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Antimicrobial effect of a cyclic peptide Nostophycin isolated from wastewater cyanobacteria, *Nostoc calcicola*

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ABSTRACT

Different types of peptides are produced by cyanobacteria of the genus *Nostoc*, which are unique in structure and have a wide spectrum of biological activities. The objective of the study is to explore different habitats of organism and study antimicrobial activities to improve their pharmaceutical application and drug like properties by structure modification. A cyclic peptide nostophycin was isolated from *Nostoc calcicola* (MK506349) through freeze dried lyophilization method. Its structure has been elucidated using FT-IR, ¹HNMR, ¹³CNMR and LC-MS. Glycine, d-glutamine, l-phenylamine, d-isoleucine, l-proline and a novel amino acid Ahoa are constituents of nostophycin. ¹HNMR, ¹³CNMR spectroscopy confirmed the number of protons and carbons, and characteristics peak determined the structure and fragmentation pattern through LC-MS. Nostophycin possess Ahoa instead of Adha which makes it different from microcystin. Nostophycin exhibits antimicrobial activity against *E.coli*, *S. aureus*, *C. albicans* and *A. niger*. Significant antifungal activity (9-52 µg/mL) and moderate antibacterial activity (concentration 18-52 µg/mL) were observed for nostophycin. In case of already known peptides, these molecules may be further exploited to improve pharmaceutical application and future drug development.

KEYWORDS: Ahoa, Cyclic Peptide, ¹HNMR, ¹³CNMR, LC-MS and Microcystine.

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INTRODUCTION

Cyanobacterial features conspicuous researchers due to their capability of synthesis of various bioactive compounds, diverse range of habitats, wide diversity and morphological variability. Cyanobacteria are gram negative, photosynthetic and ubiquitous bacteria, which known as a primary producer (Gademann & Portmann, 2008). Availability in the extreme environment and unique feature of cyanobacteria, considered it to be future pioneer for research (Kulasooriya, 2011; Potts, 1999; Scherer *et al.*, 1988; Scherer & Potts, 1989). According to Kalaitzis *et al.* (2009) cyanobacteria can produce immense range of bioactive compounds which help in survival in endurance and competitive ecological niche. Bioactive metabolites synthesized by *Nostoc* sp has been applied as a biofertilizer (Ghazal *et al.*, 2018; Win *et al.*, 2018), anticancer (Moore, 1996), antifungal (El-Sheekh *et al.*, 2014), antibacterial (Ploutno & Carmeli, 2000), antiviral (Botos & Wlodawer, 2003) and enzyme-inhibiting (Mazur-Marzec *et al.*, 2018). These bioactive compounds are explored and identified as peptides, alkaloids, terpenoids, fatty acid and lipopolysaccharides (Chorus, 2012; Dembitsky & Řezanka, 2005; Dittmann *et al.*, 2001; Nowruzi *et al.*, 2012; Parmar *et al.*, 2011). Allelochemicals influence their own growth potential,

other microbes in their vicinity, associated microorganisms, higher plants and animals. Cyanobacteria synthesize nitrogen storage material, new proteins, change pigmentation, excrete and store some other compounds, in response to environmental stress, temperature, pH, nutrient availability and light intensity (Mendes & Vermelho, 2013; Priya *et al.*, 2015; Singh, 2014).

High number of metabolites, lipid and lipid like compounds, peptides, oligopeptides and amino acid derivatives produced by different genera of the *Nostocaceae* family (Řezanka & Dembitsky, 2006). Nostocyclamide a macrocyclic peptide was first reported in *Nostoc* sp. (Jüttner *et al.*, 2001) and Nostophycin has been isolated from *Nostoc calcicola*, cyclic peptide with different amino acid including unusual amino acid (Fewer *et al.*, 2011). Extracts from various cyanobacteria have already been proved beneficial for *Triticum aestivum* (Jäger *et al.*, 2005) *Oryza sativa* (Saadatnia & Riahi, 2009), *Zea mays* (Saadatnia & Riahi, 2009), *Cucumis sativus*, *Cucurbita maxima* and *Solanum lycopersicum* (Shariatmadari *et al.*, 2011). Cyanobacteria are known to produce different allelochemicals that have the potential for multiple usages in various fields. Drug-like properties of peptide, combine effect of compounds, pharmaceutical applications of peptides can be improved and

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explored widely. The aim of the present work was to elucidate the chemical structure of extracted compound of *Nostoc calcicola* and examine its antibacterial and antifungal potential.

