

Structural Characterization and Antimicrobial Activity of a Biosurfactant Obtained From *Bacillus pumilus* DSVPI8 Grown on Potato Peels

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Background: Biosurfactants constitute a structurally diverse group of surface-active compounds derived from microorganisms. They are widely used industrially in various industrial applications such as pharmaceutical and environmental sectors. Major limiting factor in biosurfactant production is their production cost.

Objectives: The aim of this study was to investigate biosurfactant production under laboratory conditions with potato peels as the sole source of carbon source.

Materials and Methods: A biosurfactant-producing bacterial strain (*Bacillus pumilus* DSVPI8, NCBI GenBank accession no. GQ865643) was isolated from motor oil contaminated soil samples. Biochemical characteristics of the purified biosurfactant were determined and its chemical structure was analyzed. Stability studies were performed and biological activity of the biosurfactant was also evaluated.

Results: The strain, when grown on modified minimal salt media supplemented with 2% potato peels as the sole carbon source, showed the ability to reduce Surface Tension (ST) value of the medium from 72 to 28.7 mN/m. The isolated biosurfactant (3.2 ± 0.32 g/L) was stable over a wide range of temperatures (20 - 120 °C), pH (2-12) and salt concentrations (2 - 12%). When characterized using high-performance liquid chromatography (HPLC) and Fourier transform infrared spectroscopy, it was found to be a lipopeptide in nature, which was further confirmed by Matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (mass peak 1044.60) and nuclear magnetic resonance (NMR) studies. Data showed that the isolated biosurfactant at the concentration range of 30 - 35 µg/ml had strong antimicrobial activity when tested against standard strains of *Bacillus cereus*, *Escherichia coli*, *Salmonella enteritidis*, *Staphylococcus aureus* and *Paenibacillus larvae*.

Conclusions: Potato peels were proved to be potentially useful substrates for biosurfactant production by *B. pumilus* DSVPI8. The strain possessed a unique property to reduce surface tension of the media from 72 to 28.7 mN/m. In addition, it showed a stable surface activity over a wide range of temperatures, pH, and saline conditions and had strong antimicrobial activity. This potential of the identified biosurfactant can be exploited by pharmaceutical industries for its commercial usage.

Keywords: Surface Active Agents; Surface Tension; Lipopeptides; Anti-Microbial Agents; *Paenibacillus*

1. Background

Biosurfactants are a unique class of structurally diverse compounds, which are amphiphilic in nature and have pronounced surface active properties with a wide variety of applications related to environmental protection, enhanced oil recovery, cosmetic, food processing and pharmaceutical industries (1-6). These compounds have been shown to have several advantages including high biodegradability, biocompatibility, low toxicity and low irritancy over their synthetic counterparts (7). However, a major concern behind the restricted commercial application of these microbially produced biosurfactants is their high production costs (6, 8, 9). This cost can be reduced by utilizing alternative substrates, such as agricultural based industrial wastes, for economical biosurfactant production, and the choice of microorganism, which

plays an important role in biosurfactant production. A wide variety of alternative raw materials, namely various agricultural and industrial byproducts, have been evaluated for biosurfactant production (8-11). Keeping the above factors into account, modified minimal salt medium supplemented with potato peels was selected and used as the sole carbon source for production of biosurfactant. India being the third largest producer of potato in the world (10) generates large quantities of potato peels, which are generally attained by steam, abrasive or lye peeling by potato processing industries.

2. Objectives

In the present investigation the ability of a bacterial isolate to utilize potato peels, as a cost effective substrate

for biosurfactant production, was examined. Extraction, purification and characterization of the isolated biosurfactant were done to ensure the role of the isolated biosurfactant as a potent antimicrobial agent.

3. Materials and Methods

3.1. Screening and Identification of Biosurfactant-Producing Strains

A biosurfactant-producing microbial strain was isolated from motor oil contaminated soil samples, obtained from a warehouse (Roorkee, India), based on surface active measurements (12). Briefly, surface tension of the cell-free broth (50 mL) of the culture supernatant, obtained at different time intervals, was measured by a tensiometer (Sigma 703 KSV instruments Ltd., Finland), using the Wilhelmy plate measurement technique. All the measurements were done in triplicates. The selected bacterial strain (DSVP18) from the above evaluation tests was identified by microscopic and conventional biochemical tests, in accordance with Bergey's Manual of Systematic Bacteriology (13).

