

**CYANOBACTERIA: A POTENT SOURCE OF NOVEL BIO-ACTIVE COMPOUNDS****Richa Gupta<sup>1</sup>, Keshawanand Tripathi<sup>2</sup>, Dinesh Kumar Prajapati<sup>3</sup> and Pankaj Kumar Rai<sup>4\*</sup>**<sup>1, 2, 3, 4</sup>Department of Biotechnology, Invertis University, Bareilly-243123, India; (R.G.); (K.T.); (D.K.P.)

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**ABSTRACT**

*Cyanobacteria are a rich source of novel bioactive compounds with a broad spectrum of biological activities. Cyanobacteria produce many biologically active chemicals, although only a small fraction of these molecules have been identified and characterized in the last several decades. They exhibit a range of bioactivities, including antimicrobial, anticancer, antiviral, and antimalarial. They synthesized non-ribosomal peptides through non-ribosomal peptide synthetase (NRPs), which are multimodular enzyme complexes via ribosome-independent pathways. Several methods have been standardized for isolation, purification and characterization of cyanobacterial bioactive compounds. This paper provides a comprehensive review of cyanobacterial bioactive compounds, including antibacterial, antiviral, antiprotozoal, and anticancer compounds, as well as biosynthesis and extraction techniques from cyanobacteria, in response to the high demand for novel bioactive compounds.*

*Keywords: Cyanobacteria, Antibacterial, Antiviral, Antiprotozoal, Anticancer, Bioactive compounds.*

**1. INTRODUCTION**

Drug resistance is a serious problem among organisms, including methicillin-resistant *Staphylococcus aureus* (MRSA) (Reynolds et al. 2004; Qureshi et al. 2021b; Qureshi et al. 2021a; Qureshi et al. 2022), penicillin-resistant *Streptococcus pneumoniae* (Karchmer 2004), and *Pseudomonas aeruginosa* (Paterson 2006). This indicates that conventional antibiotics have lost efficacy, necessitating their substitution with newer medications. These factors together lead to the search for novel drugs and may be new sources, as many unexplored bacteria create powerful bioactive compounds. Only 1-10 % of cultivated bacteria ( $2 \times 10^5$ ) have been studied (Cowan 2000). It is anticipated that uncultured bacteria and the vast microbial diversity would provide additional sources of antibiotics. (Clardy et al. 2006).

Microorganisms, including actinobacteria, bacteria and fungi, have been explored widely over many decades, and many drugs have been originally isolated and characterized. Among the microorganisms, actinomycetes produced a tremendous amount of bioactive compounds of diverse nature (Selim et al. 2021). Bioactive compounds are secondary metabolites that do not function in the growth or reproduction of living organisms (animals, plants and microbes) (Yang et al. 2016). They have a wide range of chemical and structural diversity, such as steroids, terpenoids, alkaloids, polyketides, phenolic compounds etc., and exhibit diverse biological activity (Syta and Smetanska 2022). For the development of pharmaceutical products, this is the most fruitful source. Approximately one hundred innovative products are now being evaluated in clinical trials, mostly as anticancer and anti-infective therapies (Harvey et al. 2015). Adoption of sophisticated techniques increased the probability that novel drugs derived from bacteria, actinomycetes, or yeasts will be more broadly accessible (Newman and Cragg 2012; Harvey et al. 2015). The bulk of pharmaceuticals on the market are derived from actinomycetes or fungi.

Cyanobacteria are photosynthetic, oxygen-evolving, Gram-negative prokaryotes found in every environment, including freshwater, saline and most adverse conditions (Ward et al. 2012; Yadav et al. 2022). They are found in different environmental conditions, from symbiotic, aquatic, saline, deserts, terrestrial and hot springs (Whitton and Potts 2012). Cyanobacteria contain some specialized structure, heterocyst, and the ability to fix atmospheric nitrogen. However, few cyanobacteria, mainly marine nitrogen-fixing cyanobacteria, are capable to fix molecular nitrogen without heterocyst (Díez et al. 2008). The cyanobacteria's dispersion and colonization occur by a specialized structure known as hormogonia (Meeks et al. 2002).

Cyanobacteria are promising organisms that produce novel biochemically active chemicals known as bioactive compounds (Singh et al. 2005; Gayakwad et al. 2015; Singh et al. 2017; Begum et al. 2022; Singh et al. 2022). Several genera of cyanobacteria, including *Limbia*, *Microcystis*, *Oscillatoria*, and *Nostoc*, can synthesize bioactive chemicals (Selim et al. 2021). They potentially produce bioactive molecules essential to their ecosystems. Recent literature shows that 90 genera of cyanobacteria have synthesized bioactive compounds. Most of these organisms belong to the orders *Chroococcales*, *Synechococcales*, *Oscillatoriales*, and *Nostocales*. On the other hand, *Pleurocapsales*, *Chroococciopsale* and *Gloeobacterale* are unexplored or partially explored (Demay et al. 2019). They mainly exhibited different types of activity such as antibacterial (Reichelt and Borowitzka 1984; Cannell et al. 1987), antifungal (Kellam and Walker 1989; Hagmann and Jüttner 1996), anticancer (Carmeli et al. 1990; Kashiwagi et al. 2005), antiviral (Starr and Moran 1962; Gupta et al. 2014), and antiprotozoal compounds (Broniatowska et al. 2011).

The bioactive molecules formed within the cell biomass are referred to as end metabolites. Exo-metabolites, on the other hand, are bioactive compounds that are produced in reaction to the environment (De Morais et al. 2015). Among the several bioactive secondary endo-metabolites identified from cyanobacteria is the cytotoxic lyngbyabellin from *Lyngbya majuscula* (Han et al. 2005), pahayokolide A (anticancer) from *L. semiplena* (Berry et al. 2017), hapalindole series (antituberculosis) (Mayer et al. 2016), venturamide A, B (antimalarial) from *Oscillatoria* sp. (Linnington et al. 2008) and antimalarial linear lipopeptides from the marine *L. majuscula* (McPhail et al. 2007).