MATERIALS AND METHODS

Isolation and Identification of Cyanobacteria

Samples were collected from waste water of catchment area of Banjara Lake of Sagar, Madhya Pradesh, India, situated between 23.83° North latitude and 78.73° East longitude and unialgal culture was obtained by streak plate method. *Nostoc calcicola* were cultured in BG 11 medium, pH 6.5 at temperature $25 \pm 2^\circ\text{C}$, light intensity 2500-3000 lux for 14 h light/10 h dark (Rippka *et al.*, 1981). Morphological identification of cyanobacterium was done by microscopic analysis (Micron 36620, Carl Zeiss, Germany) using the keys and description of treaties of Desikachary (1959). Total genomic DNA extraction and 16S rRNA gene amplification were done with standard methodology. The sequence for the species was deposited to the prokaryotic rRNA submission portal of GenBank and the accession number was obtained.

Mass Culture and Crude Extract From Cyanobacterial Samples

Cells were harvested in mid log phase and centrifuged at $10000 \times g$ for 15 min and supernatant was removed and pellet was lyophilized (freeze dried) for 12 h. Furthermore, freeze dried pellet was suspended in methanol and 0.01% TFA (Trifluoroacetic acid), then shaken for 8h by orbital shaker. Later, it was kept overnight at -20°C and further sonicated in a cold room. Subsequently, centrifuged at $10000 \times g$ for 15 minute and supernatant was vacuum dried at 40°C and pellet was re-extracted twice. The mixture was stored in glass vials at -4°C . Yield percentage of extract was calculated by following formula, Yield % = Weight of evaporated extract/ Weight of cyanobacterial powder $\times 100$

SPECTRAL ANALYSIS

Spectral analysis was carried out by freeze dried extracts namely, *Nostoc calcicola* using standard procedures to identify the components.

Fourier Transform Infrared (FTIR) Spectroscopy

Fourier transform infrared spectroscopy (FTIR) performed on Shimadzu 8400S. The spectrum of solid sample was obtained using KBr (Potassium bromide) pellets. About 1 mg of sample and 100 mg of KBr were ground together, dried to remove moisture and mechanically placed in sample holder. The KBr does not absorb infrared radiation in the region 4000 cm^{-1} and 500 cm^{-1} and complete spectrum of the solid sample is obtained (Sharma, 1981).

Nuclear Magnetic Resonance Spectroscopy (NMR)

^1H NMR and ^{13}C NMR spectroscopies were performed by BrukerAvance II at 500MHz, Advanced Instrumentation

Research Facility (AIRF), Jawahar Lal Nehru University, New Delhi using DMSO as a solvent.

Liquid Chromatography- Mass Spectrometry (LC-MS)

LC-MS is sophisticated technique used for separation or purification of sample followed by fragmentation pattern. LC-MS analysis was carried out at Indian Institute of Science and Research (IISER) Bhopal, by Agilent technologies 1260 infinity and Agilent technologies 6130 quadrupole. Methanol was used as a solvent for LC-MS. Standard of amino acids were used as control for control in NMR and LCMS.

Bacterial Strain Preparation

The bacterial strain *Escherichia coli* MTCC#1591 (*E. coli*), *Staphylococcus aureus* MTCC#3212 (*S. aureus*) and fungi strains *Aspergillus niger* MTCC#9652 (*A. niger*), *Candida albicans* (MTCC#183) were obtained from The Microbial Type Culture Collection and Gene Bank (MTCC) Chandigarh, India. The test bacteria were maintained in nutrient broth media while the test fungi were maintained on potato dextrose agar plates. The test organisms were subcultured 2-8 h before the test.

Preparation of Resazurin Solution

The resazurin solution was prepared at 0.02% (wt/vol) with 0.002 g of resazurin salt powder was dissolved in 10 mL of distilled water and vortexed. The mixture was filtered by Millipore membrane filter ($0.2\ \mu\text{m}$). The resazurin solution can be kept at 4°C for 2 weeks.

