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**Gênero e espécie novos de cianobactéria homocitada  
(Synechococcales) para a costa brasileira**

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UNIVERSIDADE FEDERAL DA BAHIA  
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**Gênero e espécie novos de cianobactéria homocitada  
(Synechococcales) para a costa brasileira**

por

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## RESUMO

Cyanobacteria é um grupo filogeneticamente antigo de organismos procarióticos fotossintetizantes produtores de oxigênio, tendo surgido entre 2,7 e 3,5 bilhões de anos atrás. A diversidade desse grupo é expressa através de suas propriedades morfológicas, bioquímicas e fisiológicas, possibilitando seu estabelecimento e sobrevivência em uma grande variedade de habitats terrestres e aquáticos. O material utilizado nesse estudo foi oriundo de coleta realizada na Praia de Pirambúzios, Rio Grande do Norte, Brasil (06°00'44.62"S x 35°06'26.48"W), o qual foi mantido em cultivo em meio líquido SWBG-11, com fotoperíodo de 14:10 h (claro:escuro). Para as análises moleculares, a extração do DNA total foi realizada através do método CTAB, seguida de Reações em Cadeia da Polimerase (PCR's) para amplificar a região gênica 16S do RNA ribossomal, utilizando os *primers* 27F1/23S30R. Os produtos da PCR foram clonados usando o kit pGEM®-T Easy Vector Systems. O sequenciamento foi realizado no Centro de Energia Nuclear na Agricultura da Universidade de São Paulo (CENA/USP). Para compor a matriz de sequências para as análises filogenéticas, foi realizada busca através da ferramenta *BLAST* do *National Center of Biotechnology Information* (NCBI), totalizando 64 sequências. Foi utilizado o programa BioEdit v7.2.5 para efetuar o alinhamento, o RaxML8.2.10 para as análises de Máxima Verossimilhança (MV) e o MrBayes on XSEDE para a Inferência Bayesiana (BP). As análises ultraestruturais foram realizadas em microscopia eletrônica de transmissão na FIOCRUZ/BA. A MV e a BP evidenciaram um clado independente com 88% e 100% de suporte, respectivamente, agrupando a sequência obtida neste estudo com uma sequência denominada "Filamentous cyanobacterium" oriunda dos EUA (GenBank GQ243430). As análises ultraestruturais evidenciaram a presença de tilacoides dispostos parietalmente à membrana plasmática, que, aliados à forma filamentosa, inclui esta cepa na família Pseudanabaenaceae (Synechococcales). As características morfológicas, como a forma da célula apical, a morfometria das células, o habitat marinho e o hábito epifítico não foram descritos para a família. Esses resultados sugerem o reconhecimento de um novo gênero, representado por duas espécies, sendo uma delas, ALCB 125750, descrita neste estudo.

Palavras-chave: Cyanobacteria; ambiente marinho; taxonomia

## ABSTRACT

Cyanobacteria is a phylogenetically old group of prokaryotic photosynthetic organisms, which produces oxygen, arising between 2.7 to 3.5 Bya. The diversity of this group is expressed by the morphological, biochemical and physiology properties, making possible its establishment and persistency in a wide range of terrestrial and aquatic habitats. Brazil possess one of the greatest coasts of the world, providing a big diversity of habitats that possibilities the occurrence of benthic cyanobacteria. The material analyzed in this study was collected at Pirambúzios beach (Rio Grande do Norte, Brazil) (06°00'44.62"S x 35°06'26.48"W). It was kept in liquid medium SWBG-11 for cultivation, with photoperiod of 14:10 h (light:dark). To molecular analysis, DNA extraction was carried out with CTAB method, followed by Polymerase Chain Reaction (PCR) to amplify the 16S rRNA genic region, using 27F1/23S30R primers. PCR products were cloned using the pGEM®-T Easy Vector Systems kit. Sequencing was realized at the Nuclear Center in Agriculture of São Paulo's University (CENA/USP). In order to compose the sequence matrix to the phylogenetic analysis, a search was realized through the BLAST tool of the *National Center of Biotechnology Information* (NCBI), totalizing 64 sequences. BioEdit v7.2.5 program was used to align the sequences, and RaxML8.2.10 was used to generate the Maximum Likelihood (ML) tree, and MrBayes on XSEDE to run the Bayesian Posterior Probability (PP). Ultrastructural analysis was realized on transmission electronic microscopy on FIOCRUZ/BA.MV and PP analysis showed an independent clade, grouping the sequence obtained in this study with a "Filamentous cyanobacterium" (GenBank GQ243430), with 88% and 100% support values, respectively. Ultrastructural analysis highlighted the presence of thylakoids parietally arranged to the cross-wall. This data, allied with the filamentous form, put our strain inside the Order Synechococcales and at the family Pseudanabaenaceae. The morphological features such as the apical cell form, cell morphometry, the marine habitat, and the epiphytic habit are not found in any other genus to Pseudanabaenaceae. This results suggests the recognition of a new to science genus with two species, being ALCB 125750, described in this study.

