently accepted that azo dye reduction is due to co-metabolic reaction, in which the

biologically formed reducing equivalents can bechemically transferred to azo dyes.

Bacterial degradation of azo dyes is often mediated by azoreductase. Sulfonated azo dyes are not decolorized easily, as the permeation through the cell membrane is the rate limiting step during bacterial reduction of azo dyes. Such decolorization occurs by an oxygen sensitive azo reductase (Kodam *et al.*, 2005).

This study had reported the species which are able to decolorize the Orange G dye in aerobic condition.

## **Materials and Methods**

### **Sampling sites and Textile dyes used**

The sample was collected from the textile industry located in Gaya, Bihar, India. Orange G which is a type of azo dye which is used in this study. Stock solution was prepared by dissolving 100 ppm dye sample in 100 ml of distilled water.

### **Isolation and Screening of Bacterial Strains Decolorizing Azo dye**

The effluent and sludge samples were serially diluted and spread over minimal agar medium containing 100 ppm of Orange G. pH was adjusted to 7.0 before autoclaving and incubated at 37°C for 5 days. Colonies surrounded by halo (decolorized) zones were picked and streaked on minimal agar medium containing azo dye. The pure cultures were maintained on dye-containing nutrient agar slants at 4°C.

### **Decolourization assay**

Loopful of bacterial culture was inoculated in Erlenmeyer flask containing 100 ml of nutrient broth and incubated at 150 rpm at

30°C for 24 h. Then, 1 ml of 24 h old culture of the bacterial isolates were inoculated in 100 ml of nutrient broth containing 100 ppm of Orange G and re-incubated at 30°C till complete decolourization occurs. Suitable control without any inoculum was also run along with experimental flasks. 1 ml of sample was withdrawn every 24 h and centrifuged at 10,000 rpm for 15 min.

Decolourization extent was determined by measuring the absorbance of the culture supernatant at 475nm using UV-visible spectrophotometer, according to Hemapriya *et al.*, (2010).

Decolourization efficiency (%) =  $\frac{\text{Dye (i)} - \text{Dye (r)}}{\text{Dye (i)}} * 100$

Where,

Dye (i) refers to the initial dye concentration and Dye (r) refers to the residual dye concentration.

Decolourization experiments were performed in triplicates.

### **Optimization of culture conditions for dye decolourization by *Bacillus sp.***

#### **Effect of Temperature, pH and Dye Concentration**

The effect of temperature, pH and dyeconcentration on dye decolorizing ability of the isolate was studied. This was carried out by incubating the bacterial strains at different temperature (25-45°C), pH (5-9) and various dye concentrations (100-500 ppm).

#### **Effect of Carbon and Nitrogen source on Dye Decolourization**

To investigate the effect of various carbon and nitrogen sources, different carbon sources

such as, glucose, lactose, and sucrose (1%) and different nitrogen sources like yeast extract, beef extract, and peptone (1%) were added as a supplement individually to Nutrient broth medium for decolourization of Orange G.

#### **Enzyme assays**

Assay was carried out in cuvettes with a total volume of 1 ml. One unit per enzyme activity was defined as the amount of enzyme that transformed 1µ mol of substrate per minute (1 unit = 1U). Preparation of CellFreeExtract The bacterial strain AK3 was inoculated in Nutrient Broth containing Azodye (Orange G) and incubated at 30 °C. The cells were harvested by centrifugation at 7000 rpm for 30 min in cooling centrifuge, washed with 50 mM phosphate buffer (pH 7.0) and re-suspended in the same buffer. Then, the cells were disturbed and cell debris was removed by centrifugation at 4 °C. The resultant supernatant was used as the source of crude protein / enzyme.

#### **Laccase activity**

Assay Laccase activity was determined using 2,2'- azino-di- (-3-ethylbenzo-thiazoline-6-sulfonic acid) (ABTS) as the substrate. 5µl of 50 mM citrate buffer (pH 4.0) was mixed with 430µl of distilled water and 20µl of laccase. The reaction was started by addition of 50µl of 6 mM ABTS and increase in absorbance at 547 nm was monitored. The enzyme activity was calculated using an extinction coefficient of ABTS of  $\epsilon_{436} = 36 \text{ m mol}^{-1} \text{ cm}^{-1}$  (Ander and Messner, 1998).