Disk Diffusion Method

The antibacterial activity of extracted sample against the procured test bacteria was carried out using Kirby-Bauer Disk Diffusion susceptibility Test Method (Hudzicki, 2009). The bacterial strains were spread on Mueller-Hinton agar (Merck, Germany) using sterile cotton swab. Extracted sample was prepared at concentrations of 0.2, 0.4, 0.6, 0.8 and 1 mg/ml. Sterile blank antimicrobial disk was used in the test. The disks were loaded aliquot 5 mL with different concentrations of sample and ampicillin and miconazole was used as positive control for bacteria and fungi respectively, while methanol as a negative control.

For antifungal test, Mueller-Hinton agar supplemented with 2% glucose and $0.5\ \mu\text{g/ml}$ methylene blue was used as a medium. The inoculated plates were incubated at 37°C for 24-48 h and the diameter of the inhibition zone was measured after incubation. All the tests were performed in three replicates, and mean of inhibition zone was calculated.

Determination of Minimum Inhibitory Concentration

Minimum inhibitory concentration (MIC) was determined by using the method described in standard CLSI guidelines (Wayne, 2012). The MIC test was performed in 96 well round

bottom microtiter plate using standard broth microdilution method while MBC was performed on the MHA plates. The bacterial inoculums were adjusted to the concentration of 10^6 CFU/mL. For the MIC test, five dilutions were prepared with 4.5, 6.5, 8.5, 10.5 and 12.5 mg/mL extracted sample in methanol. A total of 50 μ L of the microbial suspension, at 0.5 McFarland concentrations, was inoculated into each well and then the microtiter plate was incubated at 37°C for 24h. The lowest concentration at which no turbidity was observed was considered the MIC. A well of the microtiter plate with no microorganisms was considered as the negative control and with no extract was considered as the positive control.

The MBC was defined as the lowest concentration of the antibacterial agents that completely kill the bacteria. MBC test performed by plating the suspension from well of microtiter plates into MHA plates. The plates were incubated at 37 °C for 24 h. The lowest with no visible growths on the MHA plate was taken as MBC value.

Time-kill Curve

Time-kill assay was done in MHB medium as described by Zainin *et al.* (2013) and Lau *et al.* (2018). The bacterial inoculums were adjusted to 10^6 CFU/mL. The nostophycin solution was diluted with MHB media containing bacterial inoculums to obtain the final concentration of 0×MIC, 1×MIC, 2×MIC, 4×MIC, and 8×MIC for each type of bacteria in the total final volume of 1 mL. The cultures were then incubated at 37°C with 150 rpm agitation. The culture (100 μ L) was spread on MHA plates at time 0, 0.25, 0.5, 1.2 and 4h in triplicates. The number of colonies on the MHA plates was quantified in CFU/mL after incubation at 37°C for 24h. For statistical analysis, SPSS (v.26) statistical package was used to determine the significant ($P < 0.05$) difference among the tested bacteria.

DATA ANALYSIS

The data are presented as mean \pm standard deviation for inhibition zones. For statistical analysis, unpaired t-test was used using SPSS version 26.

RESULT AND DISCUSSION

With the help of microscopic study and camera lucida sketching, morphological characteristics i.e. subspherical heterocyst, barrel shaped cells, and molecular study confirmed it as *Nostoc calcicola* (MK506349) (Figure 1).

SPECTRAL ANALYSIS OF NOSTOC CALCICOLA (NPC) EXTRACTED SAMPLE

FTIR Study

FTIR spectrum was recorded using KBr pellets in range 4000-500 cm^{-1} . The FTIR spectrum of the extract shows various stretching bands, hence characteristics bands were analyzed. Total 6 functional groups were detected at the wave range of

3545, 3250, 2895, 1745, 1550 and 1195 cm^{-1} . The characteristic stretching frequency for carbonyl group attributed to the presence of amide group in sample. Based on the previous literature and standard protocol, the functional classes were characterized as -OH stretching, N-H bond of amine, -C-H stretching carbonyl, C=C (aromatic ring) stretching and C-N stretching respectively (Table 1; Figure 2)

NMR Study

After confirming the presence of carbonyl group in the form of amide group, sample was further characterized by using ^1H NMR and ^{13}C NMR spectroscopic techniques. According to literature survey it was noted that the sample consists of peptide bonds (Fujii *et al.*, 1999). The expected structure of sample was further