Key words: Cyanobacteria; Marine Environments; Taxonomy

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## **1 - INTRODUÇÃO GERAL**

### **1.1 - CARACTERIZAÇÃO GERAL DAS CIANOBACTÉRIAS**

As cianobactérias, também conhecidas como algas azuis ou algas azuis-esverdeadas, é um grupo filogeneticamente antigo de organismos procarióticos fotossintetizantes produtores de oxigênio, tendo surgido entre 2.7 e 3.5 bilhões de anos atrás (Brooks *et al.* 1999; Schopf & Packer, 2010; Komárek, 2016). A principal propriedade comum a todas as cianobactérias é a sua capacidade fotossintética, aliada à liberação de oxigênio molecular (fotossíntese oxigênica) (Baulina, 2012), sendo obrigatoriamente fotoautótrofas, obtendo carbono e energia por mecanismos fotossintéticos que são similares aos das plantas superiores, o que as diferem das outras bactérias (Holm-Hansen, 1968).

O principal componente do sistema fotossintético das cianobactérias é a clorofila *a*, localizada na membrana dos tilacoides, as ficobiliproteínas, clorofilas *b*, *d* e *f*, carotenoides e xantofilas atuam como pigmentos acessórios (Miyashita *et al.* 1996; Lee 2008; Chen *et al.* 2010; Gupta *et al.* 2013). Essa variedade de pigmentos faz com que as cianobactérias tenham uma grande amplitude na absorção do espectro de luz, sendo capazes de absorver ondas entre 400 e 750nm de comprimento (Chen & Blankenship, 2011).

A diversidade desse grupo é expressa através das suas propriedades morfológicas, bioquímicas e fisiológicas, o que possibilita o seu estabelecimento e persistência em grande variedade de habitats terrestres e aquáticos, incluindo aqueles com abundância ou escassez de recursos, sendo um componente dominante da comunidade fotossintética do solo, em regiões quentes ou frias e áridas. As cianobactérias encontram-se: nas camadas superficiais de solos agrícolas, onde ocorrem tipicamente com muitos outros organismos fotossintetizantes; em lagos salinos, alguns dos quais são ricos em metais pesados, como o cobre; e em fontes termais (Whitton & Sinclair, 1975; Lee, 2008; Paerl & Otten, 2013; Palinska & Surosz, 2014;).

#### **1.1.1 - CARACTERÍSTICAS MORFOLÓGICAS**

A forma mais simples que uma cianobactéria pode apresentar é a unicelular, de vida livre, ou encerrada em um envelope mucilaginoso. Com o passar do tempo, a

evolução dos caracteres morfológicos do grupo resultou na formação de um conjunto de células, chamado de tricoma, que, quando cercado por uma bainha, é denominado filamento (Lee, 2008). As formas unicelulares e filamentosas podem viver isoladas ou agregar-se em colônias, onde a sobreposição das células e/ou dos filamentos também tem sido usada como caracteres taxonomicamente informativos (Brodie & Lewis, 2007).