### 1.1 Antimicrobial compounds

Chemicals that kill or inhibit microorganisms such as bacteria, fungi, and protozoa are known as antimicrobial compounds. Antimicrobial drugs are selective, killing or inhibiting the growth and proliferation of bacteria (Hood and Khan 2020). Secondary metabolites are produced at a certain cell age and are affected by biotic and abiotic stimuli (Ramakrishna and Ravishankar 2011). The nature of the substances varies depending on the circumstances, e.g. antibacterial in the case of bacterial pathogens, antiviral, antiprotozoal, etc. Cyanobacteria are the richest source of antibacterial compounds that belongs to a diverse chemical class, including alkaloids, fatty acids, and terpenoid. Nonetheless, a significant variety of additional substances, including macrolides, peptides, aromatic compounds, indole paracyclophanes, and others, have also been identified (Rojas et al. 2020).

Table 1 lists the antimicrobial compounds found in cyanobacteria and their biological effects. Ambiguine isonitrile A is an indole alkaloid isolated from *Fischerella ambigua* (UTEX 1903) and found to have antibacterial activity against *Mycobacterium* and *Bacillus anthracis* (Jaki et al. 2000; Mo et al. 2009). The terrestrial cyanobacterium *Nostoc commune* excreted diterpenoid molecule in the extracellular medium, identified as Noscomin, exhibits antibacterial activity against *Staphylococcus*, *Escherichia coli* and *Bacillus cereus*. *Fischerella ambigua* synthesized and secreted spent medium. Characterization of the bioassay-directed fraction parsiguine displayed antibacterial and antifungal properties (Ghasemi et al. 2004). Hapalindole-T was isolated from an organic solvent extract of *Fischerella* sp. extracted from *Azadirachta indica* (Neem bark) and showed antibacterial activity against *Mycobacterium tuberculosis*, *Enterobacter aerogenes*, *Staphylococcus aureus*, *Pseudomonas aeruginosa*, *Salmonella typhi*, and *Escherichia coli* (Asthana et al. 2006).

**Table 1.** Antimicrobial compounds synthesized by cyanobacteria

| S.N | Compounds     | Biochemical Nature | Cyanobacteria         | Activity against  | References           |
|-----|---------------|--------------------|-----------------------|---|----------------------|
| 1.  | Noscomin      | Diterpenoid        | <i>Nostoc commune</i> | <i>E. coli</i><br><i>B. cereus</i>  | Jaki et al. (2000)   |
| 2.  | Hapalindole-T | Diterpenoid        | <i>Fischerella</i> sp | <i>M. tuberculosis</i> ,<br><i>E. aerogenes</i> , <i>S. aureus</i> , <i>P. aeruginosa</i> ,<br><i>S. typhi</i> , <i>E. coli</i> | Asthana et al.(2006) |

|    |                   |               |                            |   |                        |
|----|-------------------|---------------|----------------------------|---|------------------------|
| 3. | Cryptophycin      | Desipeptide   | <i>Nostoc</i> ATCC 53789   | <i>Armillaria</i> sp. <i>S. sclerotiorum</i>  | Schwartz et al. (1990) |
| 4. | Scytoscalarol     | Sesterterpene | <i>Scytonema</i> sp.       | <i>B. anthracis</i> ,<br><i>S. aureus</i> , <i>E.coli</i> ,<br><i>C. albicans</i><br><i>M. tuberculosis</i> | Mo (2009)              |
| 5. | Lyngbyazothrins   | peptides      | <i>Lyngbya</i> sp          | <i>B. subtilis</i> , <i>E. coli</i> , <i>P. aeruginosa</i> , and <i>S. marcescens</i>                       | Asthana et al. (2009)  |
| 6. | Anachelin H       | Peptides      | <i>Anabaena cylindrica</i> | <i>M. catarrhalis</i>   | Gademann (2007)        |
| 7. | Anabaenopeptins A | Peptides      | <i>Anabaena flos-aquae</i> | <i>M. catarrhalis</i>   | Harada et al. (2004).  |

Cryptophycin was produced by the cyanobacterium *Nostoc* strain ATCC 53789, and its methanolic extract exhibited antifungal (*Armillaria* sp. *Sclerotinia sclerotiorum*), nematocidal (*Caenorhabditis elegans*), and cytotoxic (*Artemia salina*) activity, but no antibacterial activity (Biondi et al. 2008). Therefore, this cyanobacterium is utilized as a source of pesticides in the agriculture sector and is beneficial for the overcome its chemical toxicity in soil, plants, and non-target organisms. (Fan et al. 2013) noticed that photosynthetic pigments phycocyanin was extracted from *Anabaena cylindrica*, and it exhibits antibacterial activities on *Bibrio parahemolyticus*, *Bacillus mucilaginosus* and *Sarcina lutea*. Bioassay-guided fractionation was used to extract scytoscalarol, an antimicrobial sesterterpene with a guanidino group isolated from cyanobacterium *Scytonema* sp. (UTEX 1163). *Bacillus anthracis*, *Staphylococcus aureus*, *Escherichia coli*, *Candida albicans*, and *Mycobacterium tuberculosis* were all sensitive to scytoscalarol (Mo et al. 2009). Lyngbyazothrins A, B, C, and D are four novel cyclic undecapeptides isolated from cultivated *Lyngbya* sp. *Lyngbyazothrins* A and B had only modest action against *Micrococcus flavus*, but lyngbyazothrins C and D were effective against *Bacillus subtilis*, *Escherichia coli*, *Pseudomonas aeruginosa*, and *S. marcescens*.

