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RESEARCH ARTICLE

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Characterization of Autochthonous Bacteria Capable for Degradation of Residual Organic Pollutants of Pulp Paper Mill Effluent by Biostimulation Process

Pooja Sharma¹, Sonam Tripathi¹, Preeti Chaturvedi² and Ram Chandra¹*

¹Department of Environmental Microbiology, School for Environmental Sciences, Babasaheb Bhimrao Ambedkar Central University, Vidya Vihar, Rae Bareli Road, Lucknow - 226 025, India. ²Aquatic Toxicology Laboratory, Environmental Toxicology Group, Council of Scientific and Industrial Research-Indian Institute of Toxicology Research (CSIR-IITR), Vishvigyan Bhavan, 31, M.G. Marg, Lucknow - 226 001, India.

Abstract

The purpose of this study is remediation of residual organic pollutants from effluent by autochthonous bacterial community in biostimulation process. Discharged effluent showed high TDS (549 mg L-1), TSS (59 mg L-1), COD (20349 mg L-1) and BOD (25946 mg L-1), value. The level of total phenol (421 mg L-1), nitrogen (156 mg L⁻¹), sulphate (1854 mg L⁻¹), phosphorus (176 mg L⁻¹) chlorine (2.01 mg L⁻¹), sodium (75 mg L⁻¹) and potassium (8.4 mg L⁻¹) along with various heavy metals (Fe, 75.23; Zn, 15.60; Cu, 4.1; Cr, 3.12; Cd, 0.324; Mn, 13.24; and Ni, 4.01 mg L-1) were noted above the permissible limit of Environmental Protection Act. The result revealed that the reduction of the physico-chemical parameter of pollutants were above 50% after biostimulation process, this confirmed the potentiality of growing autochthonous bacterial community responsible for bioremediation. The comparative UV-Vis spectroscopy showed reduction in the absorption spectra of degraded sample. Further, GC-MS analysis showed major organic pollutants i.e. Octadecanoic acid, Hexadecanoic acid, citral, benzoic acid, and 2, 6'-Di-hydroxy acetophenone, bis (trimethylsilyl) ether, were detected in control few compounds were degraded while there was formation of some new metabolic products also. Few pollutants persisted in the degraded sample as recalcitrant toxicant and causes environmental toxicity and hormonal imbalance as endocrine-disrupting chemicals (EDCs). But the detailed knowledge and characterization of organic pollutants are not available yet regarding their properties. The SEM image showed the diversity of bacterial community in biostimulation responsible for utilisation of various detected compounds. The growing bacterial communities were identified as potential bacterial strains as Aeromonas salmonicida, BBAUPS-1 (MN294457.1) and Bacillus sp. BBAUPS-2 (MN238724.1) responsible for the remediation of residual organic pollutants. Further, Evaluation of toxicity parameter of effluent by seed germination test of Triticum aestivum and Cicer arietinum inhibited the seed germination upto 80%. Hence, this study revealed that the biostimulation process is a good technique for detoxification and degradation effluent.

Keywords: Environmental toxicity, endocrine-disrupting chemicals, organic pollutants, phytotoxicity, effluent detoxification

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*Correspondence: prof.chandrabbau@gmail.com

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INTRODUCTION

The discharged effluent from the pulp paper industry contains more than 700 organic and inorganic pollutants that are directly responsible to cause the soil and water pollution after biological and chemical treatment¹. The pulp paper industry ranks 6th among the world's most polluting industries and generate large-scale hazardous pollutants during the making of papers². Pulp paper industry effluents contain high number of complex matrices along with many other diverse compounds³. Moreover, the effluents may contain more than 250 identified chemicals and harmful components like sterols and asresin acids^{4,5}. These compounds may end up as sediments in the water bodies and serve as food for fish and benthos⁶. Although, many harmful substances are removed before release by modern wastewater purification processes but some are released unintentionally into the environment. Moreover, untreated effluent with heavy nutrient load and wood derived bioactive substances if released into the natural water bodies may cause oxygen depletion and subsequent decline in the biodiversity⁷.

In India, there are approximately 650 small and large scale paper industries discharging approximately 190-200 m³ effluent per ton of paper production as aquatic pollutants8. Furthermore, the recent study has also revealed that some of these compounds showed EDCs effects on aquatic organisms. Moreover, there is still a lack of detailed knowledge on the estrogenic and androgenic compounds from pulp paper industry effluent and their toxicity in the aquatic organism. Therefore, optimizing bacterial growth conditions by adding different nutrients and providing good environmental conditions might be an effective strategy for detoxification. However, there is lack of understanding on the controlling factors for biodegradation of complex organic pollutants derived from several polluted sites. In-situ bioremediation has been introduced as an cost effective and less tedious approach in redeveloping any contaminated site. Additionally, the identity of the microbial community and required environmental conditions should be detected before specification for bioremediation function. Three types of bioremediation processes are generally used for the in-situ bioremediation process of any complex industrial contaminants viz. natural attenuation, bio-stimulation and bioaugmentation.