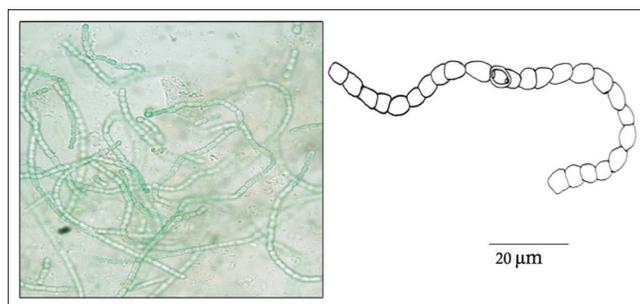


Figure 1: Photomicrograph and camera lucida sketch illustrating the morphological features of isolated *Nostoc calcicola* (MK506349). Bar denotes 20 μm

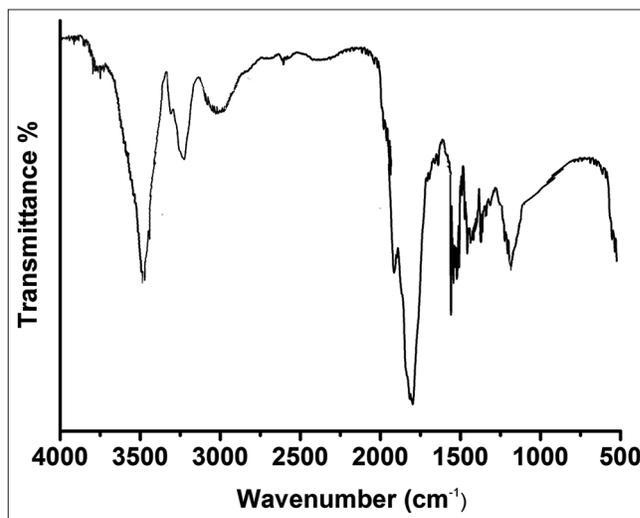


Figure 2: Characteristics IR peaks of NPC

Table 1: Characteristics IR frequencies for NPC.

Serial No.	Frequency (cm^{-1})	Functional Group	Type of Vibration
1	1745	C=O	Stretching
2	3250	N-H	Stretching
3	2895	C-H	Stretching
4	3545	O-H	Stretching
5	1550	C=C (Aromatic ring)	Stretching
6	1195	C-N	Stretching

confirmed by the presence of different types of hydrogen atoms present in it. As the literature reports that the whole structure of sample is constructed with different types of amino acids. The set of amino acids as follows proline I, isoleucine, phenylalanine, proline II, glycine and glutamine. The presence of all amino acids and formation of particular peptide bond were taken into consideration. The obtained ^1H NMR spectrum for sample was compared with ^1H NMR spectra of individual amino acids. The change in chemical shift value of $-\text{NH}_2$ in all amino acids were observed. The chemical shift values obtained in range 6-8 δppm corresponds to proton present in amino acid in the form of $-\text{NH}$ in Nostophycin molecule. The compared values suggest that the signals obtained for amino acids in sample shows larger chemical shift value (δppm) as compared to spectrum obtained for pure amino acids, it may happen due to removal of proton and formation of hydrogen atom (Figure 3).

Some remarkable changes can be mention; the original spectrum of isoleucine displays sharp singlet around 7.05 δppm but in present sample it shows some shift towards higher chemical shift value 7.37 δppm as doublet with $J=8.4\text{Hz}$. Furthermore, phenylalanine and glycine also exhibit the same behavior and their chemical shift values are around 7.39 and 6.89 δppm respectively due to removal of one proton and formation of new bond. Presence of glutamine was confirmed by singlet obtained at 8.7 δppm corresponds to $-\text{N-H}$ proton whereas singlet at 6.60 δppm value corresponds to $-\text{NH}_2$ bond. Additionally one unusual amino acid is determined to be a β -amino acid (3-amino-2,5-dihydroxy-8-phenyloctanoic acid) or (Ahoa), which exhibits two hydroxyl groups in its structure. The ^1H NMR spectrum shows following chemical shifts values 5.76 and 7.52 δppm corresponds to $-\text{OH}$ and $-\text{NH}$ proton respectively. The chemical shift value obtained around 3.62 δppm ($J=8.72$) can be assigned to protons of $-\text{CH}_2$ present in proline amino acid.