The bioactive molecule (4-[(5-carboxy-2-hydroxy)-benzyl]-8, 10-dihydroxy-2, 5, 6, 11, 11-pentamethyloctahydrocyclopenta<a>naphthalene) from the Antarctic cyanobacterium *Nostoc* CCC537 was significantly antibacterial against disease-causing bacteria, including *Mycobacterium tuberculosis* H37Rv and *Salmonella typhi* MTCC 3216, respectively (Asthana et al. 2009).

*Microcoleus lacustris* produced two abietane diterpenes that were effective against *Staphylococcus aureus*, *Staphylococcus epidermidis*, and *Vibrio cholerae*, but not other bacteria (Pérez Gutiérrez et al. 2008). The anachelin H produced by *Anabaena cylindrica* is antimicrobial. It has a moderate effect on *Moraxella catarrhalis* and is effective against bacterial and fungal diseases (Gademann et al. 2007). The thermotolerant *Phormidium* strain generated extracellular antimicrobial substances that inhibited the development of Gram-positive and Gram-negative bacteria and fungus in batch cultures (Fish and Codd 1994). The toxic effects of cyanobacterial cyclopeptides, microcystins, and nodularins pose a danger to humans.

Microcystin is a cyclic heptapeptide that has the potential to cause cellular damage when ingested into cells through various transporters such as Anion Transporting Polypeptide system (OATP). Certain OATPs are overexpressed in malignant cells relative to normal tissues, suggesting that microcystin may be a promising target for the development of cancer treatments (Sainis et al. 2010).

## 1.2 Antiviral Compounds

At the global level, spreading viral diseases such as HIV/AIDS, avian influenza, coronavirus, etc., exhibited fatal consequences for human society. The list of antiviral compound synthesized by cyanobacteria are listed in Table 2. Highly antiretroviral therapy (HAART) is used to control HIV and is approved by Food and Drug Administration (FDA). Therefore, novel drugs are essential requirements for fighting against fatal diseases.

Cyanobacterial origin of antiviral compounds exhibits activity by preventing absorption or penetration and viral replication after entry into cells (Yang et al. 1999) isolated peptides cyanovirin-N (Cv-N) from cyanobacteria, which inhibits HIV to cell fusion. Cv-N (101-amino acid peptide sequence) is an 11kDa protein derived from *Nostoc ellipsosporum*; it binds to gp120 and gp41 and suppresses many HIV and SIV strains (SiV). Calcium spirulan (Ca-Sp) is a sulfated polysaccharide antiviral agent derived from the cyanobacterium *Spirulina platensis*'s hot H<sub>2</sub>O. These antiviral medications impeded the reproduction of enveloped viruses, including Herpes simplex virus type 1, measles virus, influenza A virus, mumps virus, human cytomegalovirus, and HIV-1 (Hayashi et al. 1996). Antiviral medicines like spirulan were discovered from polysaccharide fractions of *Arthrospira platensis* cyanobacteria. These medicines displayed potent antiviral action against cytomegalovirus, human herpes virus type 6, and HIV-1, although their effect on Epstein-Barr virus and influenza A virus was negligible (Rechter et al. 2006). A polysaccharides antiviral drug 211kDa Nostoflan (Kanekiyo et al. 2005) was isolated from cyanobacterium *Nostoc flagelliforme* and active against the enveloped virus, such as influenza A and HSV, HCMV virus. These antiviral drugs inhibit the adhering of the virus to the cells (Kanekiyo et al. 2007).

In the form of a dietary supplement, *Spirulina platensis* commonly used, and exhibits antiviral and immune-stimulating properties. The natural extract of *S. platensis* was a potent inhibitor against the enveloped virus by preventing some stages of replications of the virus after penetrations into the cell (Yakoot and Salem 2012). Extracts of polysaccharides from *Gloeocapsa turgidus* and *Synechococcus cedrorum* had more antiviral activity against rabies virus than the herpes-1 virus, but extracts of polysaccharides from *Amorphonostoc punctiforme* exhibited either weak or no antiviral activity against both viruses.

**Table 2.** Antiviral compounds produced by cyanobacteria

| S.No. | Compounds        | Biochemical Nature        | Cyanobacteria                   | Activity                  | References              |
|-------|------------------|---------------------------|---------------------------------|---------------------------|-------------------------|
| 1     | Nostoflan        | Acidic polysaccharide     | <i>Nostoc flagelliforme</i>     | Anti-herpes simplex virus | Kanekiyo et al.,(2005)  |
| 2.    | Ichthyopeptins A | Cyclic depsipeptides      | <i>Microcystis ichthyoblabe</i> | Influenza A virus         | Zainuddin et al. (2007) |
| 3.    | Calcium spirulan | Sulfated polysacchride    | <i>Spirulina platensis</i>      | Anti-HIV                  | Hayashi et al. (1996)   |
| 4.    | Cyanovirin       | Cyanobacterial protein    | <i>Scytonema varium</i>         | Anti-HIV                  | Dey et al. (2000)       |
| 5.    | Scytovirin       | Cyanobacterial protein    | <i>Scytonema varium</i>         | Anti-HIV                  | Xiong et al. (2006)     |
| 6.    | Sulfoglycolipid  | Acylated sulfoglycolipids | <i>Lyngbya lagerheimii</i>      | Anti-HIV                  | Loya et al. (1998)      |