Bio-stimulation process involves adaptation to the contaminated site to provide a favourable environment for bacterial communities for effective natural degradation of different pollutants⁹. Several persistent organic pollutants (POPs) are present in industrial wastes which are known to be silent environmental killers due to their bio accumulative and longlasting existence^{10, 11}. Furthermore, isolated bacterial strains from compost soil i.e. Azotobacter sp. and Serratia marcescens might degrade and decolorize lignin-containing effluents while Bacillus subtilis and Bacillus sp. are responsible for kraft-lignin degradation¹². The quality of water is highly detoriated by surpassing TDS, TSS, BOD and COD values. Moreover, the dark colour and high turbidity of effluent due to suspended solids, TSS, TDS and TS can further cause river pollution and compromise the drinking water quality. The pulp paper industry effluent absorbs more light and reduces the oxygen concentration in water thereby affecting the aquatic life due to presence of tannins and resin acids13. While the change in the colour and water quality nearby the industrial area makes it unsafe for drinking purpose. The leaching property of different effluent pollutants is a major source of ground pollution. Pollution of our environment by heavy metals particles is a very dangerous and challenging problem for the country^{14,15}. Earlier, three potential bacterial strains viz. Panibacillus sp. Aneurinibacillus aneurinilyticus and Bacillus sp. were identified and characterized from sludge for synthetic lignin degradation and metabolic properties using gas chromatographymass spectrometry (GC-MS) analysis respectively¹⁶. Similarly, the Bacillus sp. and Serratia marcescens were also reported to cause degradation of the pentachlorophenol containing effluent in the presence of about 94% nutrients in optimized in-vitro conditions. Thus, these studies report the degradation and detoxification capability of bacteria for chlorolignins containing effluent¹⁷. The removal of heavy metals from contaminated water and soil by employing electro-coagulation for the waste management methods has been previously described^{18,19}. Therefore, the present study was designed to focus on the characterization and identification of residual organic pollutants and their phytotoxicity for ecological risk assessment. However, during the biostimulation process, the role of microbes is very interesting and important for the remediation of these pollutants. Furthermore, 16s rRNA sequencing of two potential bacterial strains isolated during the biostimulation process is important. This will lead to develop in-situ bioremediation technology for the removal of color and reduce the toxicity in the effluent to prevent pollution prevention. The industry (K R pulp paper mill Ltd) selected for sampling, discharges huge amount of effluent (200 m³ of effluent/ton of paper production) which pollutes the environmental pollution and also deteriorates the surface water quality. Additionally, the observation reveals the leaching of pollutants to groundwater and soil which adversely affect human health, but the toxicity data is not published.

MATERIAL AND METHODS Site description and sample Collection

The fresh effluent samples were collected from the M/s K R Pulp Paper Industry, Shahjahanpur, Uttar Pradesh, India (27°50′31.8″N 79°51′15.7″E). The industry uses bagasse and bamboo woods as raw materials and generates 100 tons per day (TPD) effluents after the pulping and bleaching processes. The effluent samples were collected using a sterile plastic container and stored at 4°C until further use.

Physico-chemical analysis

The physico-chemical parameters of effluent were analysed as per standard methods described for the examination of water and wastewater²⁰. The pH, TSS, TDS, TSS, electric conductivity (EC), chloride, sodium and potassium of the effluent were measured using the selective ion electrode of Thermo Orion, USA (Model 960). Lignin and the chlorophenols were estimated as per the method described by Chandra et al., (2009)²¹. While, the heavy metals i.e. Fe, Zn, Cu, Cr, Cd, Mn, and Ni were analysed using the atomic absorption spectrophotometer (AAS; ZEEnit 700, Analytic Jena, Germany)

UV- Vis spectrophotometer analysis

UV-Vis spectrophotometer (Thermo Fisher Scientific Shanghai Spectrophotometer Evolution 2001, China) was used to analyse the absorbance of the different organic and inorganic

pollutants present in the effluent. A total of 250 ml effluent sample in Erlenmeyer flask was amended with different concentrations of additional nutritional sources viz. glucose 1% (carbon) and peptone 0.5% (nitrogen, w/v), respectively. The soluble pollutants were detected by scanning the absorption spectrum in the range of 200-700 nm at room temperature.