The ^{13}C NMR spectrum of sample shows the presence of carbonyl group in structure. The carbonyl group displays its characteristic δppm value around 168-170 δppm . In this context ^{13}C NMR spectrum obtained for sample are in two different ranges. Figure 4 displays characteristics peaks for different types of carbonyl group present in compound. The chemical shift values obtained at 171.1, 170.5 and 168.1 δppm assigned for carbonyl group present in proline, isoleucine and glycine molecule. The singlet obtained for each carbonyl group suggests that no direct proton is attached to the carbon atom. The other singlets obtained at 170.1, 174.2 and 175.2 corresponds to carbonyl group present in glutamine, phenylalanine and 2,5-dihydroxy-8-phenyloctanoic acid.

Liquid Chromatography- Mass Spectrometry (LC-MS) Study

As, it is important to confirm the molecular mass of sample, LC-MS technique was applied to confirm the established structure of sample. Therefore, a confirmation of the amino acids spectra and structure of cyanobacterial peptides by LC-MS at 1000 m/z was carried out (Fujii *et al.*, 1999).

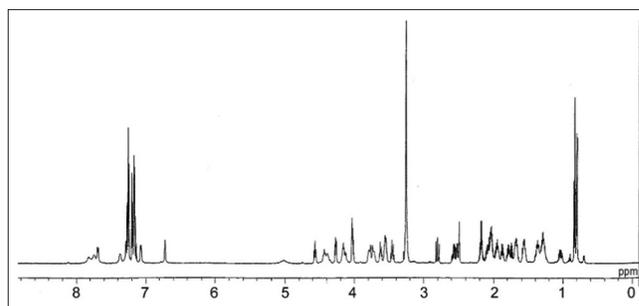


Figure 3: Typical ^1H NMR spectrum of NPC at 500 MHz in DMSO-d_6

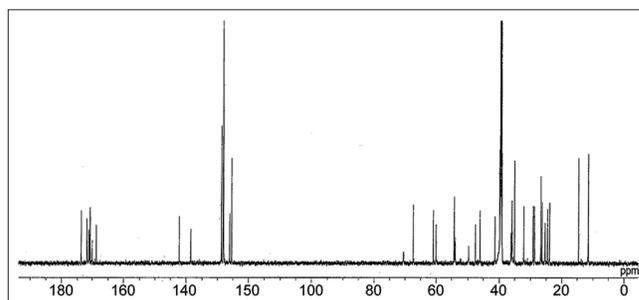


Figure 4: Typical ^{13}C NMR spectrum of NPC at 500 MHz in DMSO-d_6

In the product ion spectrum (Figure 5) for the $[\text{M} + \text{H}]^+$ at m/z 889 of sample, several ions were prominently observed, which were assigned as the ions of each constituent amino acid residue lost from the precursor ion $[\text{M-Glycine}]^+$ at m/z 833, $[\text{M-Isoleucine}]^+$ at m/z 775 and $[\text{M-Phenylalanine}]^+$ at m/z 741, $[\text{M-3-amino-2,5-dihydroxy-8-phenyloctanoic acid (Ahoa)}]$ and $[\text{M-Glycine+glutamine}]^+$ at m/z 532, $[\text{M-Glutamine}]^+$ at m/z 744.

STRUCTURE ELUCIDATION

Based on the results of FT-IR, NMR (^{13}C and ^1H) and mass spectroscopy the molecule was tentatively predicted as cyclic peptide and the structure is composed of six amino acids and a novel β -Ahoa; named as Nostophycin. Molecular formula of the nostophycin was $\text{C}_{46}\text{H}_{62}\text{N}_8\text{O}_{10}$ and molecular weight was predicted as 887. The structure was shown in Figure 6.

Antimicrobial activity of Nostophycin

The antibacterial activity of nostophycin was determined against two bacterial (*E. coli* and *S. aureus*) and two fungal species (*A. niger* and *C. albicans*). The results for disk diffusion test, MIC and MBC of the nostophycin are summarized in Table 2. For the disk diffusion test, the presence of clear zone around the nostophycin disk suggesting that the nostophycin possessed antimicrobial activity which is able to inhibit the growth of bacteria and fungus. As previous study reported that nostophycin extracted from *Nostoc* CCC537 produced a maximum inhibition zone (29 mm) for *S. aureus* followed by *P. aeruginosa* (11 mm), *S. typhi* (10 mm), *E.coli* (10 mm) and *Enterobacter aerogens* (7 mm), it has also been reported that doubling in the concentration increased the inhibition zone for

