

Short communication

Screening and characterization of bacteria for decolorizing of Azo Dye

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Azo dye is a mainly pollution, the concentration of azo dye from the waste water is very high, it is difficult to decolour and degrade azo dye in nature. Bacteria have the ability to decolour and degrade, it can be biosorption the dye. Four strains were screened and identified by the morphology of colonies and individual observation, Gram staining, flagella staining, spores formation experiment, physiological and biochemical reactions, these are belong to *Bacillus*, *kurlhia*, *Klebsiella*, the decolorization rate of four strains was more than 70%.

Key words: screening; characterization; azo dye; decolorizing

INTRODUCTION

It had been reported on many papers about bacteria on biosorption, decoloration and degradation of dyes. *Bacillus subtilis* could degrade azo dye. Now many decoloration bacteria were separated, which could use dyes as sole carbon sources. Some newly synthesized dyes developed to the direction of anti photolysis and oxidation resistance (Yang et al., 2002), which made the dye wastewater treatment more difficult. Decolorization bacteria were screened from the wastewater of printing and a dyeing factory and preliminary to identify them in our research (Khan et al., 2014)

The report on dye decolorization bacteria showed that the bacteria with a variety of dyes decolorization ability widely existed in nature, such as: *Klebsiella*, *Rhodospirillum rubrum*, *Acetobacter*, *Pseudomonas*, *Erwinia*, *Kurtzia*, *Alcaligenes*, *Escherichia*, *Plesiomonas*, *Bacillus*, *Enterobacter*, *Serratia*, *Zoogloea*, *Xanthomonas* et al., (Bazli et al., 2015) (Jaouadi et al., 2014).

MATERIALS AND METHODS

The source of water sample

A water sample was taken aeration tanks of printing and a dyeing factory in Bin Zhou City.

Dyestuff

Reactive red, reactive red X-3B, reactive brilliant orange X-GN, acid black NG, preparation of dye concentration of 10g/L solution, with 0.22um microporous membrane filtration, deposits.

The main medium

Enrichment medium

Peptone 2.0, beef extract 2.0, glucose 5.0, KH₂PO₄ 2.0, Na HPO₄ 12 H₂O 3.5, MgSO₄ 7H₂O 0.1, dyestuff 0.01-

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Figure1: The microscopic morphology of the four strains (from left to right ZB11, ZD17, ZG15, ZH23)

0.05, 1000mL water, pH 7.2.

Beef-protein medium

Beef extract 5.0, peptone 10.0, NaCl 5, 1000mL water, pH 7.2.

Screen dyestuff medium

Addition a certain amount of dye to beef-protein media in the culture medium.

Main instrument and equipment

Spectrophotometer, centrifugal machine, incubator shaker.

Water sample collection

Water samples were collected in sterile glass bottles, which were from different parts of the aeration tank, inlet, outlet in sewage treatment plant of BinZhou city.

Screening and culture

5mL the waste water addition to containing 50mL enriched culture fluid of 250mL the triangle bottle, under the 37°C 120rpm oscillation culture, every five day the 10mL culture solution was added to the 50mL fresh medium (Strittmatter et al., 2015).

Isolation and purification of resistant bacteria

Culture solution to screening and culture was diluted, coating into containing the same concentration of dye, solid plate and 37°C cultivation for a certain period of time, through comparison of colony morphology, texture, color, size, etc.,. Different strains were separated, purified and inoculated in beef extract peptone slant culture medium (Linde et al., 2014).

Screening of decolorizing bacteria

The isolated strains inoculated in the liquid dye medium (dye concentration of 30mg / L, each test tube with 5 ml

medium, each strains inoculation three tubes), the static culture for a certain period of time, 8000rpm centrifugation for 30 minutes, the supernatant in the dye maximum absorption peak was measured the OD value with spectrophotometry, dye medium without inoculation bacteria as the control, calculating the decolorizing rate, showing the ability of decolorizing of dye (Singh et al., 2015).

decolorizing rate equation = $(A-B)/100\%$

A: the OD value of without inoculation

B: the OD value of with inoculation

Identification of strains

With enrichment culture and decolorizing ability test, efficient decolorizing strains were tested by the colony morphology and individual morphological observation, Gram staining, flagella staining, spores staining et al., , then doing physiological and biochemical experiments, identification the separation of decolorizing strains.

RESULTS

Decolorization ability of different strains to the test

With the enrichment and screening and the decolorizing test, four strains (Figure 1) have the high decolorizing ability, the result showed on the table 1 As it could be seen from the table 1, the strains of bacteria of dye decolorizing ability were very strong, the decolorizing rate was more than 70%. The decolorizing ability of different strains was different, and the decolorizing rate of ZG15 strains was more than 92%.