Extraction of residual organic pollutants

Different organic solvents viz. ethyl acetate, isopropyl alcohol, n-hexane, and methanol were used to know their optimum extractability of remaining residual organic pollutants of the effluent and ethyl acetate was showed maximum solubility for the pollutants²¹. The residual organic pollutants were detected by comparing their mass spectra (m/z) with the compounds provided in the National Institute of Standards and Technology (NIST) library.

Colony-forming unit (CFU), biomass and total protein

The total microbial community during the biostimulation process was evaluated with the help of the plate count method (PCA). The sample was serially diluted and about 10 μ l of the diluted sample was used as an inoculum to spread over culture plate (Himedia, India). The inoculated plate was incubated overnight at 37°C (Chandra et al., 2018)²². The biomass estimation, 1 ml of the bacterial growth sample was centrifuged and dry in the pre-weighted eppendorf tube and after dried at 50°C, the post weight of eppendorf post weight minus the pre weight of eppendorf were estimated to be bacterial biomass per ml. The total extracellular protein of biostimulated sample was estimated using the standard method²³.

Scanning electron microscopy analysis

For scanning electron microscopy (SEM) analysis of bacterial community during the biostimulation process, the samples were centrifuged at 6000 rpm for 20 minutes and the supernatant was discarded and pellets were collected. The collected biomass was then fixed and dehydrated using 0.1M phosphate buffer (pH 7.2) solution and different concentrations of acetone for 20 minutes, respectively²⁴. Thereafter, the samples were placed on the aluminium stub and coated with platinum through a sputter coater (SC 7620 Mini Sputter Coater, Quorum Technology Ltd., UK).

Ligninolytic enzyme assay

The confirmation of degradation of lignin at different parts per million (ppm) by different ligninolytic enzymes was analysed using different substrates. Based on the oxidation of phenol red and Azure B dye in the presence of H₂O₂, the manganese peroxidase (MnP) and lignin peroxidase (LiP) were estimated and the optical density (OD) of both the enzymes was measured at 610 nm and 651 nm, respectively²⁴, ²⁵. Additionally, the activity of laccase enzyme was estimated by the presence of guaiacol at 450 nm OD and further confirmed using plate assay. The MnP, LiP and laccase enzyme assays on the plate were performed as per the standard method^{24, 26}. **Identification of bacterial strains**

The potential bacterial strains were isolated and identified as per Cowan and Steels Manual for the identification of medical bacteria²⁷. For 16S rRNA sequencing, the isolated bacterial strains were overnight incubated in Luria Bertani broth (Himedia Pvt Ltd) and the total DNA was extracted using commercial kit (real Biotech Corporation). The 16S rRNA gene was amplified using universal primers sets16Sf (5' CAGCAGCCGCGGTAATAC 3') and 16 Sr (5' TACGGCTACCTTGTTACG 3'). The PCR reaction mixture included assay buffer 5µl, forward primer 1 μl, reverse primer 1 μl, dNTP 1 μl, template 2 μl, tag polymerase 1 μl and the final total volume was made up 50 µl with milli Q water. The reagents were properly mixed and transferred to a thermo cycler (Sure Cycler 8800; Agilent Technologies, Malaysia) under the following thermal cycling conditions, denaturation at 94°C for 1 min, followed by annealing at 55°C for 1 min and extension at 72°C for 2 min, for 35 repeated cycles. The PCR amplified product of 16S rRNA gene of bacteria was confirmed by agarose gel electrophoresis. Furthermore, the PCR product in the gel was purified using QIA gel extraction kit and processed for sequencing. The sequence data were analysed by BLAST and identified based on closet similarity with the reported sequenced data. A phylogenetic tree was generated using MEGA-6.0 software²⁸.

Phytotoxicity evaluation by seed germination test with *Triticum aestivum* and *Cicer arietinum*

The phytotoxicity of the effluent with Triticum aestivum and Cicer arietinum was evaluated and analysed as per the guidelines of Organisation for Economic Co-Operation and Development²⁹ using the seed germination and root length bioassay tests. After the biostimulation process, a total of the 50 ml sample was collected and centrifuged at 8000 rpm for 20 minutes. The supernatant was collected and filtered using 0.2 µm membranes for the seed germination test. Thereafter, ten seeds each of *Triticum aestivum* and *Cicer arietinum* were placed on the three layers of filter papers (Whatman No. 1) kept in a glass petri dish (9 cm in diameter) and moisten with different concentration of the treated and untreated sample.