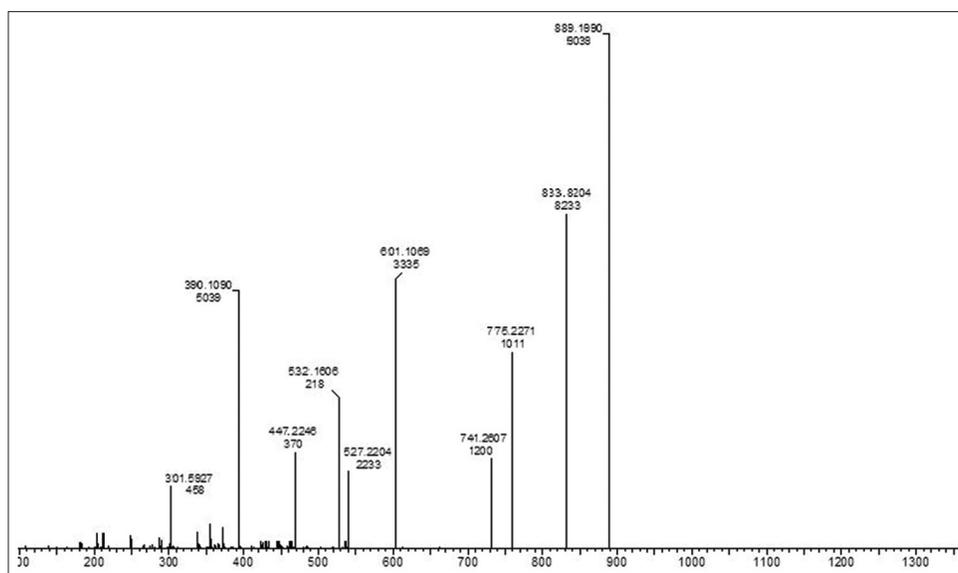


Figure 5: Partially purified molecule analyzed by LC-MS chromatograms at m/z 889 for Nostophycin (NPC) in Methanol

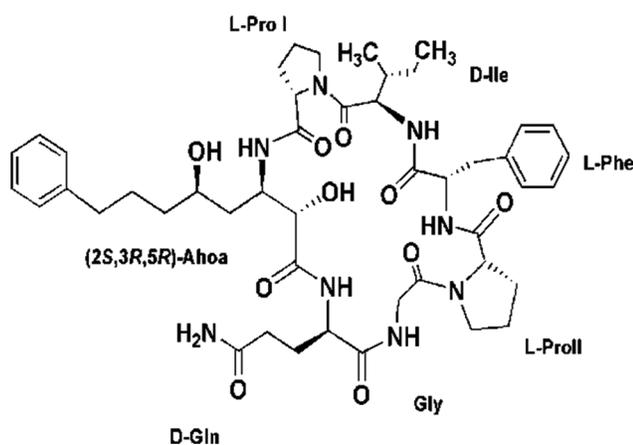


Figure 6: Proposed chemical structure of Nostophycin (NPC) by *Nostoc calcicola* (MK506349) formula $C_{46}H_{62}N_8O_{10}$ and molecular weight 887

the same microorganism (Asthana *et al.*, 2009). Agrawal (2016) also reported antibacterial and antifungal activity of extract of *N. calcicola* against different human pathogens through inhibition zone ranging from 7.5 to 20 mm.

Disk diffusion test was described as the preliminary study in screening the antibacterial activity of an antimicrobial agent; therefore, a further evaluation in determining the antibacterial activity of nostophycin using MIC value was needed. MIC was defined as the lowest concentration of the antibacterial agent to inhibit the growth of bacteria by serial dilution. As showed in Table 2, the MIC values of nostophycin against the bacteria and fungi were ranged from 4.5 to 8.5 mg/mL. It has been reported that low concentration of active principle 2.5 μ g/mL of *Nostoc* CCC 537 was effective against *M. tuberculosis* H37Rv (Asthana *et al.*, 2009). MBC is the lowest concentration of antibacterial and antifungal agent to kill the bacteria and fungus (showed no growth on the agar plate). MBC for *Candida albicans* (4.5) followed by *E.coli* and *A. niger* (6.5) and with maximum for

Table 2: The diameter of inhibition zone (mm), MIC value (μ g/mL), and MBC value (μ g/mL)