Diversas cianobactérias unicelulares, coloniais e filamentosas possuem um envelope circundando a sua membrana externa, sendo comumente chamada de bainha, glicocálice ou cápsula, e dependendo da sua consistência, pode ser referida como gel ou mucilagem. Esta camada protege as células contra dessecação em habitats que estão expostos à intensa radiação solar (Castenholz, 2001; Wehr & Shear, 2003; Lee 2008;). Nas cianobactérias terrestres, a bainha frequentemente se torna similar a um gel, sólida, ou lamelada, sendo frequentemente pigmentada. As algas desenvolvem bainhas com pigmentos sob alta intensidade luminosa. As bainhas incolores são desenvolvidas sob baixa intensidade luminosa, enquanto há aquelas que não produzem bainhas coloridas, não importando a intensidade da luz (Fogg, 1973).

Muitas das cianobactérias são diazotróficas, ou seja, são capazes de fixar o nitrogênio atmosférico (Bergman *et al.* 1997). Aquelas que são fixadoras podem ser divididas em duas formas, as heterocitadas e as homocitadas. As formas heterocitadas são capazes de fixar o N<sub>2</sub> pela ação de uma célula especializada, chamada heterocito (Bergman *et al.* 1997; Lee, 2008). Os heterocitos são fotossinteticamente inativos, ou seja, eles não fixam CO<sub>2</sub> nem produzem O<sub>2</sub>. A respiração e a fosforilação oxidativa ocorrem para que a fixação do nitrogênio não dependa exclusivamente da luz, apesar de ainda possuírem o fotossistema I para suprir a energia requerida para o processo de fixação do nitrogênio (Oren, 2014). O ambiente interno de um heterocito é virtualmente anóxico, o que é ideal para a ação da enzima nitrogenase, que é sensível à presença de O<sub>2</sub>, sendo uma especialização para a fixação do N<sub>2</sub> atmosférico ou dissolvido na água (Wher & Shear, 2003; Lee, 2008).

As formas filamentosas homocitadas (ou não-heterocitadas), também são capazes de fixar nitrogênio (Krumbein, 1975; Pearson, 1979; Gallon & Stal, 1992; Bergman *et al.* 1997; Stal & Wasmund *et al.* 2001). Nas cianobactérias homocitadas, a fixação do N<sub>2</sub> é feita através de células especializadas chamadas diazófitos, as quais produzem a enzima nitrogenase, que é sensível ao oxigênio, fazendo com que os

diazófitos realizem a fotossíntese enquanto há luz, somente fixando o nitrogênio no escuro, quando a fotossíntese cessa e o ambiente se torna anóxico (Fredriksson & Bergman, 1996; Komárek & Anagnostidis, 2005; Lee 2008).

### 1.1.2 - REPRODUÇÃO

A reprodução em cianobactéria ocorre somente de maneira assexuada, usualmente por fissão binária (Fogg, 1973). Pode ocorrer também por meio de células solitárias, que são liberadas por diversos tipos de espécies filamentosas, ou por fragmentação de um filamento ou do talo.

O modo de reprodução mais frequente nas formas filamentosas é a fragmentação, formando distintos segmentos de tricomas, nomeados hormogônios (Wehr & Shear, 2003). Os hormogônios podem ser produzidos por meio de células necridiais, que são células mortas e colapsadas, cuja morte é, provavelmente, geneticamente programada, em um processo de apoptose, para a produção de hormogônios ou ramificações falsas (Graham & Wilcox, 2000). As cianobactérias cocoides, produzem baeófitos, que se dividem sucessivas vezes, em múltiplos planos de divisão, sem crescer entre tais divisões (Lee, 2008).

### 1.1.3 - PRODUÇÃO DE TOXINAS

Algumas cianobactérias são capazes de produzir toxinas, *i. e.* cianotoxinas, como metabólitos secundários (van Apeldoorn, 2007), as quais são classificadas de acordo com o seu mecanismo de ação, podendo ser hepatotóxicas, neurotóxicas, dermatotóxicas ou promotoras da inibição da síntese de proteínas (Carneiro, 2008). As toxinas tendem a bioacumular em diferentes níveis tróficos, levando à biomagnificação trófica, uma vez que elas persistem na água mesmo após o desaparecimento da floração (Sarma, 2013).