Statistical Analysis

All data are presented as means ±SD for triplicate samples to confirm the data variability and results from validity. All the data were subjected to Statistical analysis Tukey's test³⁰ (Ott, 1984) using the Graph Pad software (Graph Pad Software, San Diego, CA).

RESULTS AND DISCUSSION Physico-chemical parameter

The physico-chemical analysis of effluent had higher values of pH (8.6 mg L-1), TS (742 mg L⁻¹), TDS (549 mg L⁻¹), TSS (59 mg L⁻¹), COD (20349 mg L-1), BOD (25946 mg L-1), was compared to treated sample (Table 1). The alkaline pH might be due to the presence of residual content of sodium hydroxide and sodium sulfide utilized in pulping process of the industry³¹. The five days BOD/COD of the effluent were 0.21 and 0.20, respectively suggesting the decreased degradability of pollutants present in the discharged effluent. The increase in TDS value might be due to the presence of dissolved lignocellulosic particles along with fine fibres plus pith particles and adequate concentration of Na⁺ and K⁺ which all together contribute to the salinity of effluent. The higher EC value might be due to increase in salt and ions content of effluent. Moreover, lignin and chlorophenols contents were noted to be very high in effluent which might be the source of the dark color. The dark color adversely affected aquatic flora and fauna. Consequently, it causes aquatic toxicity and soil pollution³². The high concentrations of total phenols (421 mg L-1), total nitrogen (156 mg L⁻¹), sulphate (1854 mg L⁻¹), phosphorus (176 mg L⁻¹), chloride (2.01 mg

Table 1. Physico-chemical characteristics of pulp paper mill effluent and their heavy metals content. All the values are Mean±SD. (n=3), Students t test (two tailed as compared to pre-treated wastewater). ^aHighly significant at p<0.001, ^bSignificant at p<0.01, ^cLess significant at p<0.05, ^{NS}Non-significant at p>0.05

Parameters	Effluent values before bioremediation (Mean±SD)	Effluent values after bioremediation (Mean±SD)	Permissible limit (EPA 2002)
pH	8.6±0.03	6.9±0.17 ^a	5-9
TS	742±110	222±12 ^b	-
TDS	549±12.34	127±7.54 ^b	-
TSS	59±2.01	20.33±2.08 ^c	35
COD	20349±205.00	3452±50.21°	120
BOD	25946±120	6248±531°	40
Total phenols	421±11.34	346.33±28.72°	0.50
Total nitrogen	156±5.34	201.66±21.03°	143
Sulphate	1854±11.70	1311±47°	250
Phosphorus	176±7.20	156±6.45 ^b	180
Cl-	2.01±0.00	1.36±0.12°	1500
Na+	75±19.90	23.33±3.05°	200
K+	8.4±0.20	1.731±0.25°	=
Lignin	47568±13.20	1628±75 ^b	0.05
Chlorophenol	201±21.00	183±20.66°	3.0
Heavy metals (mg L-1)			
Fe	75.23±1.00	1.89±0.10 ^b	2.00
Zn	15.60±0.30	0.95±0.17°	2.00
Cu	4.1±0.05	0.32±0.09 ^{NS}	0.50
Cr	3.12±0.05	0.61±0.11 ^c	0.05
Cd	0.324±0.01	0.05±0.02 ^{NS}	0.01
Mn	13.24±0.00	0.55±0.09°	0.20
Ni	4.01±0.01	0.33±0.07°	0.10

L-1), sodium (75 mg L-1), and potassium (8.4 mg L-1), were present in effluents which were above the permissible limit (Table 1). High concentration of chloride in the effluent generally causes more toxicity than sulphate to aquatic flora and fauna including microbial community. Furthermore, a significant amounts of Fe (75.23 mg L-1), Zn (15.60 mg L⁻¹), Cu (4.1 mg L⁻¹), Cr (3.12 mg L⁻¹), Cd (0.324- mg L-1), Mn (13.24 mg L-1) and Ni (4.01 mg L-1), were present in effluent which are hazardous to the environment. Thus, the pulp paper industry effluent are rendered unfit for irrigation and drinking purpose due to its adverse effect on the respiratory system, alimentary canal, nervous system, coronary system and reproductive system in the aquatic organism33. A similar observation was reported by earlier findings 34, 13. The source of heavy metals in effluent might be due to the alkaline black liquor corrosion activity produced during digestion of wood and inefficient

utilization of the organic contents by the bacterial communities in treatment plant.

