RESEARCH ARTICLE

An experimental approach for the utilization of tannery sludge-derived Bacillus strain for biosorptive removal of Cr(VI) -contaminated wastewater

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Abstract

Biosorption efficacy of Bacillus strain DPAML065, isolated from the tannery sludge, was appraised for the removal of toxic hexavalent chromium (VI) ions from synthetic wastewater. Effects of the process variable on biosorbent surface by variation in pH, metal Cr(VI) concentration and retention time were examined using batch experiments. The isolated Bacillus strain biosorbent was studied for its morphology and surface chemistry through FE-SEM, EDX and FTIR. It discloses that, the reduction mechanism of Cr(VI) during the process is mainly attributed to precipitation in addition to the functional groups (such as –COOH, –OH, C–O, P=O) present on the cellular matrix of Bacillus. Biochemical tests and 16s rRNA sequencing were also performed to identify the biosorbent at the genus level. A 95% Cr(VI) removal efficiency was procured by Bacillus strain DPAML065 biosorbent at pH 6, incubation period 24 h, 80 mg/L initial feed concentration and operational temperature 35 °C. Equilibrium behaviour of chromium binding follows the Langmuir isotherm model (R^2 = 0.968) with an adsorption capacity of 106.38 mg/g. Kinetic modelling disseminates that biosorption of Cr(VI) ions by Bacillus strain DPAML065 obeyed pseudosecond-order model (R^2 = 0.984) rather than the pseudo-first-order model. Concisely, the results indicate that the Bacillus strain DPAML065 is a potential, economically feasible and eco-friendly biosorbent which can be effectively used for removal of chromium (VI) from wastewater.

Keywords Tannery sludge \cdot Bacillus species strain $DPAML065 \cdot 16$ s rRNA gene sequencing \cdot Hexavalent chromium \cdot Cr(VI) \cdot Biosorption

1. Bacillus species strain DPAML065 was explored for chromium biosorption.

- 2. The biosorbent showed Cr(VI) removal efficiency of 95% under optimal conditions.
- 3. Maximum biosorption capacity of 106.38 mg/L was estimated for Bacillus strain DPAML065 biosorbent.

4. Chromium biosorption on Bacillus strain DPAML065 obeyed Langmuir isotherm and pseudo-second-order.

5. The study confirmed the biosorption of Cr(VI) on the bacterial cell surface through multistep mechanism process.

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Introduction

Among the various anthropogenic activities, untreated discharge of heavy metals effluent from the industries is one of the substantial problems (Gupta et al. [2000;](#page-11-0) Ghiasi et al. [2020;](#page-11-0) Prabhakar and Samadder [2020\)](#page-11-0). The major problem is its magnitude and has been perused for its effect on the environment including genetic and developmental defects in humans as well as on aquatic life (Huang et al. [2013](#page-11-0); Singh and Kumar [2020;](#page-12-0) Sharma and Malaviya [2016](#page-12-0); Sanjay et al. [2018](#page-12-0); Uniyal et al. [2013](#page-12-0)). The most common heavy metals like chromium, cadmium, lead, arsenic and mercury have a nearly meagre biological role and are detrimental to human health even at a minimal concentration (Khan et al. [2013;](#page-11-0) Singh and Kumar [2020](#page-12-0); Lata et al. [2019\)](#page-11-0). The effluent discharge having chromium has diverse use in industries like steel, cleaning agents and electroplating and also in the production of chromic acid which is extensively employed in leather industries in India (Bharagava and Mishra [2018](#page-10-0); Sanjay et al. [2018](#page-12-0); Singh et al.

[2018](#page-12-0); Torab-Mostaedi et al. [2013\)](#page-12-0). Among these industries, leather and its allied industries consume nearly 35 L of water per 100 kg of skin hides during the process. Thus, the discharged effluent consists of chromium, which is the most pernicious contaminant emancipated in an enormous quantity into the environment. Furthermore, it is documented as one of the most abhorrent anthropogenic contaminants released from the industries (Nandy et al. [1999;](#page-11-0) Ryan et al. [2002](#page-12-0); Kumari et al. [2016\)](#page-11-0). Chromium exists in multiple oxidation states, with trivalent Cr(III) and hexavalent Cr(VI) species being the most prevalent forms. Out of it, Cr(VI) is more prevalent and mutagenic compared to Cr(III), the latter being more stable. Chromium is the most lethal constituent classified by the International Agency for Research on Cancer (IARC [1990\)](#page-11-0) as a group-1 human carcinogen due to its high solubility and fast permeability in biological cell membrane as it interacts with intracellular proteins and nucleic acids leading to structural modifications (Beukes et al. [2017](#page-10-0); Cardoso et al. [2017](#page-10-0); Vinod et al. [2010](#page-12-0)). The convergence of chromium in leather effluent is about 2 to 5 g/L, which significantly is sky high when compared to the suggested allowed limit of 2 mg/L. If the concentration of $Cr(VI)$ surpass $>$ 0.05 ppm, it predominantly affects the human health, causing damage to internal organs like the kidney, lungs, liver and pancreas and even leading to bioaccumulation in organisms (Vinod et al. [2010;](#page-12-0) Cheung and Gu [2007;](#page-11-0) Belay [2010](#page-10-0)).