Gêneros formadores de *blooms* incluem membros que produzem toxinas, como, por exemplo, *Anabaena* Bory ex Bornet & Flahault, (1886), *Anabaenopsis* V.V.Miller (1923), *Aphanizomenon* A.Morren ex É.Bornet & C.Flahault (1886 '1888'), *Cylindrospermopsis* G.Seenayya & N.Subba Raju (1972), *Microcystis* Lemmermann (1907), *Nodularia* Mertens ex Bornet & Flahault (1886), *Nostoc* Vaucher ex Bornet & Flahault (1886), *Oscillatoria* Vaucher ex Gomont (1892), *Planktothrix* Anagnostidis &

Komárek (1988), *Phormidium* Kützing ex Gomont (1892) e *Raphidiopsis* F.E.Fritsch & F.Rich (1929) (Codd, 2004; Sarma 2013). Gêneros formadores de biofilme e de tapetes também possuem membros toxigênicos, como *Phormidium* (Kützing ex Gomont), *Oscillatoria* Vaucher ex Gomont e *Lyngbya* C.Agardh ex Gomont (Sarma, 2013).

#### 1.1.4 - CLASSIFICAÇÃO TAXONÔMICA

A classificação taxonômica das cianobactérias era tradicionalmente realizada apenas através dos caracteres morfológicos (Gomont, 1892; Geitler, 1932; Desikachary, 1959; Komárek & Anagnostidis, 2005). Entretanto, o uso de técnicas modernas, como microscopia eletrônica e análises moleculares para a classificação taxonômica, promoveu muitas revisões taxonômicas em níveis de gênero e supragenérico (Komárek, 2008, 2010a, 2014; Komárek & Komárkova, 2004; Strunecký *et al.* 2011).

A classificação e nomenclatura das cianobactérias segue o Código Internacional de Nomenclatura Botânica (CINB), contudo, por conta da proximidade filogenética com as bactérias, as cianobactérias também têm sido classificadas de acordo com o Código de Nomenclatura das Bactérias (Rippka, 1979; Castenholz 2001), hoje denominado Código Internacional dos Procariotos (CINP). Ambos os códigos são frequentemente usados sem que haja uma coincidência entre eles, por eles possuírem regras diferentes, fazendo com que a conciliação entre eles seja problemática (Oren, 2004; Komárek, 2005).

Ao longo das últimas quatro décadas, características ecológicas, ultraestruturais e evidências moleculares têm influenciado substancialmente o conhecimento acerca das cianobactérias (Wehr & Sheath, 2003). Para garantir uma classificação taxonômica mais verossímil, Hoffman *et al.* (2005) propuseram o uso de uma abordagem polifásica, na qual são utilizadas sequências gênicas aliadas à análise de caracteres morfológicos, ultraestruturais e os dados ecofisiológicos, provê uma identificação mais precisa para a descrição de novos táxons e revisão daqueles já estabelecidos, como apresentado em Taton *et al.* (2006), Li *et al.* (2008), Komárek *et al.* (2009, 2010a, 2014), Moustaka-Gouni *et al.* (2010), Zapomělová *et al.* (2010), Vaz (2014) e Caires *et al.* (2018).

Dentre os grupos descritos a partir de uma abordagem integrativa, encontra-se a ordem Synechococcales. Essa ordem é um grande grupo não-monofilético de cianobactérias, com mais de 70 gêneros, incluindo formas unicelulares e filamentosas. Contudo, não há dados de sequências gênicas para a maioria desses táxons. (Komárek

*et al.* 2014). A principal característica que diferencia os membros dessa ordem para os representantes de outras ordens é o arranjo parietal dos tilacóides, junto com a forma dos filamentos. Isso por que a ordem Spirulinales também apresenta o mesmo tipo de arranjo tilacoidal, mas os filamentos são em forma de espiral (Komárek *et al.* 2014). Dentro da ordem Synechococcales encontram-se as famílias Pseudanabaenaceae Anagnostidis & Komárek (1988) e Leptolyngbyaceae Komárek *et al.* 2014, que representam o caráter não-monofilético dos representantes desse grupo.