Preliminary identification of decolorizing strains

With enrichment culture and decolorizing ability test, four strains of bacteria with strong decolorizing capacity were selected, with the method of the morphology of colony and individual observation, Gram staining, flagella staining, spores formation experiment, further physiological and biochemical reactions, including acid

Table 1: Different strains decolorizing rate of different dyes

Strain number	Dye name	The decolorizing rate %
ZB11	Reactive red	73
ZD17	Reactive red X-3B	85
ZG15	Reactive brilliant orange X-GN	92
ZH23	acid black NG	85

Table 2: The result of identification of the bacteria

Strain number	Genus
ZB11	Bacillus
ZD17	Kurlhia
ZG15	Bacillus
ZH23	Klebsiella

Table 3: the colony morphology, individual character, physiological and biochemical traits of decolorizing bacterium

Strain number	Colony morphology	Individual form	Flagella staining	Gram staining	Spores staining	Acid fast staining
ZB11	round or irregular uplift opaque	straight rod 0.5-1.0×1.2-3.0um	peritrichous flagella	+	+	-
ZD17	Colony uplift, irregular edge	0.5-0.8×3-4.5um	peritrichous flagella	+	-	-
ZG15	round or irregular uplift opaque	straight rod 0.5-1.0×1.2-3.0um	peritrichous flagella	+	+	-
ZH23	colony close to circle, protruding bright	straight rod 0.5-1.0×0.5-5.0um	without	-	-	-

Table 4: The physiological and biochemical traits of decolorizing bacterium

Strain number	oxidation glucose fermentation	contact enzyme reaction	oxidase enzyme	phenylalanine deaminase	potassium cyanide (KCN)	inositol fermentation
ZB11	+	+	-	-	-	+
ZD17	-	+	-	-	-	-
ZG15	+	+	-	-	-	+
ZH23	+	-	-	-	+	+

+Positive reaction, -Negative reaction

fast staining, oxidation of glucose fermentation, contact enzyme reaction, oxidase test, phenylalanine deaminase, potassium cyanide (KCN) experiment, inositol fermentation. The results showed that the four strains were classified into the following table 2, table 3, table 4. Compared with the 16S rDNA (Figure 2). sequence of other bacterial strains in GenBank, the results of molecular biology identification validated the identification of the four strains of bacteria by morphological and physiological biochemical methods.

In the process of screening decolorizing bacteria, we found that the enrichment and cultivation of the dye gradient acclimation were very important. If there was not with the enrichment culture, because the decolorizing bacteria were low concentration, it was likely to difficult to

be separated for the high concentration of non degrading bacteria (Rekik et al., 2015) (Silva et al., 2009).

DISCUSSION

The result showed that the concentration of dye in the nutrient fluid reduced, because of the treatment by the bacteria. During treatment there were some factors which could reduce the concentration of dye, one was the biosorption, the dye was adsorbed the cell surface of the bacteria, the dissolved dye in the waste water and nutrient fluid would reduce (Hajivandi et al., 2016). The second factor was the degradation of the bacteria, which could use the dye as the nutriment, sustaining the cell

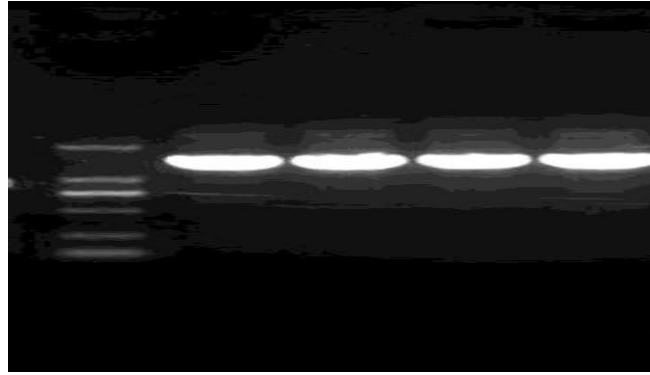


Figure 2: the 16S rDNA of four strains (from left to right ZB11, ZD17, ZG15, ZH23)

growth (Huang et al., 2015). The biosorption of treatment of the bacteria was not complete to eliminate dye; the degradation of treatment of the bacteria was complete to eliminate dye (Kadam et al., 2015), to some extent the degradation method was better, but the degradation process may produce some small molecules, which may take new pollution and hazard, we should research on the degradation new products (Patil et al., 2016), it would help reduce the harm caused by the degradation new products.

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