Organisms	Inhibition Zone Diameter (mm)	MIC (μ g/mL)	MBC (μ g/mL)
<i>Escherichia coli</i>	22	4.5	6.5
<i>Staphylococcus aureus</i>	38	6.5	8.5
<i>Aspergillus niger</i>	20	6.5	6.5
<i>Candida albicans</i>	24	4.5	4.5

S. aureus (8.5). It has been reported that lower quantity of active principle of *Nostoc* CCC 537 is more effective on high quantity of streptomycin and rifampicin against *Enterobacter aerogens* (Asthana *et al.*, 2009). Pesticidal effects of methanolic extract of *Nostoc* strain ATCC 53789 from 0.25g/L to 100g/L has been used to treat various pathogenic fungi (Biondi *et al.*, 2004). Resazurin dye was used in the study to determine cell growth, especially in cytotoxicity assays (McNicholl *et al.*, 2007). Oxidoreductase within viable cells reduced the resazurin salt to resorufin and changed the color from blue non-fluorescent to pink and fluorescent. According to McNicholl *et al.* (2007), resazurin dye has been applied for decades to check for the bacterial and yeast contamination milk.

TIME-KILL CURVE

The time kill activity of bacteria and fungi is shown in Figure 7. The reduction in the number of CFU/mL effective against bacteria and fungi were ≥ 3 Log units (99%) for bactericidal and fungicidal activity of nostophycin. The bactericidal endpoint of nostophycin for *E. coli* was reached after 2 h of incubation at 4 \times MIC (18 μ g/mL) and 8 \times MIC (36 μ g/mL), while for *Staphylococcus aureus*, the bacteria was killed after 2 h of incubation at 4 \times MIC (26 μ g/mL) and 8 \times MIC (52 μ g/mL). The fungicidal endpoint of nostophycin for *Aspergillus niger* was reached after 2 h of incubation at 4 \times MIC (26 μ g/mL) and 8 \times MIC (52 μ g/mL). *Candida albicans* was killed after 1 h of incubation at 2 \times MIC (9 μ g/mL), 4 \times MIC (18 μ g/mL), 8 \times MIC (36 μ g/mL).

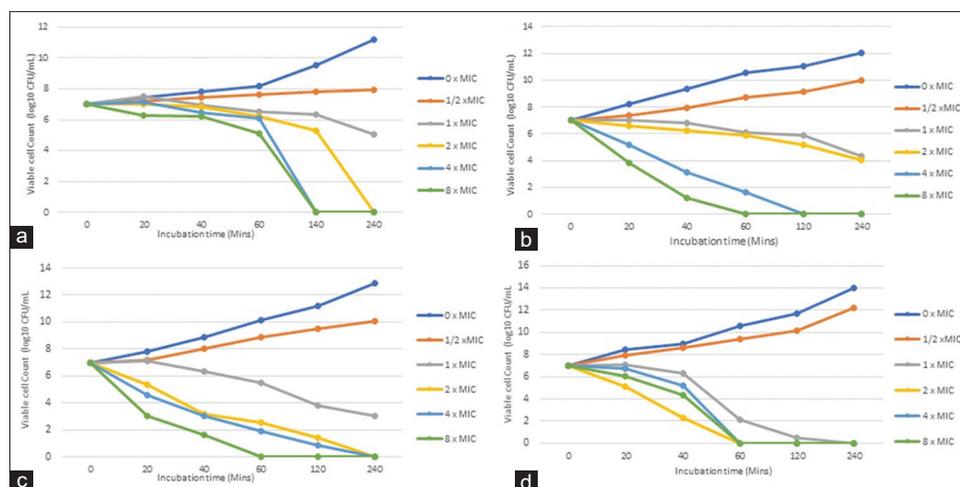


Figure 7: Time-kill plots of Nostophycin against A. *E.coli* B. *S. aureus* C. *A. niger* D. *C. albicans*