The strain was identified as *Bacillus pumilus*, DSVP18 (NCBI GenBank accession no. GQ865643), using morphological, biochemical and 16S rRNA sequence analysis. Furthermore, identification of the selected bacterial isolate DSVP18 was performed using standard 16S rRNA-specific universal primers (Genescript, India), with a forward primer (5'-AGAGTTTGATCCTGGCTCAG-3') and reverse primer (5'-AAGGAGGTGATCCAGCCGCA-3'). A reaction mixture containing approximately 50 ng of template DNA, Polymerase Chain Reaction (PCR) buffer (10 mM Tris-HCl, pH 8.3; 50 mM KCl; 2.5 mM MgCl₂; and 0.001% gelatin), a 0.2 mM concentration of each PCR primer, 0.2 mM concentration of each dNTP, and 2.5 U of Taq DNA polymerase in a total volume of 50 µL was prepared. Polymerase Chain Reaction conditions were as follows; initial denaturation at 95°C for two minutes, followed by 30 cycles of denaturation at 95°C for one minute, annealing for one minute at 58°C, and extension at 72°C for one minute. At completion of the PCR reaction, 5 µL of the PCR product was run on a 1% agarose gel containing 1 mM ethidium bromide for visualization of DNA in the amplified bands on a UV illuminator (12). The partial sequences were subjected to BLAST analysis available at www.ncbi.nlm.nih.gov/BLAST. The taxonomic position and relationship of the isolated strain, *B. pumilus* DSVP18, was determined using MEGA 6.0 by constructing a phylogenetic tree (14).

3.2. Medium and Growth Conditions

Biosurfactant production by *Bacillus pumilus* DSVP18 ($OD_{600nm} = 0.8$) was carried out using modified minimal salt medium (MSM) supplemented with potato peels (2%) as the sole carbon source under optimized growth conditions at 37°C for 10 - 12 hours. The composition of the used

mineral salt medium (MSM) (g/L) was follows: (NH₄)₂SO₄ (3.0), NaNO₃ (2.0), K₂HPO₄ (4.5), KH₂PO₄ (2.0), NaCl (0.1), MgSO₄ · H₂O (0.01), CaCl₂ (0.01) and FeSO₄ · 7H₂O (0.01) with pH adjusted to 7.0 (15). All experiments were carried out in triplicates.

3.3. Substrate Preparation and Compositional Analysis

Potato peels were obtained as waste material from a potato chips factory located in Haridwar, India. They were dried in an oven at 60°C and ground in a hammer mill. The composition of potato peels, before use for experimental work, in terms of percentage protein (16), carbohydrate (17), sugar (18) and ash contents (19) are mentioned in Table 1.

Table 1. Chemical Composition of Potato Peels

Composition	Content, %
Total Dry Matter	18.6
Carbohydrate	58.2
Crude Protein	13.6
Fat/lipid content	2.2
Ash content	7.4
pH	5.9

3.4. Dry Biomass Estimation

Dry biomass of bacterial growth was estimated at different time intervals (0, 24, 48, 72 and 120 hours) by centrifuging the samples (250 mL) at 16300x g for 25 minutes. The cell pellet obtained after centrifugation was dried overnight at 105°C and weighed (20).

3.5. Extraction and Purification of the Biosurfactant

The biosurfactant was extracted from the cell free broth using acid precipitation techniques, as described by Yakimov et al. (21). Briefly, biosurfactant precipitates were formed when cell free broth samples, obtained at different time intervals, were adjusted to pH 2 using 6N HCl and kept overnight at 4°C. Cell free broth was then centrifuged at 10 000x g for 15 minutes, and the obtained precipitate was collected and suspended in a minimal amount of distilled water, which was further neutralized to pH 7.0 using 1 N NaOH. The solution was then lyophilized, weighed and stored at -20°C (21).

3.6. Determination of Protein and Lipid Content

The total protein content of the biosurfactant was determined according to the methods described by Lowry et al. (16), using bovine serum albumin as the standard. Total lipid content was determined as described by Arutchelvi et al. (22).

3.7. Chromatographic Analysis

Thin Layer Chromatography (TLC) analysis of the isolated biosurfactant was done on silica gel (Si 60 F254, 0.25 mm, Merck) using chloroform-methanol-water 65:25:4 (v/v/v) as the solvent system. Spots were revealed by spraying with distilled water and heating at 110°C for five minutes, for detection of hydrophilic compounds. Surfactin (Sigma-Aldrich, USA) was used as the standard. HPLC of purified biosurfactant was performed using a 30 cm C18 μ -Bondapak column (Waters) at 25°C and separated using reverse phase HPLC. The used mobile phase, constituted of 80% acetonitrile with 0.1% trifluoroacetic acid and a flow rate of 1 mL/minute. The eluted peaks were detected at 210 nm (21).