UV-Vis spectroscopy analysis of effluent

The degradability of effluent after biostimulation process was confirmed by UV-Vis scanning absorption spectrum and showed elevation in overall along with generated scan peaks (Fig.1a-b). Results revealed comparative absorption spectrum by UV-Vis spectroscopy for control and degraded sample between 250 to 700 nm. Moreover, disappearances of some peaks in degraded sample and decreasing of absorption maxima indicated depolymerisation of organic compounds present in effluent. Similarly, the comparison of control and degraded sample at similar pattern of these data has been also recorded in our previous study for degradation of ligninocellulogic compound present in pulp and paper industry waste¹⁶. The result revealed a high peak absorption in different UV-region with

maximum absorbance (λ_{max}) of λ_{max} 250, λ_{max} 270 and λ_{max} 310 before biostimulation in effluent as control showed in Fig.1a-b. Moreover, after biostimulation the new peaks with maximum absorbance of λ_{max} 270, λ_{max} 310 and λ_{max} 370 were noted while some peak disappeared as shown in Fig. 1b. This change in the peak area and height indicates the conversation of compounds into complex metabolites²².

Evaluation of remediation of organic pollutants

The characterisation and identification of residual organic pollutants was done by GC-MS analysis from pulp paper industry effluent discharged after secondary treatment. The evaluation of degraded organic pollutants before and after biostimulation process revealed the change in peak height and retention time (RT) (Fig.2a-b and Table 2). The major pollutants were identified at RT-10.36, RT-10.59, RT-15.10, RT-17.82, RT-22.94, RT-25.25, RT-25.54 and RT-25.47 which are listed in Table 2. Out of these compounds, some were with EDCs nature compound identified as hexadecanoic acid, trimethylsilyl ester (RT-22.94) and pentadecanoic acid (RT-25.54) and were reported as mutagenic compound by United States Environmental Protection Agency³⁵ and Endocrine Disruptor Screening Program³⁶. The persistent organic contaminants were also extracted with similar conditions of control after degradation by biostimulation process. In addition, tetradecanoic acid and hexadecanoic acid are essential plant-based fatty acids in humic substances⁴⁵. RT-10-59 was detected as benzoic acid from effluent sample and it will be reduced during biostimulation process by potential bacterial strains. Moreover, cinnamic acids (RT-18.58) were detected in biostimulated extracted sample, which is known as the product of lignin and hemicellulose fraction of lignocellulose. Cinnamic acid and guaiacol are thought to be generated in the lignin polymer by cleavage of ester linkages in guaiacyl and p-hydroxyphenyl units³⁷. They are also capable to form ester and ether linkages via reaction of their carboxyl and phenolic groups, respectively38. Most of the ester bonds are destroyed during the alkaline extraction of the pulping process, but some cinnamic acid remain bound to the lignin by ether, which were degraded during biostimulation process. Several saturated fatty acids such as octadecanoic acid (RT-30.37) and trimethylsilyl ester were also found in effluent (control sample) and degraded finally by bacterial community after biostimulation process. This study is supported by the previous research of degradation and detoxification of effluent²². Moreover, octadecanoic acid was detected from Eucalyptus camaldulensis and is also reported as extractive of ethanol or benzene³⁹. All the above compounds have a methoxyphenol group in their structures, but at the position opposite to the hydroxyl group on the aromatic ring. Moreover, the majority of these compounds are monomers of lignin compounds. Most of these compounds are disappeared after biostimulation process. Among these compounds, octadecanoic, hexadecanoic and lactic acid are predominated

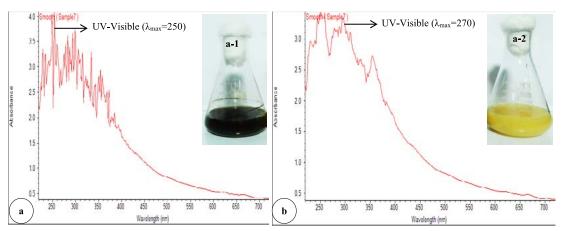


Fig. 1. UV-Vis spectroscopy analysis for investigate the degradability of pollutants at 250-700 nm before and after biostimulation process (a.a-1). Effluent before biostimulation process (b. b-2). Effluent after biostimulation process.