Na família, Pseudanabaenaceae um dos gêneros mais representativos é *Pseudanabaena* (Lauterborn), contendo 110 espécies válidas, enquanto a família Leptolyngbyaceae é constituída por 14 gêneros com 60 espécies válidas (Komárek *et al.* 2014; Guiry & Guiry 2017). Ambas as famílias apresentam tricomas cilíndricos, unisseriados e isopolares, com variação apenas entre os diâmetros celulares dos seus representantes. Além destas características morfológicas similares, a presença de bainha facultativa e ausência de aerótopos (Komárek & Anagnostidis 2005) tornam esses grupos difíceis de distinguir somente através da morfologia.

## **1.2 - ESTUDOS SOBRE CIANOBACTÉRIAS MARINHAS BENTÔNICAS NA COSTA BRASILEIRA**

O litoral brasileiro possui aproximadamente 9.000 km de extensão - entre as latitudes 4°N e 34°S (Short & Klein, 2016), provendo grande quantidade de habitats (e.g. recifes de corais, manguezais) (Leão & Dominguez, 2000), os quais possibilitam a ocorrência de cianobactérias bentônicas. Os morfotipos marinhos bentônicos filamentosos formam colônias macroscópicas, comumente, uniespecíficas e, algumas vezes, florações potencialmente nocivas (Suda *et al.* 2013). As cianobactérias marinhas bentônicas ocorrem largamente em ecossistemas lênticos e lóticos, com hábitos epipélicos, episâmicos e epilíticos, ou epifitando algas filamentosas, musgos ou plantas vasculares (Wehr & Shear, 2003).

As cianobactérias bentônicas do litoral brasileiro têm sido pouco estudadas e sem uma grande abrangência de estudos deste grupo, levando a um desconhecimento da real biodiversidade do grupo no Brasil (Crispino & Sant'Anna, 2006). Dentre as cianobactérias marinhas bentônicas comumente excluídas dos levantamentos florísticos para a costa brasileira, encontra-se a família Pseudanabaenaceae. Esta família apresenta

apenas cinco espécies referidas para este ambiente de acordo com o sítio *online* Flora do Brasil (2020), o que pode não condizer com a real diversidade desta família para o litoral do Brasil.

Considerando a escassa quantidade de trabalhos sobre a família Pseudanabaenaceae para o ambiente marinho brasileiro, atrelado ao tamanho da costa do país e à quantidade de habitats que ela oferece, pressupõem-se um enorme déficit no conhecimento a respeito desse grupo potencialmente nocivo de cianobactérias.

Segundo Komárek *et al.* (2014), uma taxonomia na qual as espécies são bem caracterizadas usando uma abordagem polifásica, é fundamental para a construção de gêneros monofiléticos, como observado em Caires *et al.* (2018), na proposição do novo gênero marinho bentônico de cianobactéria *Neolyngbya*. A existência de linhagens diferentes filogeneticamente, mas que apresentam caracteres morfológicos convergentes são, algumas vezes, difíceis de serem evidenciados sem os dados moleculares, reforçando a importância de tal abordagem para a construção de relações filogenéticas verossímeis.

## **2 - Objetivos:**

1. Caracterizar o táxon encontrado com base em aspectos morfológicos, ultraestruturais, moleculares e ecológicos; para uma delimitação específica mais precisa do mesmo;
2. Contribuir para o conhecimento da diversidade de cianobactérias marinhas bentônicas da costa brasileira

1 **CAPÍTULO ÚNICO**

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3

4 **Description of new genera and specie of marine homocytous cyanobacteria from**  
5 **the Brazilian coast**