3.8. Spectroscopic Analysis

3.8.1. Fourier Transformation Infra-Red (FTIR) Analysis

Characterization of purified biosurfactant was performed using a **Fourier Transformation Infra-Red (FTIR)** spectrophotometer (Thermo-Nicolet, USA) equipped with OMNIC software for data analysis. A pinch of isolated biosurfactant was ground with 100 mg of KBr in pestle and mortar for 30 seconds to obtain translucent pellets. The FTIR spectra was obtained at a frequency range of 4000 - 500 cm^{-1} (23).

3.8.2. Matrix-Assisted Laser Desorption/Ionization Mass Spectrometry Analysis

Furthermore, 0.5 μL of purified biosurfactant was spotted onto the anchor chip position on a Matrix Assisted Laser Desorption/Ionization (MALDI) plate, and 0.5 μL of the matrix was added to the sample spot. The matrix was a saturated solution of 2, 5-dihydroxybenzoic acid in water and 0.3 mg/ml α -cyano-4-hydroxycinnamic acid in acetone/ethanol (2:1 v/v). A peptide standard was also spotted for external calibration. The spots were left at room temperature to dry and analyzed on Applied Biosystems Voyager System 4402 Mass Spectrometer with the delayed mode and an acceleration voltage of 20 kV.

3.8.3. Nuclear Magnetic Resonance Spectroscopy Analysis

One-dimensional ^1H nuclear magnetic spectra were recorded on 298 K on a 500MHz NMR spectrophotometer (Bruker, Germany). The samples were prepared as solutions in 100% CDCl_3 , using approximately 1-3 mg of the biosurfactant (12).

3.9. Properties of the Biosurfactant

3.9.1. Effect of pH, NaCl and Temperature on Biosurfactant Stability

Temperature, pH and the effect of NaCl (%) stability on Surface Tension (ST) values of crude biosurfactant were determined (24, 25). Briefly, ST was measured by a tensiometer

(Sigma 703 KSV instruments Ltd., Finland) using the Wilhelmy plate measurement technique, and for evaluating the critical micelle dilutions, the cell free broth was diluted ten-fold (CMD^{-1}) and 100-fold (CMD^{-2}) with distilled water. The ST of these solutions was then plotted against time. To check thermal stability, 50 mL of the crude biosurfactant aq. solution (0.1%) was incubated for five hours at different temperature ranges from 20 - 120°C, followed by determination of surface activities (ST, CMD^{-1} and CMD^{-2}). The pH stability studies of crude biosurfactant aq. solution (0.1%) were done by adjusting the pH range from 2 - 12 and keeping it for 24 hours before the measurement of surface activities. The effect of salt concentration on surface active values was determined by dissolving 2 - 12% (w/v) NaCl in the crude biosurfactant aq. solution (0.1%).

3.9.2. Antimicrobial Assay of Purified Biosurfactant and Determination of Minimum Inhibitory Concentration (MIC)

Antimicrobial tests of the purified biosurfactant (50 $\mu\text{g}/\text{mL}$) were performed against standard strains, such as *Bacillus cereus* (MTCC 430), *Escherichia coli* (MTCC 1687), *Salmonella enteritidis* (MTCC 3219), and *Staphylococcus aureus* (MTCC 5021), procured from the Microbial Type Culture Collection (MTCC, IMTECH, Chandigarh, India) and *Paenibacillus larvae* (ATCC 9545), respectively. The antibacterial activity of the purified biosurfactant was determined using a growth inhibition zone assay (23). Briefly, overnight cultured bacterial strains with a standardized inoculum size of 1.5×10^8 CFU/mL were inoculated on surface of nutrient agar plates, separately. Purified biosurfactant at varying concentrations (20 - 50 $\mu\text{g}/\text{mL}$) was loaded into wells of these nutrient agar seeded plates with respective bacterial inoculum. The plates were incubated for 24 hours at 37°C. The MIC value was defined as the lowest concentration of biosurfactant that inhibited the visible growth of microorganisms after overnight incubation. The MIC values, of purified biosurfactant showing maximum zone of inhibition when tested on the above-mentioned microbial strains, were recorded. A negative control was prepared using the same solvent as employed to obtain the extract. Ofloxacin (5 μg , Sigma-Aldrich, USA) and cefoperazone-sulbactam (5 μg , Sigma-Aldrich, USA) were used for Gram-positive and Gram-negative bacteria, respectively. All the above-mentioned experiments were conducted in triplicates.