Conventional methods such as filtration, precipitation, coagulation, reverse osmosis, solvent extraction, ion exchange electrochemical treatments, adsorption, evaporation recovery, kaolinite and activated carbon were considered and were reported successful for the removal of hexavalent chromium (Balasubramanian et al. [2019;](#page-10-0) Arukula et al. [2018](#page-10-0); Deepa et al. [2019;](#page-11-0) Zambrano and Min [2019](#page-12-0); Oves et al. [2013](#page-11-0); Rathinam et al. [2010;](#page-12-0) Xiao et al. [2010](#page-12-0); Saxena et al. [2016](#page-12-0); Ahluwalia and Goyal [2007\)](#page-10-0). As these technologies involve high capital and generate metal-rich sludge which results in environmental pollution, it shifts the focus towards some economic and environment-friendly approaches (Tsekova et al. [2010;](#page-12-0) Green-Ruiz et al. [2008\)](#page-11-0). Chromium-tolerant bacteria can be considered as the most practical approach and is even eco-friendly to reduce hexavalent chromium to trivalent chromium. Cr(VI) detoxification and bioremediation due to the presence of high surface charge and contact area of the bacteria could help effectively during an interchange with the metals that alter physiologically and stay dynamic in the environment (Cervantes and Campos-García [2007](#page-10-0); Dash et al. [2014\)](#page-11-0). Bacteria can bind to a high concentration of heavy metals by some of the most common biosorption mechanisms such as physical adsorption, ion exchange, coordination, chelation, precipitation and complexation. Various familiar functional groups were incorporated in biosorption, which includes hydroxyl, carboxyl and amine groups resulting in the binding of metal ions in bacteria (Deepa and Mishra [2020](#page-11-0); Volesky [2003;](#page-12-0) Bharagava and Mishra [2018](#page-10-0)).

The main objective of the present study was to isolate and identify the Cr(VI)-resistant bacteria from the tannery sludge and to understand the performance of bacteria on the fate and mobility of chromium. The biosorptive capacity of Cr(VI) in response to initial concentration, pH and contact time was evaluated. Morphological variations prior to biosorption treatment and after biosorption of Cr(VI) on Bacillus strain DPAML065 were investigated by XRD analysis. Different functional groups involved in Cr(VI) biosorption were spotted using FTIR analysis. FE-SEM and EDX were employed to clarify the mechanism of Bacillus species in the removal of chromium through biosorption. Factors affecting biosorption isotherms and kinetic studies were also deliberated. Thus, this biosorption study offers a simple and economical solution in the reduction of hexavalent chromium.

Materials and methods

Isolation of the bacteria and culture conditions

The microorganisms were screened out in the tannery sludge from the leather industry located in Kanpur, India. The isolated bacteria were characterized through microscopic, biochemical characteristic study and 16s rRNA sequencing analysis, as shown in Table [1.](#page-2-0) The stock solution of the bacterial culture contains sodium chloride 5.0 g/L, meat extract 1.5 g/L, yeast extract 1.5 g/L and peptone 5.0 g/L. The bacterial cells used in the experimental study were harvested in nutrient broth medium for 24 h at 35 °C and then centrifuged at 10000 rpm for 12 min. Further, the pellets were washed twice with 0.1 M phosphate buffer solution.

Primary screening of chromium-resistant bacterial strain and growth conditions

The metal-resistant microbe was isolated from the tannery sludge samples by serial dilution method, in which 1 mL of the sample was serially diluted in a phosphate buffer solution. Desired diluted samples (10^{-3} to 10^{-10}) of 0.1 mL were plated on the duplicates of the appropriate media amended with varying concentrations of Cr(VI) using spread plate method and were incubated at 37 °C for 24–48 h. After incubation, the isolated strain was repeatedly sub cultured on the appropriate medium to obtain a pure culture with maximum Cr(VI)-tolerant bacterial colonies and then subjected to biochemical characterization, identification by 16s rRNA gene sequencing analysis and other studies.