Results concluded that cyanobacterial-derived polysaccharides are effective against animal as well as human viruses (Zheng et al. 2006; Mansour et al. 2011). It was found that exopolysaccharides of halophilic cyanobacterium *Aphanothece halophytica* showed effective antiviral drugs against influenza virus A (H1N1) and 30% reduction of *Diplococcus pneumoniae* infection by oral administrations of exopolysaccharides in disease suffering mice (Zainuddin et al. 2002). It was noticed that methanolic extract of genus *Microcystis* including *M. aeruginosa* and *M. ichthyoblabe* showed antiviral activity of influenza A virus. Marine cyanobacteria are natural source of cyclic, linear peptides and depsipeptides and these compounds are protease inhibitors, and exhibits antiviral activity against HIV virus (Musale et al. 2020). Lipid soluble extract of *Spirulina platensis* and *Lyngbya lagerheimii* shows antiviral agents against Herpes simplex virus (HSV) type 1, identification of extract was confirmed as sulphoquinovosyl diacylglycerols, a potent antiviral agent (Chirasuwan et al. 2009). These compounds inhibit reverse transcriptase, DNA polymerase and telomerase (Gustafson et al. 1989; Gupta et al. 2014). It was also reported that antiviral agent polyketides isolated from *Trichodesmium erythraeum* and these compounds were active against the Chikungunya virus.



Thus, innovative medications are urgently required to combat the world's most severe illnesses. Antiviral substances generated from cyanobacteria are generally bioactive by blocking viral absorption or penetration into cells and inhibiting progeny viruses' reproductive stages following infection.

### 1.3 Antiprotozoal Activity

According to the World Health Organization, approximately 1.2 billion populations of the world are infected with protozoan diseases such as malaria, leishmaniasis and trypanosomiasis, especially in tropical countries (Simmons et al. 2008). The list of antiprotozoal compounds synthesized by cyanobacteria are listed in Table 3.

**Table 3.** Antiprotozoal compounds produced by cyanobacteria

| S.No. | Compounds      | Biochemical Nature                           | Cyanobacteria                     | Activity  | References               |
|-------|----------------|--|-----------------------------------|---|--------------------------|
| 1     | Viridamide-A   | Lipodepsipeptide                             | <i>Oscillatoria nigro viridis</i> | <i>P.falciparum</i><br><i>L. mexicana</i>                   | Simmons et al. (2008)    |
| 2.    | Viridamide-B   | Lipodepsipeptide                             | <i>Oscillatoria nigro viridis</i> | <i>P.falciparum</i><br><i>L. mexicana</i><br><i>T.cruzi</i> | Simmons et al., (2008)   |
| 3.    | Symplocamide-A | Homodetic cyclic peptide                     | <i>Oscillatoria</i> sp            | <i>P.falciparum</i>   | Linington et al., (2007) |
| 4.    | Dragomabin     | Lipopeptide                                  | <i>Lyngbya majuscula</i>          | <i>P.falciparum</i>   | Mcphail et al. (2007).   |
| 5.    | Ambigol-A      | Polychlorinated triphenyl aromatic compounds | <i>Fischerella ambigua</i>        | <i>P.falciparum</i>   | Falach et al.(1993)      |
| 6.    | Ambigol-B      | Polychlorinated triphenyl aromatic compounds | <i>Fischerella ambigua</i>        | <i>Trypanosoma rhodesiense</i>                              | Falach et al.,(1993)     |
| 7.    | Ambigol-C      | Polychlorinated triphenyl aromatic compounds | <i>Fischerella ambigua</i>        | <i>Trypanosoma rhodesiense</i>                              | Falach et al.,(1993)     |

Treatment failures, mainly in the resistance to malaria and leishmaniasis, are associated with the development of resistance by the protozoa (Singh et al. 2014; Qureshi et al. 2021b; Qureshi et al. 2022). Contrary, progress in drug discovery initiatives to combat these diseases is quite slow. The Panamanian International Cooperative Biodiversity Group is researching bioactive chemicals in cyanobacteria to promote the development of effective and cost-efficient therapies for various ailments. Five kinds of antiprotozoal compounds from cyanobacteria were reported by (Singh et al. 2014). Terrestrial cyanobacteria *Nostoc commune* and *Rivularia biolettiana* showed antiprotozoal activity against *Plasmodium falciparum*, *Trypanosoma brucei rhodesiense*, and *Leishmania donovani*. Aerucyclamide C and D were isolated and characterized from *Microcystis aeruginosa* PCC 7806 and exhibited antiplasmodial activity. Aerucyclamide B and C were effective against *P. falciparum* and *T. brucei rhodesiense*, respectively. It was found that six new bioactive acyl proline derivatives, tumonoic acids D-I, derived from the cyanobacterium *Blennothrix cantharidosmum* had weak antimalarial properties (Clark et al. 2008). The lipodepsipeptides Viridamides A and B, isolated from the marine cyanobacterium *Oscillatoria nigroviridis* were antiprotozoal.

### 1.4 Anticancer Compounds

Cyanobacteria synthesized a large number of bioactive compounds which targeted the actin/tubulin filaments in eukaryotes (Jordan and Wilson 1998). The list of antiviral compounds synthesized by cyanobacteria are listed in Table 4.

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**Table 4.** Anticancerous compounds produced by cyanobacteria