4. Results

4.1. Identification of Biosurfactant-Producing Strain

Among 78 bacterial isolates obtained from oil contaminated soil samples, strain DSVPI8 was chosen based on its ability to reduce ST of the medium from 72 to 28.7 mN/m. This bacterial isolate (DSVPI8) was then subjected to biochemical tests using Bergey's manual of systematic bacteriology, and it was found to be a Gram-positive, motile, rod-

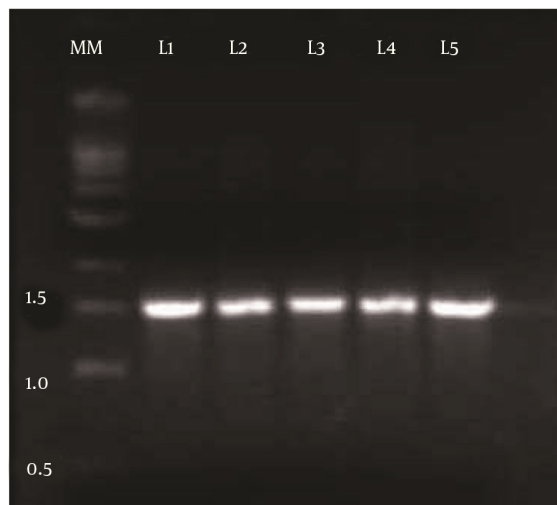
shaped and endospore-forming bacterium. This isolate also showed positive results for urease, citrate utilization and for IMViC test. Genomic DNA was isolated and an amplified product of 1.5 Kb was obtained (Figure 1). Sequence alignment of the fragment using BLAST revealed 100% identity to *B. pumilus* JS-45 16S rRNA gene partial sequence.

The physiological and biochemical characteristics, and the 16S rRNA sequence analysis confirmed that the strain was DSV18 belonging to *Bacillus pumilus* (NCBI Gen Bank accession no. GQ865643). The isolate's evolutionary history was inferred using the Neighbor-Joining method. The optimal tree, with the sum of branch length equal to 0.02602863, was constructed (Figure 2). The percentage of replicate trees in which the associated taxa clustered together in the bootstrap test (500 replicates) has been shown next to the branches (14). The tree was drawn to scale, with branch lengths in the units of the number of base substitutions per site. The evolutionary distances were computed for 19 nucleotide sequences using the Kimura 2-parameter method. There were a total of 1536 positions in the final dataset. Evolutionary analyses were conducted using the MEGA6 software (14).

4.2. Biosurfactant Extraction

Biosurfactant was extracted from *Bacillus pumilus* DSV18 using the acid precipitation technique. Data showed maximum biosurfactant level of 3.2 g/L obtained after 48 hours of culture incubation at 37°C with maximum dry biomass (1.3 g/L). The extracted biosurfactant was also capable of reducing the ST of modified MSM medium from 70 to 28.7 mN/m when 2% potato peels were used as the sole carbon source while critical micelle dilutions (CMD⁻¹ and CMD⁻²) were recorded to be 36.4 and 33.0 mN/m, respectively (Figure 3).

Figure 1. Polymerase Chain Reaction Amplified a Product of 1.5 kb Corresponding to 16S rRNA Gene



MM, Molecular Marker (1 kb DNA ladder); L1, DSV13; L2, DSV14; L3, DSV18; L4, DSV25; L5, DSV26.

4.3. Characterization of Biosurfactant

The chemical composition of the biosurfactant showed that it was composed of 14.7% protein and 18% lipid. The proportion of the surfactant ratio of protein and lipid remained constant in the acid precipitate at different