6

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20 **ABSTRACT**

21 Cyanobacteria is a group particularly challenging to classify because of its long and  
22 complex evolutionary history, and had its classification performed on morphological  
23 characters. Its evolutionary history is better reflected using an integrative approach  
24 combining morphological, ultrastructural and molecular data, because once the  
25 convergent evolution occurs, it often leads to taxonomical errors. This study aimed to  
26 analyze the genetic and morphological features of a Brazilian strain of cyanobacteria.  
27 The ALCB 125750 strain was collected at Pirambúzios beach (Rio Grande do Norte  
28 state, Brazil) (06°00'44.62"S x 35°06'26.48"W). The strain was put in SWBG-11 liquid  
29 medium for cultivation, under 14:10h (light:dark) photoperiod and temperature of  
30 25±1°C. To molecular analyzes, DNA extraction were realized followed by PCR to  
31 amplify the 16S rRNA. PCR products were cloned using the pGEM®-T Easy Vector  
32 Systems. To phylogenetic analyzes, the sequence obtained was added to the sequence  
33 matrix acquired from NCBI. Ultrastructural analysis was realized on transmission  
34 electronic microscopy. ALCB 125750 strain constituted a clade with a “Filamentous  
35 cyanobacterium” sequence from USA (GenBank GQ243430). Ultrastructural analysis,  
36 highlighted the presence of thylakoids arrangement parietally to the cross-wall. This  
37 data with the filamentous form put our strain into family Pseudanabaenaceae.  
38 Morphological features, such as the apical cell morphometry, marine habitat, and  
39 epiphytic habit, aren't found in any other genus of this family, suggesting the  
40 recognition of new genus and two species, one of which obtained in this study.

41 Key words: Biodiversity, Cyanobacteria; Marine Environments; Phylogeny; Taxonomy  
42 Abbreviations: TEM – Transmission Eletronic Microscopy; PCR – Polimerase Chain  
43 Reaction; NCBI – National Center for Biotechnology Information.



## 44 INTRODUCTION

45 Cyanobacteria is a group particularly challenging to classify mostly because of its long  
46 and complex evolutionary history (having appeared between 2.7 to 3.5 Bya). Because of  
47 this, the morphological identification between individuals of this group became difficult  
48 (Schopf and Packer, 1987; Broks et al. 1999; Komárek et al. 2014).

49 The Brazilian tropical coast offers one of the most diverse habitats that propitiate the  
50 occurrence of many cyanobacterial eco and morphotypes (Leão and Dominguez, 2000).  
51 However this area is very poorly studied, thus very little of the diversity and distribution  
52 of these organisms are known. The cyanobacteria constitute a very important  
53 component in a variety of communities, as epiphytic forms, epilithic and also as  
54 microbial mats including coral reefs (Charpy et al. 2012).

55 The marine benthic morphotypes form macroscope colonies, commonly unispecific and  
56 sometimes potentially harmful blooms (Suda et al. 2013). In this kind of environment,  
57 the morphological variations are the result of the characteristic of site where they occur,  
58 such as luminosity, water temperature and substrate. Cyanobacterial classification was  
59 performed based mostly on morphological characters (Gomont, 1892; Geitler, 1932;  
60 Desikachary, 1959; Komárek and Anagnostidis, 2005), however similar morphotypes  
61 can occur under very different ecological conditions, while different morphotypes can  
62 occur under the same ecological conditions, which generates a taxonomical confusion  
63 (Kling and Watson, 2003).

64 The classification system of this group was radically modified with the introduction of  
65 electronic microscopy and genetic methods to cyanobacterial taxa characterization,  
66 (Komárek et al. 2014). Hoffmann et al. (2005) suggested an integration between

67 morphological, ultrastructural and molecular data, which they denominated *polyphasic*  
68 *approach*, to better reflect the evolutionary inside cyanobacteria. However, this  
69 approach hasn't been used to examine the homocyted form of marine cyanobacterial  
70 strains, however has been mostly applied to examine the filamentous heterocytous form  
71 (Genuário et al. 2017).

72 Despite the length of the Brazilian coast, and the importance of a polyphasic  
73 characterization to study cyanobacteria, works with homocyted cyanobacteria are  
74 extremely rare, condition is not exclusive to Brazilian marine environment, to the  
75 marine environment worldwide, there is an equally poor number of published works,  
76 who deals with the homocytous form. Due the poor number of research, the size of the  
77 Brazilian coast and the quantity of habitats that's provided by them, there's a gap about  
78 the knowledge of this group, underestimating its real diversity.