Antibacterial activity of *Nostoc* sp. is already reported by Asthana *et al.* (2009), and antifungal activity is reported by Agrawal (2016). Structurally diverse group of Nostophycin exhibits antibacterial activity against gram negative bacteria *E.coli* and gram positive bacteria *S. aureus*, similarly antifungal activity against *A. niger* and *C. albicans*. No significant differences were found between gram positive and negative bacteria, and fungus, which indicates that nostophycin is broad spectrum antimicrobial and antifungal agent. In this study, nostophycin compound extracted from *Nostoc calciola* include a structurally diverse group i.e. cyclic and linear peptides, phenolics and fatty acid. Different bioactive compounds like nostocine A (Hirata *et al.*, 1996) and tenuencyclamide A- D (Banker & Carmeli, 1998) from *N. spongiaeformae*, noscomin (Jaki *et al.*, 1999) and comnostin A-E (Jaki *et al.*, 2000) from *N. commune*, borophycin (Hemscheidt *et al.*, 1994), muscoride (Nagatsu *et al.*, 1995), cryptophycin (Biondi *et al.*, 2004) and nostocarboline (Becher *et al.*, 2005) from *N. linkia*, *N. muscorum*, *N. elliposporum*, *Nostoc* ATCC 53789 and *Nostoc* 18-12A respectively have been reported. Methanol as a control was not able to inhibit microbial activity but methanol extracted nostophycin exhibits antimicrobial activity against bacteria, different workers also adopted methanol extracts for evaluating antibacterial activity (Asthana *et al.*, 2009); Jaki *et al.*, 2000; Mundt *et al.*, 2001). The inherent capacity to produce specific bioactive compounds from methanolic extract of *Nostoc* strain ATCC 53789 seems to play an antifungal role against a variety of pathogens of different agricultural importance fungi such as *Armillaria* sp., *Colletotrichum coffeanum*, *C. trifolii*, *Fusarium solanii*, *F. oxysporum* f. sp. *melonis*, *Penicillium expansum*, *Phytophthora cambivora*, *P. cinnamomi*, *Rhizoctonia solanii*, *Rosellinia* sp. *Sclerotinia sclerotiorum* and *Verticillium albo-atrum* (Biondi *et al.*, 2004). Various *Nostoc* species behave as an antifungal agent such as extract of *N. linkia* inhibit the growth of wilt disease causing *Fusarium oxysporum* f. sp. *lycopersici*, which help in biological control and better yield for tomato plants (Alwathnani & Perveen, 2012). Extract of *Nostoc commune* FA-103, *Nostoc endophytum* and *Nostoc muscorum* suppress the effect of *Fusarium oxysporum* f. sp. *lycopersici* and soyabean root rot causing *Rhizoctonia solanii* respectively (Ismail & Ismail, 2011; Kim & Kim, 2008).

CONCLUSION

Cyanobacterial cyclic peptide was isolated and identified from *Nostoc calcicola* through ¹³CNMR, ¹HNMR and LC-MS and named as nostophycin, it contains Gln, Gly, Pro I, Pro II, Phe, Ile and Ahoa. Nostophycin exhibits resemblance with microcystin with all variables except amino acid ADHA ((2S,3S,8S,9S)-3-amino-9-methoxy-2,6,8-trimethyl-10-phenyldeca-4,6-dienoic acid), which is replaced by Ahoa ((2S,3R,5R)-3-amino-2,5-dihydroxy-8-phenyloctanoic acid). The β-amino acid Ahoa from nostophycin is structurally quite similar to (Ahda) in scytonemin. Similar β-amino acids are also reported in several peptides isolated from cyanobacteria. But unusual β-amino acid Ahoa is only found in nostophycin (Fewer *et al.*, 2011). Cyclic peptides from natural resource exhibits a variety of significant biological profiles. The anti-microbial activity of cyanobacterial secondary metabolites has been reported by many researchers. However, the MIC values from previous study showed a large variation. Therefore, the comparison of the result is difficult as there is no standard method for determination of antimicrobial activity of nostophycin. In this study, nostophycin exhibits a good antimicrobial activity against gram positive and gram negative bacteria and fungi. Lower concentration of nostophycin is able to kill bacteria and fungi in a short time, the reproduction time of bacteria is main cause of infection and treating with nostophycin could be a viable way to prevent infections. Unique structure of nostophycin and a wide spectrum of biological activity shows a remarkable biotechnological potential of the *Nostoc*. Newly discovered or already known bioactive peptides can be used to improve drug-like properties through structure alteration or conjugation with antibodies or small molecules. It is necessary to explore chemical synthesis, exact mechanism of secondary metabolites against bacteria and fungi and search for new chemicals from cyanobacteria.

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