

Polyphasic characterization of four species of *Pseudanabaena* (Oscillatoriales, Cyanobacteria) from China and insights into polyphyletic divergence within the *Pseudanabaena* genus

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Abstract

Current cyanobacterial taxonomic studies are mainly performed through polyphasic characterization at the genus and/or species levels. In this study, eleven *Pseudanabaena* strains isolated in China were taxonomically and phylogenetically evaluated by investigating several characteristics, including morphological features, cellular ultrastructures, pigment composition, and 16S rRNA gene sequences. The strains were identified as four morphospecies, namely, *Pseudanabaena mucicola*, *P. galeata*, *P. limnetica*, and *P. minima*. The cellular ultrastructures of these species showed that peripheral thylakoids with 4–8 parallel layers were parietally distributed. Phylogenetic results based on 16S rRNA gene sequences indicated that most of the *Pseudanabaena* strains used in this study, including the eleven from China and three *Pseudanabaena catenata* strains from GenBank, gathered into one large cluster that represents the core of the genus *Pseudanabaena*. Several *Pseudanabaena* species/strains that represent different extreme environments were phylogenetically located outside the main stream cluster and should be removed under the genus *Pseudanabaena* on the basis of the role of the main stream species. *Limnothrix* strains were divided into two clades, and the taxonomic solution for these clades was discussed. This study is the first to report on the morphology and phylogeny of *Pseudanabaena* species in China, providing new insights into the taxonomy of the genera *Pseudanabaena* and *Limnothrix*.

Key words: China, cyanobacteria, *Limnothrix*, morphology, phylogeny, *Pseudanabaena*, taxonomy

Introduction

Comprehensive studies on water-bloom-forming cyanobacteria have been magnified because of the harmful effects of toxin production and the unpleasant taste and odor caused by these cyanobacteria (Peterson *et al.* 1995, DeVries *et al.* 2012). It is well known for those bloom-forming cyanobacteria, which have relatively large colony/filament sizes and can float on water surface through their gas vesicles. However, tiny cyanobacteria species, including those related to water blooms, have been overlooked because of difficulties in observation brought about by their relatively small cell/filament sizes. Occasionally, these cyanobacteria dominate phytoplankton biomass (Romo and Miracle 1994, Sthapit *et al.* 2008), however, they were usually described with few characters or only recorded as species names probably because of poor understanding of their traits (Mischke and Nixdorf 2003, Villena and Romo 2003). These cyanobacteria are important in ecosystem dynamics and primary productivity.

Anagnostidis and Komárek (1988: 380) described the genus *Pseudanabaena* Lauterborn (1915: 437) as a group of tiny cyanobacteria. With *Pseudanabaena catenata* Lauterborn (1915: 437) (Chang 1988) as the type species, *Pseudanabaena* is characterized by the presence of filaments with conspicuous constrictions at cross-walls consisting of cylindrical cells. Some species that contain both phycocyanin (PC) and phycoerythrin (PE) display complementary chromatic adaptation that allows the regulation of PC/PE ratio to maximize the absorption of available light (Kehoe and Gutu 2006, Stomp *et al.* 2004, 2008).

As described above, the *Limnothrix* strains shown in the phylogenetic tree were mainly divided into two clades (Fig. 4). The first clade included three strains of *Limnothrix redekei* (NIVA-227, CCAP1443/1, and CCAP 1459/29), whereas the second clade included strains of *L. redekei* (2LT25S01 and 165c) and *L. planktonica* CHAB753. The first *Limnothrix* clade was fully embedded in Cluster I with the type species and most *Pseudanabaena* strains. The *L. redekei* strains in this clade should be transferred under *Pseudanabaena* on the basis of the real *Pseudanabaena* cluster described above and later time of establishment of *L. redekei*. The second *Limnothrix* clade was positioned outside the *Pseudanabaena* cluster, suggesting that these *Limnothrix* strains should remain under *Limnothrix*. Such a result partially agreed with the treatment by Whitton (2011), who used *Pseudanabaena redekei* (Goor 1918: 258) B.A. Whitton (2011: 110) as the synonym of *Limnothrix redekei*. Future studies on the taxonomic ambiguity of *Limnothrix* are needed. Investigation of other genes (e.g. *rbcLX*, *rpoCI*, *cpcBA* etc.) might be helpful.

In conclusion, the polyphasic characterization of *Pseudanabaena* strains isolated in China allowed us to establish the main stream species of *Pseudanabaena*. The phylogenetic relationship shown in this study by adding *Pseudanabaena* strains from Chinese waters provided new insights into the taxonomic revision of both *Pseudanabaena* and *Limnothrix*.

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