Chemicals and reagents

Nutrient Agar and Nutrient Broth were provided by Hi-Media Laboratories Pvt. Ltd., India. Nitric acid $(HNO₃)$,

Morphological study	Biochemical tests Results Results		Acid production from carbohydrates	Results	Growth at pH	Result	
Cell shape	Rods	Indole test		Dextrose	$+$	4.0	$+(W)$
Size (μm)	$2 - 4$	Methy reductase test	$+$	Mannitol	$^{+}$	5.0	$+(W)$
Elevation	Raised	Voges proskauer test	$\overline{}$	Xylose	$\ddot{}$	6.0	$+$ (G)
Surface	Flat	Urea hydrolysis	$+$	Use of sugar as a sole carbon source		7.0	$+$ (F)
Margin	Entire	Casein hydrolysis	$+$	Fumarate		8.0	$+(W)$
Gram stain	$\ddot{}$	Catalase test	$^{+}$	Glycerol	$\ddot{}$	9.0	
Opacity	Opaque	Oxidase test	$^{+}$	Lactose			
Motility	$^{+}$	Nitrate reduction					
		Gelatin hydrolysis	$+$				

Table 1 Morphological/ physiological and biochemical characteristics of the isolated bacteria Bacillus species DPAML065

W weak, F fair, G good

hydrochloric acid (HCl), sodium hydroxide (NaOH), potassium dichromate $(K_2Cr_2O_7)$ and 1, 5-diphenyl carbazide (DCP) were purchased from Loba Chemie and Merck, India. All the reagents used in this experiment were of analytical grade with high purity levels (about 99%), without further purification.

Biosorption experiments

The biosorption experimental studies were carried out at room temperature, and the experimental protocol was accomplished. Preparation of chromium stock solution of 1000 ppm was done by adding 2.835 g of potassium dichromate $(K_2Cr_2O_7)$ salt to 1000 mL de-ionized water. The working solution (20, 40, 60, 80, 100 and 120 mg/L) was prepared by diluting the stock solution with Millipore distilled water (Merck, Germany). All the experiments were conducted in triplicates. Two hundred fifty milligrams (wet weight) of bacterial cells was added to the varying concentrations of the chromium mentioned above, and batch study was done. Then, the flasks were incubated in a rotary shaker with 120 rpm at 35 °C for 24 h (REMI CIS-24 PLUS, India). Control was maintained with Cr(VI) solution without bacterial culture. After, 24 h of growth, bacterial culture from each test concentration was centrifuged at 10000 rpm for 12 min (REMI R 24, India). The optical density (OD_{600}) was monitored using UV Spectrophotometer (Lab-Tech, China). Further evaluation of Cr(VI) concentration reduction was done by the DPC method at 540 nm as per the method de-scribed by (APHA [2012\)](#page-10-0) and was calculated through the calibrated curve developed from $K_2Cr_2O_7$. Cr(VI) was measured by taking 5 mL of the sample in a beaker with an adjusted pH of 1.0 which was done by using 0.1 N HNO₃; the sample was made up to 100 mL using Millipore water. Further 2 mL of DPC (1,5-diphenylcarbazide $(C_{13}H_{14}ON_4)$) solution was added in the sample (prepared by dissolving 500 mg in 100 mL of acetone) and left to stand for 5 min to develop a pink colour. The absorbance was noted down in a UV spectrophotometer. Removal percentage of Cr(VI) biosorption ($R \%Qmg/g$) was calculated from the following Eqs. (1) and (2)

$$
\% \text{Biosorption of Cr (VI)} = \frac{C_0 - C}{C_0} \times 100 \tag{1}
$$

$$
Q = \left(C - C \right) \times \frac{V}{W} \tag{2}
$$

where C_0 = initial metal concentration (mg/L), C = final concentration of the metal (mg/L), $V =$ volume of the solution (litres) and W is the weight of the bacterial cells (grams).

Determination of minimum inhibitory concentration of heavy metals (MIC)

The Cr(VI) minimum inhibitory concentration of Bacillus species strain DPAML065 was determined. A stock solution of heavy metal was prepared at increasing concentrations (20, 40, 60, 80, 100 and 120 mg/L); the solutions were incubated with separate batches of the cultured bacterial strain. The result was then measured by agar dilution method for the complete inhibition zone. Thus, MIC was noted when no growth was observed in the petri dishes even after 48 h of incubation, and the results are given in Table S1.

FE-SEM and EDX analysis

Figure [4](#page-6-0) depicts the morphological and surface characterization of the metal precipitates of chromium-loaded and chromium-unloaded Bacillus species. Quantification of chromium in the Bacillus species was obtained from energydispersive X-ray analysis and field emission scanning electron microscopy (FE-SEM) (Hitachi S-4300). For FE-SEM and EDX analysis, the chromium-loaded and chromiumunloaded samples were prepared by using 2.5% (W/V) of glutaraldehyde in sodium phosphate buffer having pH 7.0 with 0.1 molarity. Further, the pellets were washed thrice with the phosphate buffer and then were fixed for 1 h at 4 °C in the buffer solution before dehydration. Different concentration of ethanol (W/V) was used in turn to dehydrate the samples: further, the samples were fixed on the cover slip, and these cover slips were supported on the copper grids for further morphological studies (Nwinyi and Owolabi [2019](#page-11-0)).