#### **Tyrosinase assay**

Tyrosinase activity was determined in reaction mixture of 2 ml containing 500 µl of 0.01% catechol in 500 µl of 0.1 M phosphate buffer (pH 7.4) and 1ml of cell free culture at 475 nm (Zhang and Flurkey, 1997).

### **Lignin peroxidase (LiP) assay**

LiP (Lignin Peroxidase) activity was determined by monitoring the formation of propanaldehyde at 475 nm in a reaction mixture of 2.5 ml containing 500 µl of 100 mM n-propanol, 500 µl of 250 mM tartaric acid, 500 µl of 10 mM H<sub>2</sub>O<sub>2</sub> and 1 ml of cell free culture (Shanmugam *et al.*, 1999) at 475 nm.

### **MnP (Manganese Peroxidase) assay**

The reaction mixture contained 500 µl of 50 mM sodium malonate buffer (pH 4.5), 25µl of 20 mM MnCl<sub>2</sub> solution, 415µl of distilled water and 50 µl pf MnP. The reaction was started by adding 20 µl of 10 mM H<sub>2</sub>O<sub>2</sub>. The extinction of the solution was measured photometrically at the wavelength 475 nm ( $\epsilon_{270}= 11.59 \text{ mmol}^{-1} \text{ cm}^{-1}$ ) (Wariishi *et al.*, 1992).

### **Azo reductase assay**

Assay was carried out in cuvettes with a total volume of 1 ml using colorimeter. The reaction mixture consists of 400 µl of potassium phosphate buffer with 200 µl of sample and 200 µl of reactive dyes (500 mg/l). The reaction was started by addition of 200µl of NADH (7mg/ml) and was monitored photometrically at 547 nm. The linear decrease of absorption was used to calculate the azoreductase activity. One unit of azoreductase can be defined as the amount of enzyme required to decolorize 1µ mol of Orange Gperminute.

### **High Performance Liquid Chromatography (HPLC)**

This method was carried out according to (Sahasrabudhe *et al.*, 2014) with some alteration. The removal of Orange G azo dye and a generation of decolourization products

were monitored using an HPLC system equipped with an ODS C18 column (Shimadzu SPD 20A) with HPLC grade methanol 60 % + 40 % deionized water as mobilephase at the flow rate of 0.6 mL/min for 10 min at 265 nm. A portion of the sample (10 microliters) was manually injected into the injector port, and then the metabolites were analysed using a dual absorbance UV detector.

### **Results and Discussion**

Isolation, Screening and Identification of bacterial strains decolorizing textile dyes the results shown in Table.1 revealed that 03 bacterial isolates, designated as AK1 to AK3 were found to becapable of decolorizing Orange G (Fig. 1).

Out of 03 isolates, AK3 was found to be the superior strain with the highest decolourization efficiency (96.60 %).

Morphological, cultural and biochemical characteristics of AK3 strains were tabulated in Table.2. On the basis of the above mentioned characteristic features and by the comparison with “Bergey’s manual of Determinative Bacteriology”, the isolate AK3 was identified as *Bacillus* sp. Strain AK3 (fig.2).

The extent of dye decolourization of Orange G by the bacterial isolates (AK1 to AK3) is shown in Fig. 3.

### **Optimization of Dye Decolorizing Ability of AK3 Isolate effect of Incubation time**

Dye decolourization by *Bacillus* sp. Strain AK3 was found to be growth dependent, since considerable dye decolourization was noticed in the fermentation broth as soon as the bacterial strains entered the late exponential phase and the activity reached the maximum

level in stationary phase after 72 h of incubation (Fig. 4).

### Effect of Temperature

The influence of incubation temperature on the decolourization of Orange G by *Bacillus sp.* AK3 was studied at temperatures ranging from 25-45 °C. The color removal efficiency of the bacterial isolate (AK3) achieved highest levels (95.02 %) at 35°C, after 72 h of incubation. However, incubation at temperatures below 30°C and above 40°C

was found to be down regulating the decolourization percentage of the isolate (Fig. 5).

### Effect of pH

Dye decolourization efficiency of *Bacillus sp.* AK3 against Orange G was detected over a broad range of pH (5.0-9.0), with optimum decolourization of (94.54 %) being exhibited at neutral pH (7.0). At slightly alkaline pH (8.0), decolourization efficiency of the isolate was found to be effective (82.54%) (Fig. 6).