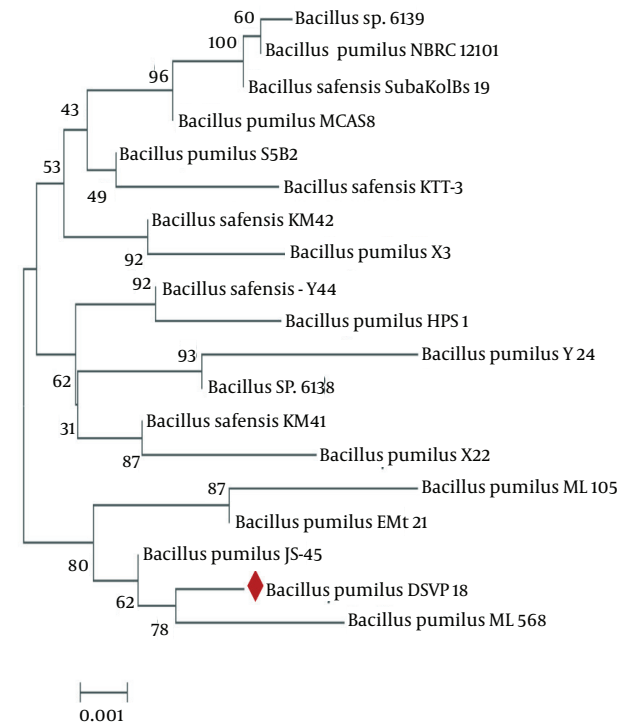


Figure 2. The Phylogenetic Tree of *Bacillus pumilus* DSV18 Constructed Using the Neighbor-Joining Method (MEGA 6 Software)

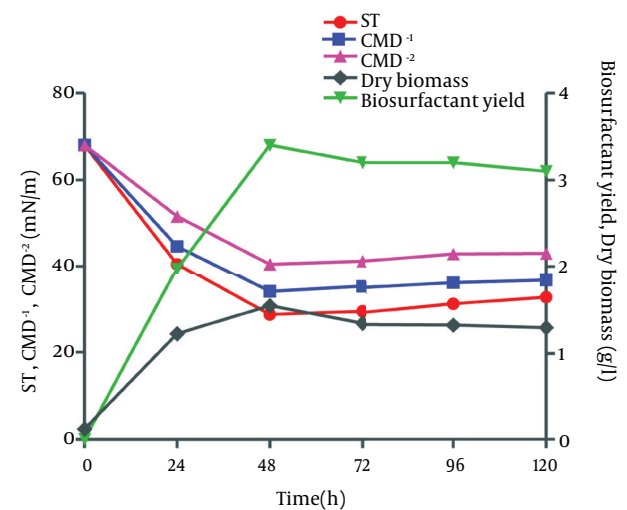


Figure 3. Surface Tension (ST), Critical Micelle Dilutions (CMD⁻¹, CMD⁻²), Dry Biomass and Biosurfactant (BS) Production by *B. pumilus* DSV18 at Different Time Intervals

periods of harvesting. *Bacillus pumilus* produced a lipopeptide, which was stable when checked for surface activities (ST, CMD⁻¹ and CMD⁻²) at a wide range of pH (2.0 to 12.0), temperatures (20 - 120°C) and 2% to 12% salt conditions (Table 2). HPLC analysis of the partially purified biosurfactant showed six major peaks at retention times: 13.65, 19.18, 20.58, 22.61, 25.62, and 26.37, respectively (Figure 4). Fourier Transformation Infra-Red spectrum of the biosurfactant from *Bacillus pumilus* DSVP18 (Figure 5) showed strong absorption bands of peptides at 3405 cm⁻¹, 1604 cm⁻¹ and 1541 cm⁻¹ resulting from N-H stretching, C = O stretching and combined C-N stretching modes. The predominant adsorption bands were 1423 cm⁻¹ and 2964 cm⁻¹, which indicated the presence of aliphatic chains (CH₂, CH₃) in the sample. The intense band at 1604 cm⁻¹ corresponds to the CO-NH-R group. The absorption region at 1730 cm⁻¹ was due to an ester carbonyl band. Data obtained from MALDI-TOF-MS of purified biosurfactant showed well-resolved groups of peaks at m/z values 1000 - 1096 (Figure 6). The mass peaks at m/z 1047.6, 1030.5, 1044.6 and 1058.6 indicated a lipopeptide with mixture of structural analogs. Mass spectrometry confirmed isomers of surfactin and iturin in purified fractions. The mass peak at 1044.60 was attributed to a surfactin isoform containing a β-hydroxy fatty acid with a chain of 14 carbon atoms. Figure 6 shows the assignments of peaks obtained from MALDI-TOF-MS.

Table 2. Surface Activities of Isolated Biosurfactant Aqueous Solution (0.1%) at Different pH (2 - 12), Temperature (2 - 120 °C) and NaCl Concentrations (2 - 12%)

Parameters	Surface Tension, mN/m	CMD ⁻¹ , mN/m	CMD ⁻² , mN/m
pH			
4	34.5	38.2	46.3
6	32.4	36.7	43.8
8	34.6	38.8	46.5
10	36.2	39.4	48.8
12	36.4	39.4	49.2
Temperature, °C			
20	34.7	36.5	41.8
40	30.5	32.5	40.5
60	36.2	36.2	42.0
80	38.0	43.1	51.5
100	42.8	45.0	54.0
120	42.8	43.4	54.0
NaCl, %			
2	36.0	42.1	49.8
4	36.2	42.0	53.0
6	38.8	48.0	56.0
8	45.6	53.4	59.8
10	48.1	55.4	60.2
12	48.4	56.6	60.8