FTIR and XRD analysis

Fourier transform infrared spectroscopy (FTIR) analysis for Bacillus strain which was grown in the nutrient broth medium was recovered through centrifugation (10,000 rpm for 10 min); further, the cells were washed with Millipore water and lyophilized. Chromium-loaded and chromium-unloaded bacterial cells were finally grounded with solid KBr, and then, the infrared spectra were recorded at 500–4000 cm−¹ using FTIR spectrometer (Perkin Elmer) which is represented in Table [3.](#page-7-0) X-ray diffraction studies (XRD) have been carried out with anhydrous chromium-treated and non-treated chromium pellets and were examined with Bruker (Model: Kappa APEX II).

Results and discussion

Identification of the isolated bacterial species

In the present study, morphological and biochemical characteristics were carried out for the isolated bacterium from the tannery sludge and are listed in Table [1](#page-2-0). A 16s rRNA gene analysis of the stain DPAML065 was carried out by DNA sequencing facility, Yazz Xenomics Pvt. Ltd., Chennai, Tamil Nadu using universal primers, 8F (5′AGAGTTTG ATCCTGGCTCAG3′) and drive back primer; 1541R (5′ AAGGAGGTGATCCAGCCGCA3′). Further, nucleotide sequence data were deposited in the Gen-Bank sequence database. The online program BLAST was used to find out the

Fig. 1 Phylogenetic tree based on the 16s rRNA sequence of gene strain compared with that of analogous organisms accession number MT250939.1

related sequences with known taxonomic information in the databank at the NCBI website [\(http://www.ncbi.nlm.nih.gov/](http://www.ncbi.nlm.nih.gov/BLAST) [BLAST](http://www.ncbi.nlm.nih.gov/BLAST)) to accurately identify the strain, as shown in Fig. 1; the phylogenetic bonding of the bacterial isolate was identified. The results revealed that the isolated Bacillus species DPAML065 bacteria was homogenous with 99% of 16s rRNA gene sequencing. Further, the strain was identified as Bacillus species, and the gene sequence was deposited in the Gene bank under accession number MT250939.1.

Effect of pH on Cr(VI) reduction

A pivotal role in the biosorption of chromium ions is manifested by pH (Lata et al. [2015\)](#page-11-0). The pH of the medium primarily influences the biosorption ability as it affects the valency of metals in solution and functional groups present on the biomass. Moreover, pH also affects the network of negative charges on the surface of the biosorbing cells (especially in case of Gram-positive bacteria) and the chemistry of the walls, as well as physicochemistry and hydrolysis of the metal (Kalola and Desai [2019\)](#page-11-0). The pH optimization of chromium by Bacillus species in the range of $(pH 1–9)$ at an initial concentration of Cr(VI) 20 mg/L with a contact time of 24 h at 35 °C was studied. As shown in Fig. [2](#page-4-0), the biosorption of Cr(VI) increased gradually with an increase in pH value from 1 to 5 and maximum Cr(VI) biosorption of 97% was observed at pH 6. Whereas in the case of pH values < 6 , chromium biosorption was less due to the decreased growth of cell in the medium and the functional anionic active sites protonate at the surface of biomass and attract a large number of anionic groups. Depending on the pH, chromium exists in the form of chromate (CrO_4^2) , dichromate $(Cr_2O_7^2)$ and bichromate (HCrO4 −) (Acharya et al. [2020](#page-10-0); Babel and Kurniawan [2004\)](#page-10-0). At low acidic pH < 6, chromium is in the form of $Cr_2O_7^{2-}$. The stable form of Cr(VI) existing at pH 6 is HCrO₄⁻which possess negative charge; hence, an electrostatic attraction between chromium species and the positively charged functional groups present on the bacterial cell surface of the biomass occurs

Fig. 2 The effect of pH on reduction of Cr(VI)

(Anah and Astrini [2017\)](#page-10-0). In addition to it, at pH 6, favoured cell growth in the medium takes place in which the propinquity of positively charged ions of the utility group available on the cell of the biomass results in the electrostatic attraction, precipitation, ion exchange and complex formation between the anionic Cr(VI) and biomass (Nguema et al. [2014](#page-11-0); Jobby et al. [2019;](#page-11-0) Hu et al. [2012\)](#page-11-0). Researchers also reported that B. coagulans and Halomonas species have also reduced Cr(VI) at an optimum pH 6–7 with an initial concentration of chromium at 60 mg/L with 96.7% of Cr(VI) reduction (Murugavelh and Mohanty [2013](#page-11-0); QuiIntana et al. 2001). At pH > 6, chromium exists in the form of CrO_4^2 ; hence, the reduction capacity of Cr(VI) was drastically decreased which might be due to the development of negative charge on the cell surface developed as a result of deprotonation of carboxylate groups. Even metals start precipitating at high pH, which reduces the solubility and affinity of metal ions leading to reduced biosorption (Vijayaraghavan and Balasubramanian [2015\)](#page-12-0).