| S. No. | Compounds      | Biochemical Nature  | Cyanobacteria                        | Target Site   | References                                     |
|--------|----------------|---------------------|--------------------------------------|---|--|
| 1      | Dolastatin-10  | Linear pentapeptide | <i>Leptolyngbya</i> sp               | Inhibition of microtubule Assembly  | Kalemkerian et al. (1999)                      |
| 2      | Dolastatin-12  | Linear pentapeptide | <i>Symploca</i> sp                   | Induced cell cycle arrest at the G2-M phase                                     | Catassi et al. (2006)                          |
| 3      | Curacin-A      | Polyketide          | <i>Lyngbya majuscula</i>             | Inhibition of polymerization of tubulin in eukaryotes                           | Gerwick et al. (1994). Yoo and Gerwick, (1995) |
| 4      | Curacin-B      | Polyketide          | <i>Lyngbya majuscula</i>             | Inhibitors of microtubule polymerization  | Gerwick et al. (1994). Yoo and Gerwick, (1995) |
| 5      | Curacin-C      | Polyketide          | <i>Lyngbya majuscula</i>             | Inhibitors of microtubule polymerization  | Gerwick et al. (1994). Yoo and Gerwick, (1995) |
| 6      | Curacin-D      | Polyketide          | <i>Lyngbya majuscula</i>             | Colchicines binding and tubulin polymerization inhibitors                       | Márquez et al. (1998)                          |
| 7      | Veraguamides-A | Hexadepsipeptides   | <i>Symploca</i> cf. <i>hydroides</i> | Cytotoxic activity against colon Cervical carcinoma                             | Salvador et al. (2011).                        |
| 8      | Obyanamide     | Cyclodepsipeptide   | <i>Lyngbya confervoides</i>          | Antineoplastic activity   | Williams et al. (2002).                        |
| 9      | Malevamide-D   | Peptide ester       | <i>Symploca hydroides</i>            | Antineoplastic activity   | Horgen et al. (2002)                           |
| 10     | Majusculamide  | Amino acid amide    | <i>Lyngbya majuscula</i>             | Antineoplastic activity   | Pettit et al. (2008)                           |
| 11     | Lagunamide-A   | Cyclodepsipeptide   | <i>Lyngbya majuscula</i>             | Cytotoxic activity against Murine Leukemia cell lines and antimalarial activity | Tripathi et al. (2010)                         |
| 12     | Lagunamide-B   | Cyclodepsipep       | <i>Lyngbya</i>                       | Cytotoxic and   | Tripathi et                                    |

|    |                            |                          |   |  |                        |
|----|----------------------------|--------------------------|---|--|------------------------|
|    |                            | tide                     | <i>majuscula</i>                            | antimalarial properties                            | al (2010)              |
| 13 | Lagunamide-C               | Cyclodepsipeptide        | <i>Lyngbya majuscula</i>                    | Cytotoxic, antimalarial and anti-swarming activity | Tripathi et al (2011)  |
| 14 | Lagunamide-D               | Macrocyclic Depsipeptide | <i>Lyngbya</i> sp and <i>Dichothrix</i> sp. | Induced apoptosis                                  | Luo et al., (2019)     |
| 15 | Digalactosyldiacylglycerol | Glycerides               | <i>Phormidium tenue</i>                     | Antitumorogenic                                    | Tokuda et al. (1996)   |
| 16 | Caylobolide-B              | Macrolactone             | <i>Phormidium</i> sp.                       | Cytotoxic  | Salvador et al. (2010) |
| 17 | Bisebromoamide             | Peptide                  | <i>Lyngbya</i> sp                           | Protein kinase Inhibitors                          | Teruya et al. (2009)   |
| 18 | Largazole                  | Cyclic depsipeptide      | <i>Symploca</i> sp.                         | Anticancer properties and osteogenic activities    | Taori et al. (2008)    |
| 19 | Hormothamnin-A             | Cyclic undecapeptide     | <i>Hormothamnion enteromorphoides</i>       | Cytotoxicity                                       | Malloy et al. (2012)   |

Two anticancer compounds Dolastatin 10 and 12, were isolated from *Leptolyngbya* sp. and *Symploca* sp., respectively, and these compounds are linear pentapeptides (Kalemkerian et al. 1999; Catassi et al. 2006). *Lyngbya majuscula* is a rich source of bioactive chemicals from which Curacin A, B, C, and D are manufactured. They possess powerful antiproliferative and antimetabolic properties (Nagle et al. 1995). Lyngbyabellin B was extracted from a different species of *L. majuscula* and showed cytotoxic activity against LoVo and KB cell lines. It was reported that two anticancer compounds, Malynamide 3 and Cocosamides B, are anticancer from *L. majuscula*, and these compounds exhibited poor cytotoxicity against colon and breast cancer (Linnington et al. 2008). Cryptophycin inhibited leukemia cell line (L-1210) in a dose-dependent manner and isolated from *Nostoc* sp. (Smith et al. 1994). *Calothrix* sp. is the source of Calothrixin A and B and was extracted from the organic extract. It is a potent inhibitor of human cancer cells of the HeLa cell line (Rickards et al. 1999). Nine variants of Apratoxin, such as A, B, C, D, F, D, E, F and G, were isolated and characterized from *Lyngbya* spp. These variants exhibited potent cytotoxicity against different cancer cell lines of HT-29 colon adenocarcinoma, H-460 lung cancer, HCT-116 colorectal, HeLa cervical carcinoma and KB oral epidermoid (Luesch et al. 2001; Matthew et al. 2008; Pérez Gutiérrez et al. 2008; Tidgewell et al. 2010; Oftedal et al. 2010) evaluated 41 marine cyanobacteria, and are effective against acute myeloid leukemia cell lines. Approximately 50% exhibited apoptosis-inducing activity in the aqueous extract of cyanobacteria. A mixture of cyanobacterial extract acts as a synergistic effect with daunorubicin (anthracycline anticancer drug anticancer) in AML cell lines, and cardiomyoblasts were inhibited. Therefore, marine cyanobacteria are a novel source of antileukemic drugs (Oftedal et al. 2010). Largazole (I) is a macrocyclic depsipeptide and isolated *Symploca* sp. The natural products of Largazole showed specific bio-activity with differential growth inhibition (2-10%) of mice cell lines and transformed, non-transformed human cell lines (Bowers et al. 2008).

Veraguamides, cyclic hexadepsipeptides generated from the ocean cyanobacterium *Symploca cf. hydroides* and collected from the Micronesian island, were cytotoxic against colon adenocarcinoma HT29 and cervical cancer

cell lines HeLa (Salvador et al. 2011). Freshwater cyanobacterial extracts of *Carotinosum* and *N. linckia* showed antiproliferative and cytotoxic effects on the rat Glioma C6 cell line (Karan and Aydin 2018).