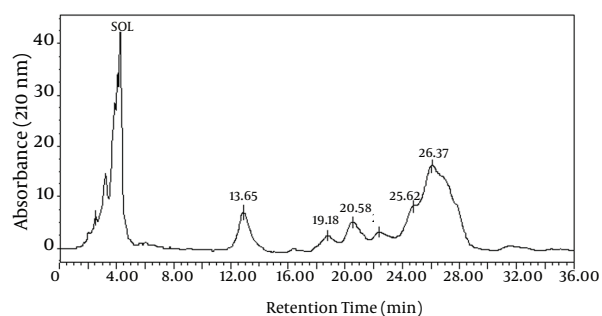


Figure 4. High-Performance Liquid Chromatography (HPLC) Chromatogram of Biosurfactant Obtained From *Bacillus pumilus* DSVP18

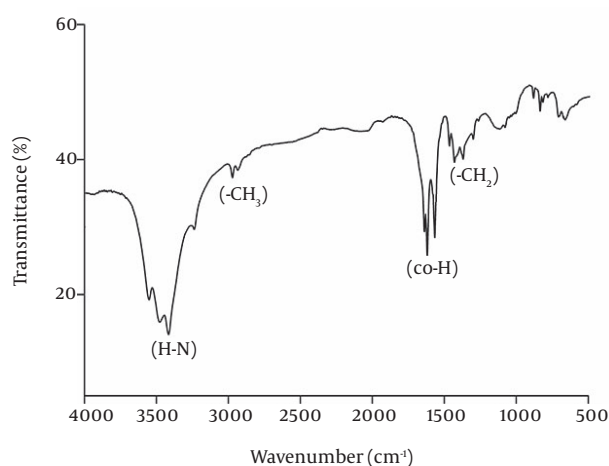


Figure 5. Fourier Transformation Infra-Red Spectrum of Purified Biosurfactant Obtained from *Bacillus pumilus* DSVP18

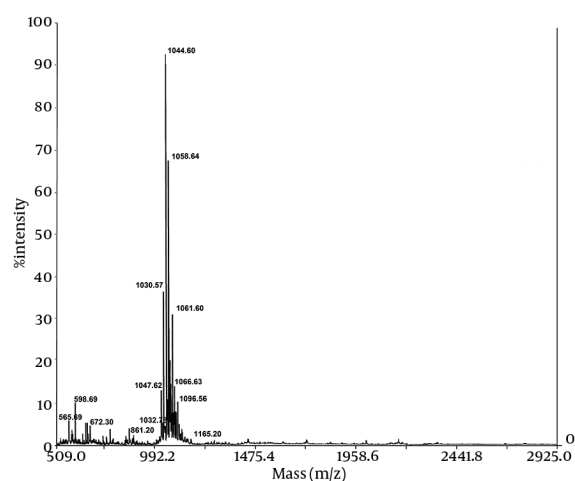


Figure 6. Matrix-Assisted Laser Desorption Ionization-Time of Flight (MALDI-TOF) Mass Spectrometry Spectra of Purified Biosurfactant Obtained From *Bacillus pumilus* DSVP18

Results obtained with 500 MHz for ^1H NMR (Figure 7), clearly indicated that the molecule being studied was a lipopeptide. The spectrum confirmed the presence of a long aliphatic chain (CH_2 at 1.5 - 1.2 ppm) and that the peptide backbone-amide-NH groups were at the region from 7.95 to 7.2 ppm. The doublet signal obtained at 0.859 ppm corresponded to the $(\text{CH}_3)_2\text{-CH}$ group; specifying terminal branching in the fatty acid component. The spectrum indicated the resonance of the following amino acids: valine with peptide CH at 4.6, CH_2 at 1.7, and 2 CH_3 at 0.9 ppm; leucine with peptide CH_2 at 1.7, CH at 1.3 and 2 CH at 0.9 ppm; isoleucine with peptide CH at 4.2, CH at 2.2, CH_2 at 1.9 and 2 CH_3 at 0.9 ppm; aspartic acid or asparagines with peptide CH at 4.7 and CH_2 at 2.8 ppm; and glutamic acid or glutamine peptide CH at 4.3, internal CH_2 at 1.9, and CH_2 α to the carbonyl at 2.1 ppm.