Biosorption behaviour of Cr(VI) on Bacillus sp. strain DPAML065

The initial metal concentration effect of Cr(VI) on Bacillus strain DPAML065 was examined at an optimum pH of 6.0. Figure [3](#page-5-0) shows the optimum (pH 6) with varied concentration of Cr(VI) in the range from (20–120 mg/L). Results revealed that in low concentration, the reduction was nearly high. As the concentration increased, the biosorption efficiency significantly decreased to 120 mg/L, which might be due to the increased toxic effect of cell growth in the medium. Additionally, elevated mass transfer with aggravating kinetic energy due to the collision between the metal ions and biosorbents could also be responsible for it. (Uniyal et al. [2013\)](#page-12-0). Besides this, it can also be observed that the concentration of Cr(VI) was drastically reduced within 24 h by Bacillus species. Similar studies have been documented (QuiIntana et al. [2001\)](#page-11-0) in Thiobacillus ferroxidans in which the concentration of chromium was considerably decreased from 60 to 1.89 mg/L within 48 h. Thus, the results conclude that the biosorption of Cr(VI) by isolated culture Bacillus strain *DPAML065* is a time-dependent activity which removed the maximum concentration of chromium ions, i.e., 80 mg/L within 24 h, and depicted a biosorption efficiency of 95.4%. Besides this, the potential strain was compared with the other Bacillus species of live biomass in the removal of Cr(VI) and is represented in Table [2](#page-5-0).

The minimum inhibitory concentration of heavy metals (MIC)

The Cr(VI) MIC for Bacillus strain DPAML065was provoked by agar dilution method in solid media which ranged from 20 to 160 ppm of hexavalent chromium and showed a maximal degree of resistance to chromium in 80 ppm when compared with other increased concentrations as shown in the Table S1.

FE-SEM and EDX analysis

Field emission scanning electron microscopy (FE-SEM) and energy-dispersive X-ray analysis (EDX) were used to determine the quantification of chromium ions on the biosorption surface. In Fig. [4d,](#page-6-0) the spectral image clearly exhibit that chromium was adsorbed on the surface of the Bacillus strain DPAML065. Similar quantification and accumulation studies of chromium was also documented with the help of FE-SEM and EDX by other researchers in fungal and bacterial strains (Aspergillus niger and Serratia sp.; Halomonas sp. (Srivastava and Thakur [2006](#page-12-0), [2012;](#page-12-0) Kalola and Desai [2019](#page-11-0)).

Different types of live biomass	Heavy metals removed	Concentration (mg/L)	Working pH	$%$ reduction	References
Bacillus pumilis	Cr(VI)	100	2	87.79	Sultan et al. (2012)
Bacillus cereus	Cr(VI)	100	2	86.79	Sultan et al. (2012)
Bacillus sphaericus	Cr(VI)	5.		66.6	Al-Daghistani (2012)
Pantoea agglomerans	Cr(VI)	100		83.64	Sultan et al. (2012)
Bacillus DPAML065	Cr(VI)	80	6	95.4	Present study

Table 2 The comparison of live biomass in removal of $Cr(VI)$

High biosorption of chromium was observed in the isolated potential microbe, as shown in Fig. [4d](#page-6-0) and was in good agreement with the reports of Guo et al. [2012](#page-11-0). However, chromium biosorption of Bacillus species from various sources was below the optimal conditions (Sultan et al. [2012](#page-12-0); Al-Daghistani [2012\)](#page-10-0). In the present study, very high biosorption of chromium was remarked when compared with different experimental conditions performed by other researchers (Sultan et al. [2012](#page-12-0); Al-Daghistani [2012](#page-10-0)). This phenomenon can be assigned to facts that the functional groups present on the surface of the bacterial cell wall hold polysaccharides as building blocks of proteins and lipids which perform ion exchange properties as well as precipitation mechanism. Thus, as a result, these functional groups are proficient in the binding of chromium ions (Hu et al. [2012;](#page-11-0) García-Mendieta et al. [2012](#page-11-0)).

FTIR analysis

The infrared spectra were performed for control (without chromium) and chromium-loaded sample in the range of 500– 4000 cm^{-1} at room temperature. The analysis was done to investigate predominant groups present on the cell wall of Bacillus

biosorption shown in Table [3](#page-7-0). Figure [5](#page-8-0) represents the FTIR spectra of before and after biosorption of chromium peaks which showed different bonds due to the interactive action of bacteria on the mineral surface with copious functional groups. In the control sample, a broad peak was observed at 3432.95 cm⁻¹ that designates the presence of amines (N–H) bonded group with a stretching and medium intensity. But in case of a chromiumloaded sample, a sharp peak was observed at 3426.75 cm⁻¹ which shifted the functional group, indicating a secondary amide protein representing the participation of membrane proteins in the binding of chromium. The peak at 2960.18 cm^{-1} in control (Bacillus microorganism) represents the presence of aldehyde or other =C–H with strong stretching vibration. After treatment, the band appeared at 2942.07 cm⁻¹ in a reduced form which revealed that symmetric or asymmetric C–H stretching vibrations of aliphatic acids involved in the reduction of chromium ions. The band at 2925.77 cm⁻¹ in control biomass (C–H) bond with stretching vibration along with a strong intensity has been shifted after biosorption of chromium to 2914.07 cm^{-1} depicting that C– H covalent bond participated in the reduction of chromium ions (Ramrakhiani et al. [2011\)](#page-11-0). The existence of spectral peaks has

strain DPAML065, which is responsible for chromium

Fig. 3 The biosorption behaviour of Cr(VI) by Bacillus species DPAML065 at an optimum pH 6 with different concentrations of Cr(VI)

