

Taxonomic Distribution of Bioactive Metabolites in *Nostoc* and *Nostoc*-like Genera

Maria Christodoulou*

Department of Microbiology, University of Helsinki, Helsinki, Finland

Introduction

Cyanobacteria address a gathering of morphologically different oxyphototrophic microscopic organisms found in practically all environments on Earth including those which are viewed as threatening to life and are known as 'outrageous' conditions. The enormous variety of cyanobacteria alludes not exclusively to their morphological changeability and living space inclinations yet additionally to an assortment of practically different and primarily complex metabolites they produce. A consequence of this serious level of synthetic variety is the expanded chance of finding substances with biomedically fascinating properties with regards to cyanobacteria [1].

Description

As per similar examinations, the significant justification for this peculiarity is the grouping of cyanobacteria dependent exclusively upon morphological symptomatic highlights, consequently bringing about a huge misstatement of cyanobacterial variety. This morphology-just based approach, which was customarily utilized in cyanobacterial scientific categorization, is these days supplanted by a cutting edge framework in which new taxa are depicted in view of a mix of morphological, ultrastructural, sub-atomic and environmental information. This 'polyphasic approach' drove not just to the foundation of a few new monophyletic taxa yet additionally considered more designated regular item revelation endeavors [2].

Nostoc, with *Nostoc* cooperative as its sort species, is one of the most notable, cosmopolitan cyanobacterial genera. In spite of being quite possibly of the earliest cyanobacterial family portrayed, its scientific classification stays tricky because of the absence of clear morphological attributes. Ongoing phylogenetic investigations have shown that *Nostoc* is polyphyletic and a few new *Nostoc*-like genera including *Mojavia*, *Desmonostoc*, *Halotia*, *Komarekiella*, *Aliinostoc*, *Desikacharya*, *Compactonostoc*, *Parakomarekiella* and *Pseudotaliinostoc* have proactively been laid out [3].

Notwithstanding its hereditary heterogeneity, strains relegated to *Nostoc* have been accounted for to create various normal items going from poisons to organically dynamic particles, proposing that *Nostoc* is a productive wellspring of bioactive mixtures. *Nostoc*-like strains have been separated from a large number of natural surroundings including basic lakes, which address a novel and stable climate portrayed by high pH levels that have remarkable microbial networks.

Heterocytous cyanobacteria detached from Brazilian basic lakes including Salina Verde were concentrated utilizing a polyphasic approach [4]. The

morphological and phylogenetic examination led in that uncovered, bury alia, the presence of a clever cyanobacterial heredity that was morphologically comparative yet phylogenetically far off from *Nostoc sensu stricto* clade. The bunch was described by the presence of motile hormogonia with gas vesicles that was considered an autapomorphic diacritical include isolating this bunch from other heterocytous genera. *Nostoc*-like variety *Aliinostoc* from a eutrophic lake in India and proposed the arrangement of the previously mentioned strains under this book family. From that point forward, two more *Aliinostoc* species secluded from soil territories in Iran were laid out utilizing a polyphasic approach. These three taxa are the just truly depicted *Aliinostoc* species and not a single one of them have been considered in regard to their bioactive potential [5].

Conclusion

The point of this study is to grow the biological and geological dispersion of *Aliinostoc* in view of a polyphasic approach. Besides, this study investigates the bioactive capability of the new species against bacterial and parasitic potential microorganisms as a feature of our continuous endeavors to find novel bioactive metabolites with drug applications. At long last, this study uses generally accessible hereditary and bioactive metabolite data to explore whether family *Nostoc* is without a doubt wealthy in bioactive mixtures or whether this apparent compound extravagance of *Nostoc* rises out of the absence of appropriate ordered examinations in view of the previously mentioned polyphasic approach.

References

1. Swain, Shasank S., Sudhir K. Paidesetty and Rabindra N. Padhy. "Antibacterial, antifungal and antimycobacterial compounds from cyanobacteria." *Biomed Pharmacother* 90 (2017): 760-776.
2. Mazard, Sophie, Anahit Penesyan, Martin Ostrowski and Ian T. Paulsen, et al. "Tiny microbes with a big impact: The role of cyanobacteria and their metabolites in shaping our future." *Mar Drugs* 14 (2016): 1-19.
3. Singh, Rahul Kunwar, Shree Prakash Tiwari, Ashwani K. Rai and Tribhuban M. Mohapatra. "Cyanobacteria: An Emerging source for drug discovery." *J Antibiot* 64 (2011): 401-412.
4. Jones, Martin R., Ernani Pinto, Mariana A. Torres and Fabiane Dörr, et al. "CyanoMetDB, a comprehensive public database of secondary metabolites from cyanobacteria" *Water Res* 196 (2021): 1-12.
5. Leão, Pedro N., Niclas Engene, Agostinho Antunes and William H. Gerwick, et al. "The chemical ecology of cyanobacteria." *Nat Prod Rep* 29 (2012): 372-391.

*Address for Correspondence: Maria Christodoulou, Department of Microbiology, University of Helsinki, Helsinki, Finland, Tel: 9273349874; E-mail: MariaChristodoulou892@gmail.com

Copyright: © 2022 Christodoulou M. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Date of Submission: 02 June, 2022, Manuscript No: jpeb-22-71727; **Editor assigned:** 04 June, 2022, PreQC No: P-71727; **Reviewed:** 09 June, 2022, QC No: Q-71727; **Revised:** 14 June, 2022, Manuscript No: R-71727; **Published:** 19 June, 2022, DOI: 10.37421/2329-9002.2022.10.221

How to cite this article: Christodoulou, Maria. "Taxonomic Distribution of Bioactive Metabolites in *Nostoc* and *Nostoc*-like Genera." *J Phylogenetics Evol Biol* 10 (2022): 221.