Table 2. Identified residual organic pollutants by GC-MS analysis in the TMS derivatized ethyl acetate extracts of pulp paper industry effluent

Retention Time (RT)	Identified compounds	Toxicity
Before Bioremediation	on	
6.88	2-Butoxyethanol	Acute toxicity
10.10	9-decenoic acid, trimethylsilyl ester	Data not reported
10.36	D-Lactic acid-DITMS	Acidosis
10.59	Benzoic acid, Trimethylsilyl ester	Gastric pain, vomiting, and allergic reactions
15.10	2,3,6-trimethyl phenol	Unknown
17.82	Citral	Reproductive toxicity
18.58	Cinnamic acid	Eco toxicity
22.94	Hexadecanoic acid, trimethylsilyl	Endocrine disrupting chemicals
	ester (Palmitic acid)	(EDCs)
25.25	2',6'-Dihydroxyacetophenone,	Data not reported
25.54	bis(trimethylsilyl) ether Pentadecanoic acid	EDCs
27.47	Phenol-4-ethyl-2-methoxy or 4-Ethylguaiacol	Data not reported
30.37	Octadecanoic acid, trimethylsilyl ester	EDCs
48.67	Phenol, 2,6-dimethoxy or syringols	Aguatic toxicity
49.98	Pthalatic anhydride	Data not reported
After Bioremediation	•	Data not reported
22.41	Phenol-2-methoxy-4-(1-propenyl or isoeugenol)	Data not reported
32.27	1,2-benzendicarboxylic acid disononyl ester	Data not reported
33.29	Benzyldehyde,4-(acetyloxy)-3-methoxy or acetyl vanillin	Data not reported
35.80	2,6-bis[trimethylsilyl]-3,4-dimethylphosphinine	Unknown
51.19	1,2-benzenedicarboxylic acid,bis (2-ethylhexyl)ester	Unknown
53.69	5,8-dimethoxy-6-methyl-2,4-bis(phenyl methyl) naphthalene	Data not reported
55.77	9-[2,6-diethylphenyl]2,8-dimethyl-9-h- purin-6-amine	Unknown

most in effluent⁴⁰. The toxicity of hexadecanoic acid is reported as an endocrine-disrupting chemical which is responsible for the hormonal disorder in fishes.

CFU, protein and biomass production

The measurements of CFU and biomass during the biostimulation process revealed continuous increase in comparison to the control (Fig. 3). The enhanced microbial community was observed after the addition of nutrients at different environmental conditions. The reduction of color was also observed in comparison to control, attributed to reduction of lignin and its metabolite during biostimulation process. The increase in biomass and CFU after enhancement of the carbon and nitrogen supply showed that

the autochthonous microbial communities do not have sufficient organic compounds present in the effluent as a nutrient. Hence, increase in the biomass could only be possible after addition of nutrients. The biomass increases the assisted biostimulation process. The gradual increase in the bacterial cell with diversified morphology also shows a strong confirmation of detoxification of effluent after biostimulation process. Further, it confirmed that in the presence of optimum nutrient and environmental conditions, the increasing bacterial community has sufficient potential for degradation of residual complex organic compounds. The degradation and detoxification of complex organic compounds from anaerobically digested effluent distillery

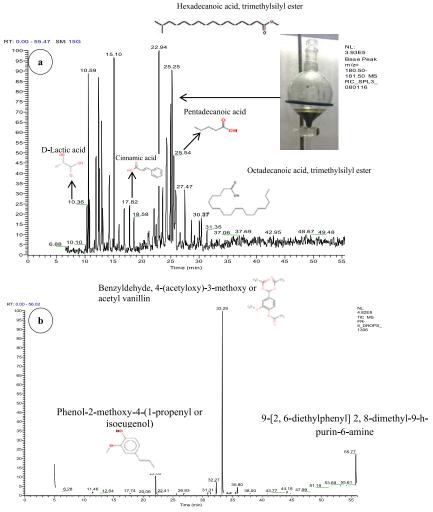


Fig. 2. Total Ion chromatogram (TIC) of TMS derivatized detected residual organic pollutants from wastewater by ethyl acetate extract of biostimulation process (a). Effluent before the biostimulation process (b). Effluent after the biostimulation process.

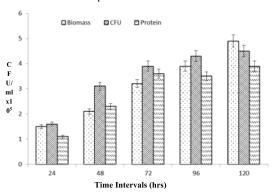


Fig. 3. Pattern of biomass, colony forming unit and protein contents in the pulp paper industry effluent after biostimulation process.

was observed after supplementing adequate $nutrients^{24}$.