Fig. 4 The FE-SEM and EDX analysis of Bacillus sp. strain DPAML065. A(i) Microscopic image of rod-shaped bacteria, A(ii) morphological features and surface characteristics of biosorbent before Cr(VI) adsorption

and (B) after Cr(VI) adsorption. C, D EDX analysis of Cr(VI)-unloaded and Cr(VI)-loaded Bacillus species during biosorption

been observed in chromium-loaded biomass at 2850.76 cm−¹ and 2355.28 cm⁻¹ on the surface of Bacillus strain DPAML065 that might be due to the attraction of positive electrons donated by chromium species which are favourable for the reduction of chromium ions during the biosorption process (Han et al. [2007\)](#page-11-0). The bands appeared in a raw sample at 1722.17 cm^{-1} , 1652.67 cm⁻¹ and 1640.97 cm⁻¹ depicting the presence of (C=O) carbonyl acid groups with strong and stretching vibrations. While in chromium-loaded biomass, the peaks were observed at 1629.96 cm⁻¹ which are attributed to –NH bending of primary and secondary amides from the peptide bond coupled with –COO anion. Several other peaks in the treated sample at 1474.43 cm⁻¹ and 1439.34 cm⁻¹ wavelengths designate the presence of aromatic functional groups which mainly participated in the reduction of Cr(VI) ions (Sun et al. [2014\)](#page-12-0). The FTIR spectral band observed at 1329.92 cm^{-1} in control has been shifted to 1399.42 cm⁻¹ indicating that aromatic amine (CN stretching) exhausted the chromium ions during the biosorption process. The peak present at 1191.60 cm⁻¹ and 1059.47 cm⁻¹ in control biomass has been shifted to 1082.18 in the chromium-

loaded biomass which suggests that reduction of chromium ions with the involvement of polysaccharides P=O and (C–O) groups of alcohol with strong and stretching vibration (Sun et al. [2014\)](#page-12-0). A small and sharp peak appeared at 978.96 cm^{-1} and 794.53 cm^{-1} that corresponds to bending aromatic compounds which affect revelation and suppression of functional groups on the microbial cell wall (Arivalagan et al. [2014;](#page-10-0) Rajesh et al. [2014](#page-11-0)). Therefore, these results suggest that chromium binding on to the bacterial biomass is because of the involvement of carboxyl, hydroxyl and amine groups, which is accountable for the electrostatic attraction and complexation mechanisms while reducing chromium ions (Ramrakhiani et al. [2011;](#page-11-0) Chakravarty and Banerjee [2012;](#page-10-0) Javanbakht et al. [2014\)](#page-11-0).

XRD analysis

Figure [6](#page-8-0) shows the powder X-ray diffraction patterns of (A) control biomass and (B) chromium-loaded biomass of Bacillus strain DPAML065 grown in nutrient broth medium, supplemented with chromium, and the results were analysed using Table 3 FTIR transmittance spectra (500–4000 cm⁻¹) of Bacillus species strain DPAML065 and their possible assignments

Jade 5 software. In Fig. [6](#page-8-0), it can be noticed that the XRD pattern of control biomass (A) revealed the significant peak around $(2) = 27.53$, 45.47 and 56.73 which attributes to the presence of fatty acids and some polysaccharides in the cell wall of Bacillus strain DPAML065 (Das et al. [2014;](#page-11-0) McLean et al. [2000](#page-11-0)). Additionally, the spectrum showed several other peaks of (2) at 31.69 and 45.45 with a d- spacing values of 2.82 and 1.99 and was referred with ICCD-00-005-0628 as well as through the peak information from JCPDS file. It revealed the presence of sodium crystals that might be due to the existence of NaCl in the nutrient broth medium. In Fig. [6b](#page-8-0), the Cr(VI)-loaded biomass showed some peaks which appeared in the XRD pattern of the bacterial cell, which corresponds with the precipitation of reduced Cr(III) species. Various low-intensity peaks at 27.53, 56.65 and 66.45 are responsible for the existence of Cr (III) in the bacterial cell after the chromium biosorption process (Dhal et al. [2010;](#page-11-0) Karthik et al. [2017\)](#page-11-0).