Similarly, *A. oryzae* extract suppressed the growth of MCF7 breast cancer cells (Karan and Aydin 2018). Apratoxin A, a cyclic depsipeptide produced from *Lyngbya* spp., has a fatal impact on human cervical carcinoma HeLa cell lines by interfering with the cell cycle (Malve 2016). Peptides were isolated from cyanobacteria *Nostoc* sp. and *Lyngbya* sp. exhibited anticancer activity by inhibiting dissociation of microfilaments and secretory pathways (Malve 2016). *Synechocystis* sp. and *Synechococcus* sp. aqueous extracts showed anticancer activities against HL-60 human leukemia cell lines by activating apoptosis and membrane budding (Martins et al. 2008).

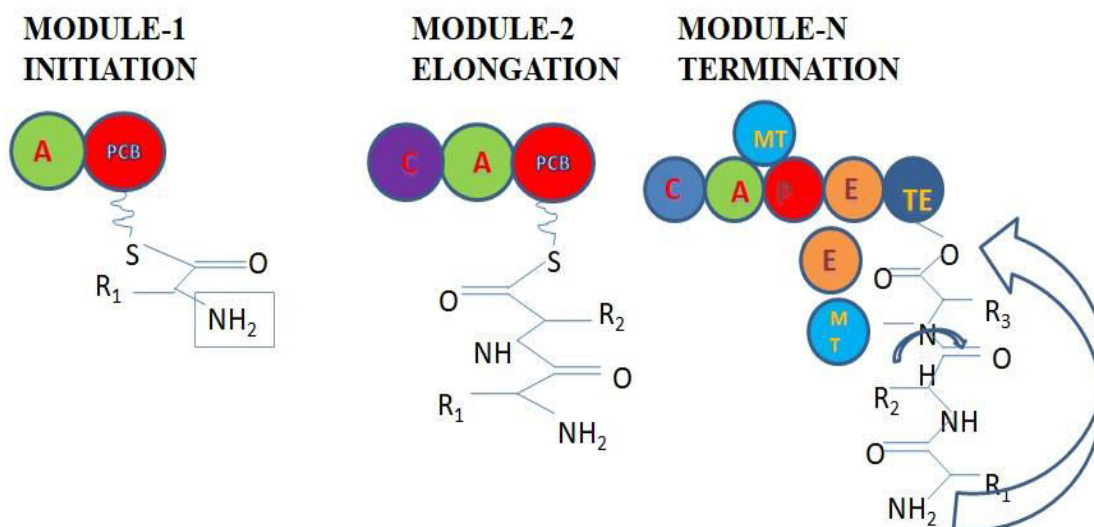
Several reports have been published that photosynthetic pigments also act as anticancer activity. C-phycoyanin (CP) is a photosynthetic pigment that shows anti-tumor activity by suppressing the growth of tumor in a concentration and time-dependent manner. CP exhibited potent anticancer properties in several cancer cell lines, including breast cancer (Li et al. 2010), colon cancer (Wang et al. 2007), and bone marrow cancer (Gardeva et al. 2014); multiple processes are involved, including the production of reactive oxygen species (ROS), programmed cell death, and cell cycle inhibition (Pan et al. 2015; Hao et al. 2021).

The extract of *Arthrospira platensis* containing phycoyanin act as an anti-cancer drug for various tumor cells (Jiang et al. 2017; Braune et al. 2021) found that phycoyanin is a promising source of anticancer agents due to the inhibition of cell-cycle arrest, stimulation of cell apoptosis, autophagy and reduction of ant-proliferating activity. Human prostate cell line LNCaP cured by combined application of C-phycoyanin and 10% topotecan as compared with C-PC treatment and single topotecan. It activates a series of caspases by producing radical oxygen (ROS) that leads to the apoptosis of tumor cells and inhibits toxicity generated by topotecan (Gantar et al. 2012).

### **1.5 Biosynthesis of bioactive compounds**

Cyanobacteria synthesized different groups of natural products and showed different chemical natures. Alkaloids, polyketides, terpenes, fatty acids, and UV-absorbing chemicals are bioactive compounds. Peptide biosynthesis operates into ribosome-independent pathways and is operated by enzymes known as non-ribosomal peptide synthetase (NRPS), and further peptides are modified during post-translational modification and processing. NRPs are a kind of secondary metabolite found in various microorganisms, including bacteria, cyanobacteria, fungi, and plants (Singh et al. 2019). Non-ribosomal peptides synthetases are multimodular proteins, and it synthesized by known as NRPs genes. Single module domain proteins are also reported, and approximately 3 to 3.5 Kb genetic sequences are required. They are multifunctional multimodular enzymes; that consist of non-ribosomal peptide synthetases (NPSs) and polyketide synthases (PKS), which are responsible for the biosynthesis of non-ribosomal peptides and polyketides, respectively (Ref. figure 1 & 2).





**Figure.1.** Nonribosomal peptides synthetases (NPSs) consist of different domains; adenylation domain (A), peptidyl carrier domain (PCB), condensation domain (C) thioesterase domain (TE), epimerization domain (E), and methyltransferase (MT) domain. These enzymes catalyzed synthesis of nonribosomal peptides.

Diversity of these bioactive compounds (peptides and polyketides) is maintained by NRPSs (Süssmuth and Mainz 2017). NRPs activated and incorporate amino acids (D and L), hydroxy acids, keto acid, heterocyclic and other functional groups. They contribute structural and chemical diversity of the peptides (McErlean et al. 2019; Kudo et al. 2019). Consequently, NRPSs are extraordinary macromolecular machines that generate various physiologically and therapeutically significant compounds. (Bloudoff and Schmeing 2017).