Morphological view of bacterial community during biostimulation

The SEM analysis of the biostimulated sample showed the diversity of bacterial community (Fig. 4a-b). Besides, there was also a reduction in color, lignin, and organic compounds compared to control over the bacterial growth period. Addition of nutrient increases the bacterial community. Furthermore, SEM analysis of degrading samples also supported the evidence that the organic compounds are utilized as nutrients through the process of co-metabolism for cell multiplication. The significant increase in

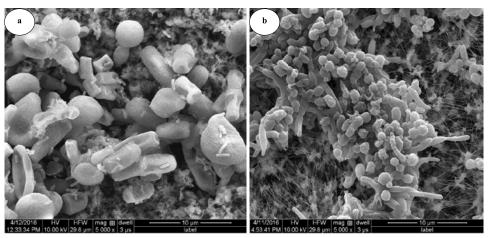


Fig. 4. Morphological view of bacterial strains observed under scanning electron microscopy analysis during bioaugumentation process at 5000x magnification (a-b).

the bacterial cells with diversified morphology strongly indicated that the bacterial species are essential for detoxification of different pollutants found in the effluent. The result revealed complex irregular structure mixed with an elongated rod or cylindrical shaped bodies and organic polymer i.e. lignin, cellulose and heterogeneous compound on the surface of effluent. A similar observation for the granulated appearance of lignin with the complete structure of different size has been reported in an earlier study⁴¹. Furthermore, the irregular shape is also indicating the lignin complexion with different

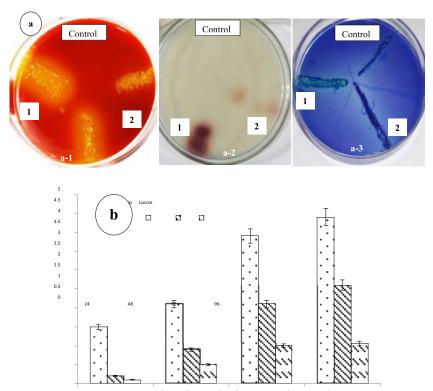


Fig. 5. The ligninolytic i.e. a-1 (Mnp), a-2 (Lip) and a-3 (Laccase) enzyme activity isolated bacterial strains a (*Aeromonas salmonicida*) and b (*Bacillus* sp) on plate assay (a) and for the extracellular enzyme activity was measure by biostimulated sample (b).

heavy metals and another carbonyl, hydroxyl and phenolic compounds⁴².

Ligninolytic enzyme activity

The role of extracellular enzyme activity by the bacterial community during the biostimulation is shown in Fig. 5. The maximum MnP was noted as 520 U/ml between 90-92h of the biostimulation process. The Lip and MnP were recorded as dominating enzymes at the initial stage of bacterial growth during the biostimulation process. The highest Lip was recorded as 0.0079 and 0.0091 U/ml, respectively at 48 and 96 h of incubation during bacterial growth in order to degrade organic pollutants. However, the incubation of laccase at the later stage of growth phase i.e. at 120 h indicated that phenolic compound was dominant and laccase could contribute very vital role. Moreover, maximum induction of laccase (41.0 U/ml) was noted at 120 h. The role of extracellular enzyme activity by the autochthonous bacterial community was observed during the degradation of effluent. The enzymes secreted by bacteria during the biostimulation

process were measured in the supernatant after centrifugation of sample. Moreover, Lip and MnP were recorded as dominating enzymes at the initial stage of bacterial growth. The observation of ligninolytic enzyme activity was observed on plate assay based on the selective agar qualitative screening and different indicator dye (Fig.6). The strong zone across the bacterial colony indicates the production of ligninolytic enzymes on the phenol red plate by the potentially isolated Mnp on phenol red plate, laccase on the guaiacol plate and Lip on the azure-B plate (Fig.5a-1, a-2, a-3). The ability to produce Mnp enzymes of isolated bacterial strains (Aeromonas salmonicida and Bacillus sp.) was measured by comparing the diameter of the colony and yellow-brown circles. Moreover, laccase is a multicopper oxidase and forms a brown zone due to the presence of guaiacol by the single-electron oxidation of organic compounds for the respective radical species. The bacterial cell produces extracellular oxidative enzymes including peroxidases that have been involved in lignin degradation. The combination of

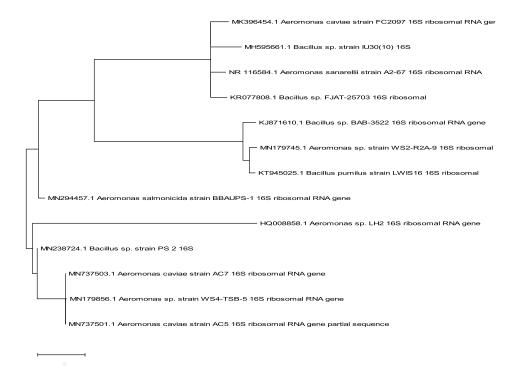


Fig. 6. Phylogenetic tree showing the relationship of bacterial communities growing during biostimulation process. The accession numbers for all strains used in tree construction are indicated in the figure.

these extracellular peroxidases has applications in the removal of phenolic compounds from effluent along with lignin degradation⁴³.