Biosorption isotherm

Biosorption isotherm curve can be used for better understanding the equilibrium distribution of metal ions between the aqueous and solid phase, as well as the capacity of biosorption at a constant temperature. The most frequently used models are Langmuir and Freundlich, which illustrate the biosorption process (Sonal et al. [2020](#page-12-0); Joo et al. [2010](#page-11-0); Volesky [2003;](#page-12-0) Sala et al. [2002;](#page-12-0) Volesky and Holan [1995\)](#page-12-0). Langmuir isotherm is the supreme model that explained the mechanism in which the maximum adsorption capacity corresponding to homogeneous monolayer biosorption of metal ions on the surface of the biomass, by electrostatic attraction. The mathematical formula of non-linear Langmuir equation can be expressed as Eq. (3):

$$
q_{eq} = \frac{q_{\text{max}}bC_e}{1 + bC_e} \tag{3}
$$

where q_{max} (mg/g) is the maximum adsorption capacity of the chromium ions per unit weight of biomass to form a complete monolayer, Ce (mg/L) represents equilibrium concentration of metal, q_{eq} (mg/g) represents the amount of metal uptake occurs at equilibrium and b (L/mg) is the Langmuir constant, which also reflects the affinity of the chromium for binding sites on the biosorbent.

The Freundlich isotherm is the earliest known to describe non-ideal and reversible adsorption, but later its empirical model used to explain the multilayer adsorption with nonuniform distribution of adsorption heat and affinities over heterogeneous. The mathematical formula of non-linear Freundlich equation was represented in Eq. (4)

$$
q = K_f C_e^{1/n} \tag{4}
$$

where K_f (mg/L)ⁿ represents the Freundlich constant and *n* is the dimensionless Freundlich intensity parameter that defines the magnitude of adsorption driving force. Fig. S1 delineates the biosorption of chromium by Langmuir and Freundlich isotherm.

Fig. 5 a Control and **b** Cr(VI)loaded biomass shows the FTIR analysis of Bacillus species DPAML065

The parameters of the Langmuir and Freundlich were estimated by non-linear analysis is presented in Table [4](#page-9-0). Based on the obtained results of the correlation coefficient (R^2) , it has been concluded that the Langmuir isotherm model gave an acceptable fit to the experimental data than the Freundlich. Hence, the monolayer biosorption mechanism occurs during the biosorption of chromium on the surface of Bacillus DPAML065.

Biosorption rate kinetics

The kinetic study is important to understand the mechanism and rate of the adsorption process. The kinetics designates solute uptake, as well as its dominant contact period of the biosorbent in the solid medium of the interface. The adsorption of metal ions on the surface of biosorbent was evaluated by pseudo-first-order, pseudosecond-order and intraparticle diffusion model. Pseudofirst and pseudo-second-order kinetics models are the two kinetic models which were used for experimental biosorption of chromium on the surface of Bacillus strain DPAML065.

The pseudo-first-order model is expressed as:

$$
\log(q_e - q_t) = \log q_e - \frac{k_1}{2.303}t\tag{5}
$$

Fig. 6 Characterization of X-ray diffraction (XRD) of Bacillus species DPAML065. a Control biomass. b Cr(VI)-loaded biomass

Table 4 The Langmuir and Freundlich adsorption isotherm constants for Cr(VI) biosorption by Bacillus species strain DPAML065 Metal Biosorbent Langmuir **Freundlich** q_{max} (mg/g) b (L/mg) R^2 K_f n R^2 Cr(VI) Bacillus species 106.382 0.4277 0.968 42.35 0.2413 0.8479

where q_e and q_t represent the quantity of Cr(VI) absorbed in the active biomass (mg/g) at equilibrium and at time t (min) and k_1 is the first-order rate constant (min^{-1}) .

Similarly, the pseudo-second-order model can be expressed as:

$$
\frac{dq}{qt} = k_2 \left(q_{eq} - q \right)^2 \tag{6}
$$

The amount of metal adsorbed on the active sites of adsorbent at equilibrium is denoted by $q_{eq}(mg/g)$, or at any time t, the adsorption capacity of adsorbent is denoted by q_t and t is the contact time (minute), k_2 (g mg⁻¹ min⁻¹) is the rate constant of pseudo-second-order kinetics (Senthilkumar et al. [2006](#page-12-0); Lutke et al. [2019](#page-11-0)). A straight line was obtained by plotting t/q_t as the function of t (min) as shown in Fig. S2. The intraparticle diffusion model (Weber and Morris [1962a](#page-12-0), [b\)](#page-12-0) is expressed as:

$$
q_t = k_i t^{1/2} + C \tag{7}
$$

where k_i (mg g⁻¹ min^{-1/2}) represents the intraparticle diffusion model's initial rate constant and C (mg/g) is the intercept representing a constant related with the thickness of the boundary layer. The values of q_e , k_2 and correlation coefficient (R^2) were calculated using the slope and a straight-line intercept perceived in the graph and are listed in Table 5.