**Table.1** Bacterial Strains Decolorizing

S.No	Isolates	% of Decolourization
1	AK1	87.66
2	AK2	80.64
3	AK3	96.60

**Table.2** Morphological, Cultural and Biochemical Characteristics of *Bacillus sp.*

S.No.	Test characteristics	Bacterial Isolate (AK3)
<b>I.</b>	<b>Morphological characteristics</b>	
1.	Colony morphology	Round
2.	Cell morphology	Rod shape
3.	Gram reaction	Positive
4.	Motility	Motile
<b>II.</b>	<b>Physiological characteristics</b>	
5.	Growth under aerobic condition	+
6.	Growth under anaerobic condition	+
7.	Growth in Liquid medium	+
<b>III.</b>	<b>Biochemical characteristics</b>	
8.	Indole Test	-
9.	Methyl Red test	-
10.	Voges Proskauer's	+
11.	Citrate utilization	-
12.	Glucose	-
13.	Adonitol	-
14.	Arabinose	-
15.	Lactose	-
16.	Sorbitol	-
17.	Mannitol	-
18.	Rhamnose	-
19.	Sucrose	-

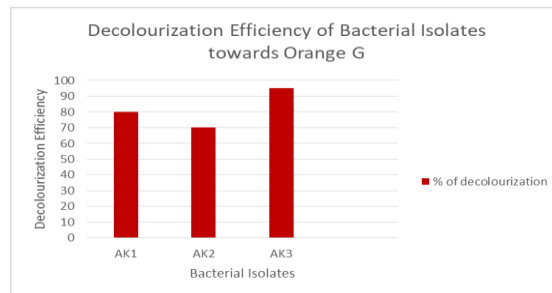
**Fig.1** Orange G before and after decolorization in Nutrient Broth



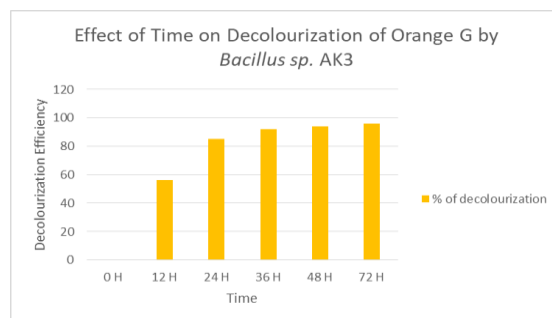
**Fig.2** Biochemical Characteristics of *Bacillus* sp.



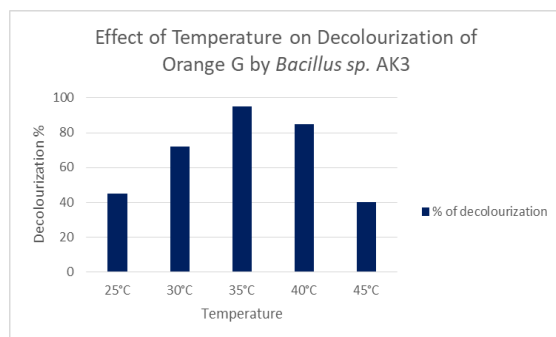
**Fig.3** Decolourization efficiency of bacterial isolates towards Orange G



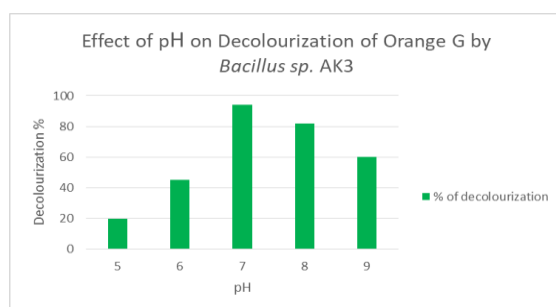
**Fig.4** Effect of Incubation Time on Decolourization of Orange G by *Bacillus* sp. AK3