The several modules of NRPS are responsible for the progressive incorporation of amino acids. The majority of modules are composed of three domains: the adenylation domain (A) (amino acid activation domain), the peptidyl carrier domain (PCB), and the condensation domain (C) (Fig.1) (Koglin and Walsh 2009; Marahiel and Essen 2009). NRPs may add up to 300 proteinogenic and non-proteinogenic substrates, in addition to domain-specific tailoring modifications and amino acid epimerization as substrates (Miller and Gulick 2016). These groups of peptide biosynthetic processes displayed diverse natures and considerable activity. Ribosome-mediated peptides can add only 20 amino acids, and leader and core peptides are an integral part of proteins.

Amino acids are selected and activated by the adenylation domain (Finking and Marahiel 2004; Marahiel and Essen 2009). The peptide carrier protein and condensation domains entail the formation of peptide bonds between two amino acids and the binding to peptidyl carrier proteins. The adenylation and peptidyl carrier protein domains are components of an initiation module. The thioesterase domain catalyzes the production of a macrocycle and the release of molecules (TE) (Finking and Marahiel 2004; Marahiel and Essen 2009). The presence of tailoring domains might increase the number of peptide modifications possible. A few of the tailoring domains are the epimerization domain (E), which converts L- to D-amino acids; the methyltransferase domain (MT), which methylates amino acid residues; and the monooxygenase domains (OX), which adds one hydroxyl group to substrates. (Finking and Marahiel 2004; Marahiel and Essen 2009). Glycosyltransferases are the enzymes responsible for adding sugar units to free hydroxyl groups on polypeptides. (Finking and Marahiel 2004; Welker and Von Döhren 2006)

Similarly, PKS-type I modules must have at least the following domains: acyltransferase (AT), acyl carrier protein (ACP), and ketosynthase (KS). Acyltransferase picked one unit and incorporated it into the peptide malonyl-, methyl-, ethyl-, and propylmalonyl-CoA (Katz 1997). The malonyl-CoA unit is transferred to the acyl carrier protein, and the ketosynthase domain enables the development of a Claisen-condensation link between the

two units. During the last stages of elongation, the thioesterase domain (TE) cleaves and macrocyclizes the product in every PKS domain. It was reported that NRPS and PKS domains are present in the hybrid system and are involved in the secondary metabolites production of cyanobacteria (Wang et al. 2014). A fatty acid part can be found in the chemical structure of a few non-ribosomal peptides and polyketides. Fatty acid synthesis can take two forms; type I, which involves large multifunctional proteins, and type II, which involves working specific proteins (Campbell and Cronan 2001). Type II is more prevalent in bacterial systems and has more conserved or less conserved protein sequences, such as enoyl reductases, Fab fatty acid biosynthesis, acyl-CoA carboxylase, or (ER), respectively. (Campbell and Cronan 2001). In addition, other proteins, including acyl carrier protein (ACP) and biotin carboxyl carrier protein (BCCB), are involved in fatty acid production (Campbell and Cronan 2001).

Fatty acyl-CoA ligases (FACLs) might produce fatty acetaldehyde, an intermediate unit for the derivative of coenzyme A (Trivedi et al. 2004; Arora et al. 2009). The production of lipid molecules is promoted by activated fatty acyladenylates (Arora et al. 2009)

Curacin A is a secondary metabolite formed by the addition of fatty acids by PKSs or the condensation domain of NRPS, which acts as an acceptor of a  $\beta$ -hydroxyl fatty acid (Rausch et al. 2007). Surfactin (Arima et al. 1968; Rausch et al. 2007), lichenysin (Horowitz and Griffin 1991), fengycin (Tosato et al. 1997), arthrofactin (Morikawa et al. 1993), polymyxin (Choi et al. 2009), and pelgipeptin (Horowitz and Griffin 1991) are examples of biosynthetic gene clusters.

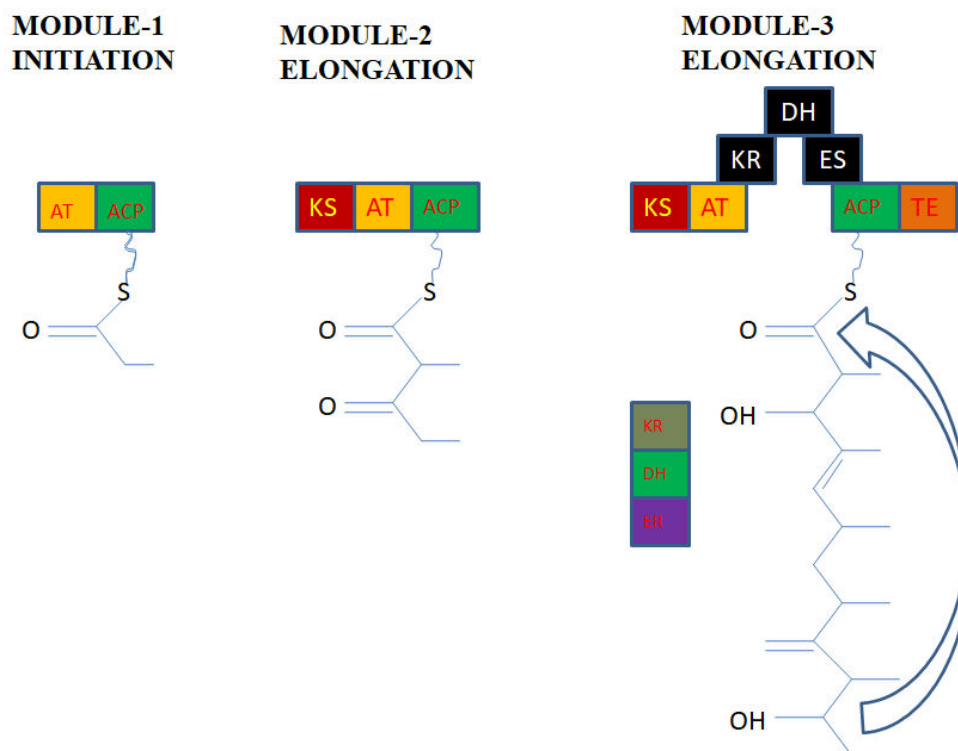
### **1.6 Methods for Isolation and Characterization of Bioactive Compounds**