79 This underestimated diversity highlights the necessity of further investigating  
80 cyanobacterial species using combined tools to show the real biodiversity of the  
81 homocyted cyanobacteria on the Brazilian coast. Thus, the goal of this study is describe  
82 a new genus and a new species (ALCB125750) using a polyphasic approach.

## 83 MATERIALS AND METHODS

### 84 Cyanobacterial cultivation and sampling

85 The sampling was realized in the tropical region of the Brazilian coast at Pirambúzios  
86 beach (Rio Grande do Norte state) (06°00'44.62"S x 35°06'26.48"W). The collecting  
87 was performed in mediolittoral zone during spring tide. Metal spatula was used to  
88 collect the epilithic samples. The refractometer (Biobrix<sup>®</sup> 211) was used to measure the  
89 water salinity. The samples was stored in seawater and put in liquid medium SWBG-11

90 after being transported to the laboratory (Rippka et al. 1979). To stimulate the growth of  
91 the strain, vitamin B<sub>12</sub> (2 mL L<sup>-1</sup> of a stock 1 mg mL<sup>-1</sup>, Sigma Aldrich®) was added to  
92 the medium. Filaments were isolated and put to grow under a photoperiod of 14:10  
93 (light:dark), and irradiation of 30 μmol photons m<sup>-2</sup> s<sup>-1</sup> (fluorescent white bulbs), and  
94 temperature of de 25 ±1°C (Jacinavicius et al. 2012). All isolated strains are kept in  
95 cultivation chambers maintained at the Biotechnology Laboratory (LaBBiotec) at the  
96 Universidade Federal da Bahia (UFBA), Brazil. Part of the strains were preserved at  
97 formaldehyde (4%) and laid up at Alexander Leal Costa Herbarium (ALCB), at the  
98 same institution.

#### 99 Morphological and ultrastructural analyzes

100 To morphological and ultrastructural analyzes, the strains were studied under a  
101 microscope with QImaging GO-3 digital câmera (Olympus CX31RTS5®, Tokyo,  
102 Japan). The filaments of every strain were measured using the AxioVs40 v4.8.2.0  
103 imaging system (Carl Zeiss, Jena, Germany) program. To the Transmission Electronic  
104 Microscopy (TEM) the cyanobacteria filaments were fixed overnight in Karnovsky  
105 (final concentration: 2% of glutaraldehyde, 0,05 M of sodium cacodylate – pH 7.2,  
106 0,001 M calcium chloride and paraformaldehyde at 2%) (Karnovsky, 1965). Biomass  
107 was washed three times with 0.01 M sodium cacodylate buffer and post-fixed with 2%  
108 osmium tetroxide in the same buffer for 2 h at room temperature. Using the same buffer,  
109 we repeated the washing, and the biomass was maintained at 4°C in 2.5% uranyl acetate  
110 overnight. We carried out with dehydration of the filaments with acetone series (30, 50,  
111 70% for 5 min each, and 90 e 100% for 20 min each), and embedded the filaments in  
112 Poly/Bed® 812 Embedding Kit resin (Polysciences, Warrington, USA). We obtained  
113 ultrathin sections (ultramicrotome; Leica®, Mannheim, Germany), which were post-

114 stained with 2.5% uranyl acetate and lead citrate. We analyzed the samples with a  
115 JEOL® 1230 transmission electron microscope (JEOL, Tokyo, Japan).

116 The identification of genus and species was accomplished using specific literature for  
117 the group. The classification system used was the proposed by Komárek *et al.* (2014).