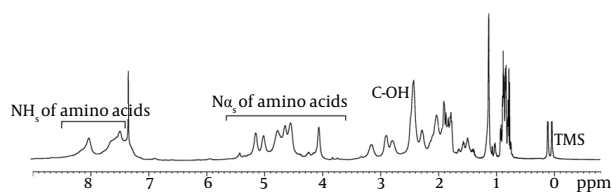


Figure 7. Nuclear Magnetic Resonance Spectra of Purified Biosurfactant Obtained from *Bacillus pumilus* DSVP18

4.4. Antimicrobial Action of Lipopeptide

Lipopeptide produced by *B. pumilus* DSVP18 showed antimicrobial activity against *E. coli*, *B. cereus*, *S. enteritidis*, *S. aureus* and *P. larvae* at different concentrations using the well diffusion method. The MIC value of the lipopeptide varied from 30 to 35 $\mu\text{g}/\text{mL}$. The maximum zone of inhibition recorded with respect to lipopeptide concentration was 30 $\mu\text{g}/\text{mL}$ against *E. coli* (23 ± 0.3 mm) followed by *B. cereus* (19 ± 0.5 mm), *S. enteritidis* (18.6 ± 0.3 mm) and *S. aureus* (18 ± 0.7 mm) while for *P. larvae* (15.3 ± 0.8 mm) this was noted at 35 $\mu\text{g}/\text{mL}$ lipopeptide concentration.

5. Discussion

Potato peels hold a major portion of processing waste and cause severe disposal problems for the potato industry (10). In the present study, these waste products of the potato industry were used for biosurfactant production, as they are a rich source of carbohydrate and protein, which eventually can be used as a potential energy source, thereby alleviating the waste disposal problem (26). Previous researches have made numerous attempts to use potato peels and mushrooms to obtain products, such as enzymes, organic acids, ethanol and polysaccharides with little success (26).

The production of compounds from such agricultural waste requires the selection of microorganisms with relevant enzymatic activities. Most *Bacillus* sp. have a wide

range of hydrolytic enzyme systems and are often capable of utilizing organic matter consisting of complex mixtures (27). In our study, soil samples contaminated with motor oil were collected for the isolation of biosurfactant-producing bacterial strains capable of utilizing potato peels as a cheap carbon source. Amongst various known screening tests for biosurfactant production, such as blood agar lysis, drop collapse assay and emulsification activity, ST reduction assay was chosen for the screening of potential microorganisms that produce biosurfactant (28). The *B. pumilus* DSVP18 isolate was selected for further studies, as this microorganism possessed efficient ST reduction ability and fulfilled the above criteria thus suggesting that it could be an efficient biosurfactant producer. Physiological and biochemical tests and 16S rRNA sequencing allowed the identification of the potential biosurfactant producer, *Bacillus pumilus* DSVP18, at the species level with 100% identity to *Bacillus pumilus* JS-45 from the GenBank database (29).

The selected strain, *Bacillus pumilus* DSVP18, was capable of utilizing potato peels as the sole carbon source. In our study biosurfactant production by *Bacillus pumilus* DSVP18 was growth-associated because a good correlation was observed between the maximum yield of biosurfactant and biomass at 48 hours of incubation (at late log phase) and the surface-active compound was stable over time, which is in good agreement with earlier findings (30). Maximum biosurfactant production (3.2 ± 0.32 g/L) was detected during late log growth phase with the least ST value of 28.7 mN/m. These results suggest that expressions of the gene responsible for biosurfactant production are involved in reducing ST values (31). Previous studies have also reported the maximum yield of surfactin to be approximately 110 mg/l by the *Bacillus subtilis* S 499 strain (32). Similarly, biosurfactant production of 1.74 g/L was observed when the microbial consortium of *Enterobacter cloacae* and *Pseudomonas* sp. (ERCPP1-2) was grown on minimal salt medium supplemented with olive oil (33).

The lipopeptide produced by *B. pumilus* was quantified using HPLC (34), for which six major peaks were attained at the retention time (13.65 - 26.37). The HPLC data strongly suggest that the biosurfactant obtained in this study had a lipopeptide structure, which is in support of previous findings (35). The molecular mass of the purified lipopeptide biosurfactants was attained using MALDI-TOF-MS. The groups of obtained mass spectra resembled that of surfactins and iturins, which represent well-known biosurfactant families produced by *Bacillus* sp. (36). The nature of the biosurfactant was evaluated comparing the FTIR spectra of this fraction to that of pure surfactin. Analysis by HNMR and FTIR of isolated biosurfactant led us to suggest their structural relatedness to lipopeptides. Production of different biosurfactant isoforms by *Bacillus* sp. can be attributed to a set of conserved genes responsible for these enzymes by non-ribosomal synthesis. These changes in structure confer an advantage to the

microorganism to survive under drastic environments. Considering the importance of bioremediation of oil-contaminated sites (oil spills), it is necessary to isolate and screen potent biosurfactant producers, which can survive in harsh conditions.