Purified isolated bacterial strains were identified based on 16S rRNA sequencing. The 16S rRNA sequence data were submitted to National Center for Biotechnology Information Search database (NCBI) using the Basic Local Alignment Search Tool (BLAST) tool and the obtained sequences were then further compared with sequences of bacteria from known taxa. Further strains BBAUPS-1 (Aeromonas salmonicida) and BBAUPS-2 (Bacillus sp.) were isolated

based on 16S rRNA sequencing. Moreover, the partial sequences were submitted to the GenBank public database under the accession number MN294457.1 (BBAUPS-1) and MN238724 (BBAUPS-2), respectively (Fig.6).

Phytotoxicity evaluation after biostimulation process

The seed germination test of *Triticum* aestivum and *Cicer arietinum* was inhibited most by 88 and 96% after irrigation with effluent from the pulp paper industry site in comparison to the control samples in which it is least inhibited i.e. 8 and 9%, respectively (Fig.7a-b). The comparative

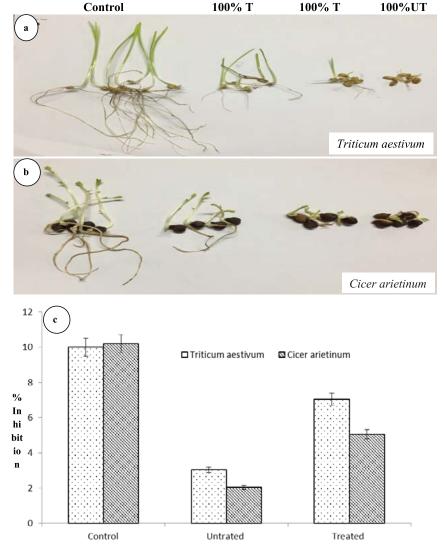


Fig. 7. Seed germination test by effluent of pulp paper industry untreated and bacterial treated on *Triticum aestivum* (a) *Cicer arietinum* (b) percent inhibition of seed germination (c), T-treated, UT-untreated

toxicity assessment of biostimulated samples revealed reduction in the toxicity of effluent and increase in the growth of plants. Further, the result also revealed that Cicer arietinum was found more sensitive than *Triticum aestivum* (Fig.7.c). The increased toxicity in seed germination was observed due to the presence of metals and ions in the effluent in high concentration in the control sample. The comparative cytotoxic and genotoxicity effect of effluent before and after biostimulated sample on A. cepa roots were also determined based on chromosomal aberrations⁴⁴. Reducing effluent toxicity has supported the removal of pollutants after the bacterial growth on toxic compounds. The seed germination tests showed inhibition in the seed germination attributed to the toxic compounds which were inhibitory for the α-amylase enzyme. Similar observations were also reported in earlier studies²⁴. The inhibitions shown in this assessment were underdeveloped root length and the lateral root due to some hormonal disturbance in the treated plant. These results clearly indicated that the increase in the toxicity of effluent was directly related to the load of pollutants.

CONCLUSION

The study concludes that the biostimulation process is suitable for the optimization of degradation and decolorization of the pulp paper industry effluent after secondary treatment. The effluent contains various organic and inorganic pollutants along with high concentration of ion and salts that cause phytotoxicity in *Triticum aestivum* and Cicer arietinum. The discharged effluent is reported as the main source of soil and aquatic pollution in the large area of agricultural land. These effluents have inhibitory effects over seed germination and in turn reduce the productivity of land and subsequently reduce farmer's income. The biostimulation process enhances the remediation strategy of residual organic pollutants by the bacterial community and is suitable for detoxification and decolorization of effluent for safe environmental disposal. In addition, disposal of effluent after such treatment and processes will not harm the environment much and provide better support to the agricultural system of India.

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CONFLICT OF INTEREST

The authors declare that there is no conflict of interest.

AUTHORS' CONTRIBUTION

This work is done by PS as PhD Scholar, while bacterial sequencing was done by PC and Deepshi Chaurasia. ST has supported to draft this manuscript and Prof. Ram Chandra has worked as overall mentor of the work.

ETHICS STATEMENT

This article does not contain any studies with human participants or animals performed by any of the authors.

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DATA AVAILABILITY

All datasets generated during this study are included in the manuscript and the Supplementary Files.

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