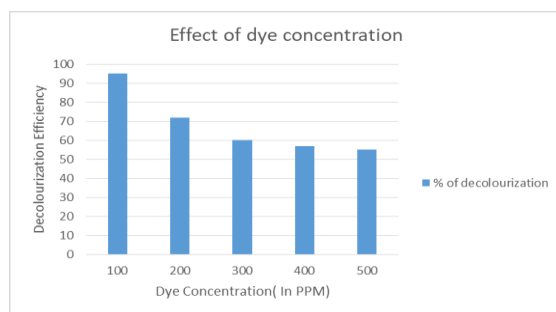
**Fig.5** Effect of Temperature on Decolourization of Orange G by *Bacillus sp. AK3*



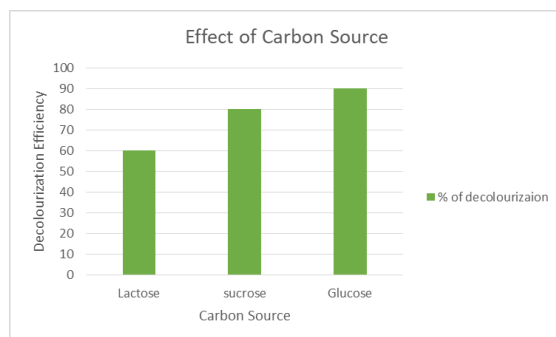
**Fig.6** Effect of pH on Decolourization of Orange G by *Bacillus sp. AK3*



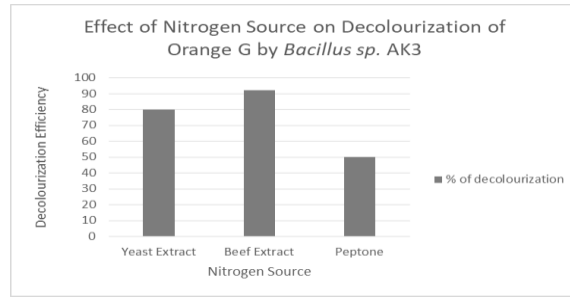
**Fig.7** Effect of Dye Concentration on Decolourization of Orange G by *Bacillus sp. AK3*



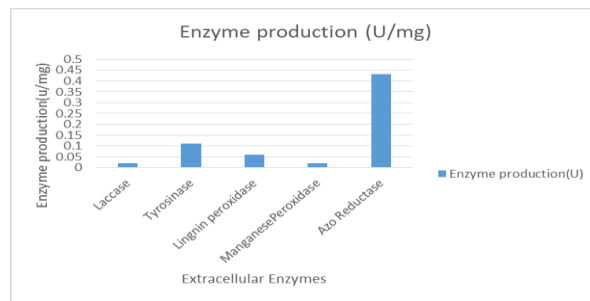
**Fig.8** Effect of Carbon Sources on Decolourization of Orange G by *Bacillus sp. AK3*



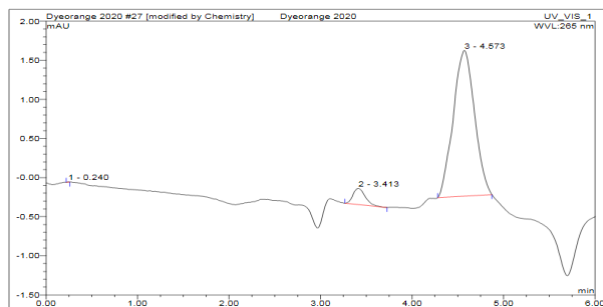
**Fig.9** Effect of Nitrogen Sources on Decolourization of Orange G by *Bacillus sp.* AK3



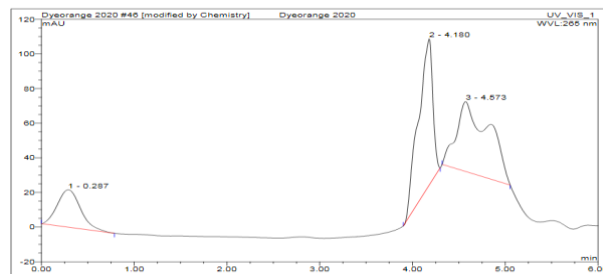
**Fig.10** Extracellular Decolorizing Enzyme production by *Bacillus sp.* AK3



**Fig.11 (A)** HPLC profile of 100 PPM Orange G and **(B)** HPLC profile of Orange G treated by *Bacillus sp.*, showing dye metabolites after decolourization



Control



After Decolourization



### Effect of Dye Concentration

The results revealed that the decolourization rate of the isolates was optimized in the presence of initial dye concentration of 100 ppm (Fig. 7). As the dye concentration increased in the culture medium, a gradual and directly proportional decline in color removal was attained. At high concentration (500 ppm), Orange G greatly suppressed decolourization ability of *Bacillus sp.* AK3.