A variety of approaches are used to extract valuable secondary metabolites. Extracellular and intracellular metabolites can be extracted simultaneously (extracellular and intracellular metabolites) or sequentially (intercellular metabolites). Both extraction methods start with liquid nitrogen, quenching to freeze the metabolic activity. Organic solvent (butanol, acetone, hexane, chloroform) and a mixture of water and organic solvent in different ratios is generally used for immediate quenching in the simultaneous extractions. Classical extraction methods generally used solid-liquid (SLE) and liquid-liquid (LLE). SLE method Soxhlet is often used, and solutes are extracted from solids or matrices in a liquid solvent.

In contrast, in LLE, both phases are liquid, and the separations of the solutes are based on the distribution coefficient of the solute in the liquid solvent. All these extraction methods, SLE and LLE, need large amounts of solvent and time. The drawback of the methods is their poor reproducibility.

Some other low-cost methods, such as extraction, supercritical fluid extraction (SFE), and pressurized liquid extraction (PLE), are also applied. Solute are extracted in SCF by employing gases above the critical temperature. Carbon dioxide is most popularly used as a supercritical fluid, and sometimes methanol or ethanol is also used as a co-solvent. Some researchers extracted antibacterial components from *S. platensis* using carbon dioxide and 10% ethanol (co-solvent) (Mendiola et al. 2007). Using a combination of pressurized-liquid and solid-phase extraction (PLE-SPE), Onofrejova et al. extracted bioactive phenols from freshwater and marine algae (Onofrejová et al. 2010).

Several analytical methods are available for detecting and purifying bioactive molecules from the extract of cyanobacterial (Fig. 2).



**Figure.2.** Multifunctional and multimodular polyketides synthases (PKS), consist of different domains (acyltransferase (AT), acyl carrier protein (ACP), ketosynthase (KS), thioesterase domain (TE). These enzymes are involved in biosynthesis of polyketides.

The simplest approach for identifying and separating bioactive compounds is thin layer chromatography (TLC), followed by spectrophotometric analysis. Pelander et al. separated short cyanobacterial peptides utilizing high-performance TLC plates (HPTLC) (Pelander et al. 2000). Conversely, TLC/HTLC separation is nonspecific and poorly sensitive. The application of high-performance liquid chromatography (HPLC) for detection and characterization has grown considerably. Ultra-performance liquid chromatography (UPLC), a more modern method, is now available, and it might be a better alternative than HPLC. Liquid chromatography is followed by mass spectrometry (LC-MS) to accurately identify a bioactive compound (Harada et al. 2004). The different configurational approaches utilized are atom bombardment (FAB-LC-MS) and electrospray ionization (ESI-LC-MS). Zhang et al. used electrospray ionisation with LC-MS-MS to identify cyanobacterial cyclopeptides (Zhang et al. 2021). Liquid-liquid extraction can be coupled with quadruple time-of-flight tandem mass spectrometry (LC/Q-TOF-MS).

Immunoassay methods are widely used to characterize biomolecules due to their better sensitivity, specificity, and easy operation. Lindner et al. reported that enzyme-linked immunosorbent assays are highly sensitive methods for characterizing cyanobacterial cyclopeptides (Lindner et al. 2004). Matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (MALDI-TOF MS) is also used to characterize the metabolites. This approach uses a minimal sample size and does not need sample separation or measurement (Welker et al. 2002).

Identification and characterization of intact cyanobacterial cells using MALDI-TOF-MS were reported by (Welker et al. 2002). MALDI-TOF MS is frequently used to identify peptides but may also be used to identify alkaloids. Desorption electrospray ionization mass spectrometry (DESI-MS) is also utilized for chemical profiling, characterization, and quantification of lower molecular weight biomolecules (Bayona et al. 2022). Direct analysis in real-time mass spectrometry (DART-MS) is another approach that may be used for the chemical profiling and characterization of bioactive chemicals. Tripathi et al. used DART-MS to identify the

*Nostoc* sp.-based chemical components (chemical fingerprinting) (Tripathi et al. 2021). These approaches are useful in identifying and characterizing bioactive molecules.

## CONCLUSION

Cyanobacteria are a unique group of photosynthetic autotrophic bacteria, primary producers and distributed worldwide. They provide food to many micro and macro-organisms, and their food habitats differ from heterotrophic bacteria. During the last two decades, they have been recognized as an emerging source of therapeutic bioactive compounds. In contrast to protein biosynthesis, NRPs are large multimodular enzymes with multiple subunits. Secondary metabolites produced by cyanobacteria include alkaloids, polyketides, and non-ribosomal peptides. Non-ribosomal peptides are secondary metabolites synthesized by prokaryotes, especially cyanobacteria and lower eukaryotes; they are diverse in nature, such as pigments, toxins, siderophores and others. Thus, the studies of cyanobacterial-based bioactive compounds are promising endeavour. We have explained several bioactive compounds derived from cyanobacteria and have a broad spectrum of activity, such as antibacterial, antiviral, anticancer and antiprotozoal. Due to simple microbes, cyanobacteria are an excellent option for drug discovery because it requires inorganic nutrient media and is easy to cultivate. Consequently, we advise expanding cyanobacterial drug discovery efforts and enhancing biotechnological and bioprocess engineering for chemical production.

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