The outcome values are in good agreement and best fits with pseudo-second-order model as coefficient values, $R^2 = 0.984$, which has a higher correlation than the pseudo-first-order model $(R^2 = 0.921)$ is given in Fig. S2(A) and (B), indicating the chemisorption behaviour of Cr(VI) on the surface of Bacillus strain DPAML065 (Sinha et al. [2018](#page-12-0)). Similarly, numerous reports have been published on the chemisorption behaviour of heavy metals like chromium, cadmium, zinc, arsenic and copper along with lead by the Klebsiella, Arthrobacter and Rhodococcusopacus species, respectively (Bueno et al. [2008](#page-10-0); Muñoz et al. [2015](#page-11-0); Prasad et al. [2013\)](#page-11-0). Microbial biomass generously authoritative on the metal ions for their dynamic site, which could be better understood by chemisorption nature of the adsorbent surface (Munoz et al. 2015). Also, in Fig. $S2(C)$, it has been found that in the intraparticle diffusion model ($R^2 = 0.8124$), the straight lines did not pass through the origin, implying that the process of adsorption is controlled by multistep mechanism (Sonal et al. [2020](#page-12-0)). Hence, it can be concluded that the pseudo-second-order adsorption model would be the most appropriate to describe the adsorption kinetic model.

Generally, there are two distinct mechanisms which explain the Cr(VI) biosorption on the bacterial biomass: (i) underlying binding of Cr(VI) ions (i.e., adsorption of Cr(VI)) at the protonated active sites on the surface of the biosorbent through electrostatic interactions and (ii) through a complex Cr(VI) reduction step, by contact with the electron-donor groups of the biomass, resulting in binding of reduced chromium form to the biomass (Albadarin et al. [2011;](#page-10-0) Park et al. [2006\)](#page-11-0). In our present study, the integrated analysis of models and the experimental and instrumentational data, it has been observed that biosorption of Cr(VI) on the biomass adsorbent has been accomplished through more than one step mechanism, including active electrostatic attractions of Cr(VI) with the protonated sites (such as amino, carboxyl) along with the reduction of Cr(VI) by reductive groups (such as hydroxyl, carbonyl) on the biomass surface. Ramrakhiani et al. [2011](#page-11-0) studied the biosorption mechanism of Cr(VI) by heatinactivated fungal biomass of Termitomycesclypeatus and reported that biosorption of Cr(VI) on the biomass involved more than one mechanism.

Table 5 Kinetic parameters obtained from pseudo-first-order and pseudo-second-order and Intraparticle diffusion for Cr(VI) biosorption by Bacillus species strain DPAML065

Metal	Biosorbent	Initial conc. $q_{e \text{ exp}}$ (mg/L)	(mg/g)	Pseudo-first order		Pseudo-second order		Intraparticle diffusion			
				Λ_1 $(min^{-1}$			q_e cal. (mg/g) R^2 K_2 (g mg ⁻¹ min ⁻¹) q_e cal. (mg/g) R^2		C		R^2
	$Cr(VI)$ Bacillus species 80		76.071 0.301		78.41		0.921 0.0035	82.64	0.984 13.482 1.8338 0.8124		

Conclusion

In the present study, we have explored chromium biosorption and the involved mechanism between chromium and Bacillus strain DPAML065 derived from the tannery sludge. Maximum Cr(VI) reduction (95%) was observed at pH 6.0 with Cr(VI) concentration of 80 mg/L at a retention time of 24 h and operational temperature of 35 °C. Biosorption of Cr(VI) by Bacillus strain was well fitted with Langmuir adsorption isotherm, and the maximum adsorption capacity (q_{max}) was calculated to be 106.38 mg/g. The biosorption kinetics of Cr(VI) was represented with pseudo-second-order kinetics $(R^2 = 0.984)$. FE-SEM and EDX analysis were carried out to find out the biosorption competence of the Bacillus strain DPAML065 as well as to discover the surface adsorption of the metals on the bacterial cell surface. FTIR spectra revealed the participation of carboxyl, hydroxyl and amine groups involved in binding of chromium ions and suggested that ion exchange, precipitation and electrostatic attraction following complex reduction steps are the possible mechanisms during the biosorption of Cr(VI) by the isolated Bacillus strain DPAML065. XRD results confirmed that the peaks at 27.53, 56.65 and 66.45 depict Cr (III) on the surface of the bacterial cell, the plausible reason being precipitation of Cr(VI) during the biosorption process. Hence, this study concludes that Bacillus strain DPAML065 can be used as a low cost and highly efficient biosorbent to decontaminated wastewaters from the leather industries. Additionally, it can be utilized as a promising biosorbent in bioremediation strategies.

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Author contributions Arukula Deepa did all the experiments and the analysis and interpreted the results. Astha Singh interpreted the isotherm study. Aakansha Singh interpreted the biosorption study. Brijesh Kumar Mishra formulated the experimental design.

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Data availability All data generated or analysed during this study are included in this published article (and its supplementary information files)

Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethics approval and consent to participate Not applicable.

Consent for publication Not applicable.

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