### Effect of Carbon and Nitrogen Sources

The bacterial isolate AK3 was able to utilize most of the carbon sources tested, whereas glucose instigated maximum decolourization efficiency (90.93 %) (Fig. 8). Among the various nitrogen sources tested, yeast extract was found to be the superior source in maximizing decolorizing ability (92.90%) (Fig. 9).

### Enzymatic assay for decolourization of Orange G

The culture supernatant of AK3 cells that mediated the decolourization of Orange G was screened for the presence of dye decolorizing enzymes such as Azoreductase, Laccase, Tyrosinase, Lignin Peroxidase, MnP (Manganese peroxidase). Azoreductase was found to be the dominant enzyme (0.43 U mg<sup>-1</sup>protein), whereas Laccase, Tyrosinase, Lignin peroxidase, MnP (Manganese Peroxidase) were found to be secreted in very trace amounts (0.02, 0.11, 0.06, 0.02 U mg<sup>-1</sup>protein respectively) (Fig. 10).

### High Performance Liquid Chromatography (HPLC)

Fig.11 (A) HPLC profile of 100 PPM Orange G and (B) HPLC profile of Orange G treated by *Bacillus sp.*, showing dye metabolites after decolourization

After 5 days of aerobic treatment process, HPLC analysis for biodegradation products was carried out. HPLC analysis of Orange G as a control showed a peak at the retention time of 4.573 min. (Figure 11A), After the decolourization the *Bacillus sp.* showed two metabolites peaks at the retention times of 4.180 and 4.573 min. (Figure 11B). All samples showed the appearance of dye peak in the metabolites along with appearance of new peaks with different retention times which supported dye biotransformation to different metabolites.

In conclusion the environmental biotechnology is constantly enlarging its efforts in the biological analysis of coloured textile discharge, which is an environmental friendly and low-cost substitute to physico-chemical decomposition method. The textile industries are chemical utilizing concerns, of which various dyes are of importance. Assembly of dye stuff creates environmental pollution and also health and aesthetic trouble. With the advent of civilization, the application of dye has become unavoidable and its release in the environment poses a severe problem. Therefore, an urgent need for technically practicable and cost-effective treatment method to be devised to overcome this problem is the need of hour. Microbial decolourization and degradation of various azo dyes have outstanding prospective to address various problem due to their environmental friendly and economical features. In addition to these bacteria have various other properties like growth rate is fast, hydraulic retention is high and therefore they are productive for the treatment of high strength waste waters. The literatures analysed in this research paper mainly focused on various experiments which have been carry out on decolourization of azo dye by the use of pure bacterial culture. In this study we have seen that, under aerobic state the subtractive division of these azo bonds

form the toxic aromatic amines compound, so it is still very necessary to evaluate the efficient aromatic amine bacterial degraders. Since various other parameters also affect the performance of microorganisms and therefore, optimization of these is very necessary. To conquer the practicable process of dye decolorization, amplify knowledge of enzymatic study and comprehensive characterization of the intermediary and metabolites which are produced during the process of biodegradation using different scientific techniques. Further, to guarantee the safety of environment and to decolorize dye effluent, studies must be organized on the toxicity due to untreated effluent dye solution. Temperature variation had a significant effect on the decolorization of Orange G by *Bacillus sp.* strain AK3. The rate of decolorization was found to be optimized at 35°C after 72 h of incubation. The rate of decolorization decreased with the decrease in temperature. This fact implies that the local temperature in the microenvironment of the effluent samples has a very significant effect on the decolorization activity (Moosvi *et al.*, 2005). Decolorization activity of *Bacillus sp.* strain AK3 was significantly suppressed at temperatures more than 40°C, which might be due to the loss of cell viability or denaturation of the enzymes responsible for the decolorization at elevated temperatures. The most biologically feasible pH for the decolorization of Orange G by *Bacillus sp.* strain AK3 was found to be 7.0.

The foregoing results suggest the potential of utilizing *Bacillus sp.* strain AK3 to degrade textile effluent containing synthetic textile dyes via; appropriate bioreactor operation.

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