The high stability of the biosurfactant in a wide pH (2 - 12) range, temperature (20 - 120 °C) and salinity conditions (2 - 12 %), makes it very suitable for these extreme conditions as depicted by Nitschke et al. using cassava wastewater (2). The pH and thermal stability and stability over high saline conditions of the biosurfactants increase the scope of its application in microbial enhanced oil recovery (MEOR) processes. Several lipopeptide biosurfactants produced by *Bacillus* sp. have been noted for their antibacterial activity (23). Lipopeptide biosurfactants are most prominently known as surfactin, produced mostly by the *Bacillus* sp. despite similar global structures; surfactins, iturins and fengycin differ in some aspects regarding their biological activities. The antimicrobial potential of DSVP18 biosurfactant against several standard test pathogens, including *S. enteritidis*, *S. aureus*, *B. cereus*, *E. coli* and *P. larvae* was evaluated in this study. The obtained zone of inhibition clearly represents the potential of DSVP18 biosurfactant as an antimicrobial agent. The response of these microorganisms to biosurfactants may be attributed to their different membrane structures and their relative permeability. Sensitivity towards biosurfactants was in the following order as per their zone of inhibition *E. coli* > *B. cereus* > *S. aureus* > *S. enteritidis* > *P. larvae*. This application potential of the biosurfactant from *B. pumilus* DSVP18 is of great importance as in the future its medicinal use may further be assessed to protect against gastrointestinal pathogenic infections.

The antimicrobial activity increased with increasing concentration of biosurfactant. Our results are in accordance with data for biosurfactants obtained from *B. subtilis* natto (37), *B. circulans* (38) and *B. licheniformis* (39). The biosurfactant from *B. pumilus* DSVP18 showed good inhibitory action against Gram-negative bacteria. This is in contrast to previous reports by Singh and Cameotra (40), where they found the lipopeptide biosurfactant to be active mostly against Gram-positive bacteria having little or no effect on Gram-negative bacteria. A *Bacillus subtilis* isolate showing antagonistic activity against food-borne pathogens was also reported recently (41). In their study, Moore et al. (41) reported three *B. subtilis* strains with proficient biosurfactant activity against test pathogens, i.e. *Salmonella* and *Staphylococcus* cultures, also known as clinical multi-resistant strains. Sabate et al. (42), reported on the antibacterial action of surfactin against *P. larvae*. In their studies they stated that vegetative cells of *P. larvae* were affected as soon as they came in contact with the surfactin.

The current work proposes for the first time the use of potato peels as the sole carbon source for biosurfactant production. Our results are of great importance, since the uniqueness of the compound and the possibility of

cost effective production and stability of biosurfactant makes it a potential candidate for use in environmental applications. The biosurfactant produced by the *B. pumilus* DSVP18 isolate was characterized as a lipopeptide with bioremediation activities under extreme conditions. Purification and study of the mode of action of this low mass compound could lead to the development of an innovative antimicrobial compound in the medical sector. With an alarming increase in drug resistant microorganisms, such new biomolecules can provide an alternative to the existing antimicrobial agents. The potential of this microorganism utilizing other agricultural-based cheap raw materials, for cheaper and large-scale production of biosurfactant, can also be evaluated. Biosurfactant-producing bacterial strain *Bacillus pumilus* DSVP18 (Gene Bank Accession No. GQ865643) screened from oil contaminated soil was identified by 16S rRNA gene sequencing. The strain showed unique ability to utilize potato peels as a carbon source for biosurfactant production and was able to reduce surface tension of the media from 72 to 28.7 mN/m. Structural characterization of the stable (pH, temperature and salt concentration) biosurfactant using HPLC, FTIR, NMR, GC-MS and MALDI-TOF/MAS analysis showed that it was lipopeptide in nature, which possess strong antimicrobial activity. This potential of biosurfactant can be exploited by pharmaceutical industries for its commercial usage.

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Authors' Contributions

All authors supervised and managed the research project. Deepak Sharma, Mohammad Javed Ansari and Sonam Gupta conducted the research. Ahmad Al Ghamdi, Parul Pruthi and Vikas Pruthi helped in writing the manuscript. Deepak Sharma and Mohammad Javed Ansari contributed equally to this work.

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