118 Molecular and phylogenetic analyzes

119 The strain was macerated in liquid nitrogen for the genomic DNA extraction following  
120 Fiore *et al.* (2000). In PCR reactions, almost complete 16S rRNA and 16S-23S ITS  
121 regions was successfully amplified using the primers 27F1 (Neilan *et al.* 1997) and  
122 23S30R (Taton *et al.* 2003), in a Veriti® 96 -Well Thermal Cycler (Applied  
123 Biosystems®, Foster City, California, USA). For PCR reactions, we used 10 ng of  
124 genomic DNA, 0.5 µM of each primer, 200 µM of dNTPs, 2.0 mM of MgCl<sub>2</sub>, 1 × PCR  
125 buffer, and 1.5 U Taq DNA polymerase (Invitrogen, São Paulo, Brazil); the total  
126 volume was 25 µL. Amplifications were achieved under the following conditions: one  
127 cycle of 5 min at 94 °C; 10 cycles of 45 s at 94 °C, 45 s at 57 °C, and 2 min at 72 °C; 25  
128 cycles of 45 s at 94 °C, 45 s at 54 °C, and 2 min at 72 °C; the final elongation step of 7  
129 min at 72 °C. The PCR products were analyzed by running agarose gels (1 % w/v),  
130 using the Low DNA mass ladder (Invitrogen, Carlsbad, CA, USA), and they were  
131 cloned using the pGEM-T Easy vector system (Promega, Madison, WI, USA),  
132 according to the manufacturer's protocol. For blue-white colony screening, *Escherichia*  
133 *coli* chemo-competent DH5α cells were transformed and plated (Sambrook and Russell,  
134 2001). Plasmids containing gene fragments were extracted using the alkaline lysis  
135 method (Birnboim and Doly, 1979). We sequenced plasmidial DNA with the plasmidial  
136 primers M13F/M13R, and the 16S rRNA with the follow internal primer sets to cover

137 the entire fragment: 357F/357R, 704F/704R and 1114F/1114R (Lane, 1991). Genetic  
138 sequencing was done at Instituto Gonçalo Muniz (FIOCRUZ – Bahia State/Brazil).

139 To compose de matrix data, 63 sequences of the representatives of the orders of  
140 cyanobacteria were obtained from GenBank (www.ncbi.nlm.nih.gov) following the  
141 division by Komárek et al. (2014) were used to the phylogenetic analyzes. The total of  
142 64 operational taxonomic unities was achieved. The bacteria *Escherichia coli* was  
143 selected as the outgroup.

144 The evolutionary model (GTR+I+G) was identified by Mr.Modeltest 2.3 (Nylander,  
145 2008) using the Akaike Information Criterion (AIC) (Posada and Crandall, 1998). The  
146 reconstruction of the phylogenetic trees by the Maximum Likelihood (ML) algorithm  
147 (Stamatakis *et al.* 2008) was done using the RaxML 8.2.10 (Stamatakis, 2006), which  
148 was tested with 1.000 bootstrap. MrBayes 3.1 software (Ronquist et al. 2012) was used  
149 to do the Bayesian analyzes (Mau *et al.* 1999). The results generated by this  
150 phylogenetic analyze were expressed in bootstrap percentages (BP) to Maximum  
151 Likelihood, and posterior probability (PP) to the Bayesian analyzes. Values less than  
152 75% are not shown.

## 153 RESULTS

154 Based on a polyphasic characterization, we described a new genus of homocytous  
155 marine cyanobacteria which is included in Pseudanabaenaceae family. This genus  
156 includes two new species, one from USA and another from the Brazilian coast. Only the  
157 Brazilian new species is described in this study. The new species from the USA based  
158 on the “Filamentous cyanobacterium” sequence (GenBank GQ243430), is not described  
159 in present study, because we could not access the sample for either morphological,

160 ultrastructural nor ecological analyzes, because the study that describes this  
161 “Filamentous cyanobacterium” was not published yet.

162 Morphological and Ultrastructural Analyzes

163 *Genus nov.* ALCB 125750 (Pseudanabaenaceae, Synechococcales) V. L. Araujo, J.M.C.  
164 Nunes, T. A. Caires

165 Diagnosis: thallus forming green mats, filaments straight to curved, unbranched, cells  
166 always short, apical cells slightly to intensively rounded.

167 *Species nov.* ALCB 125750 (Pseudanabaenaceae, Synechococcales) V. L. Araujo,  
168 J.M.C. Nunes, T. A. Caires (Fig. 1)

169 Description

170 Thallus forming mat olive-green to moss-green. Straight filaments, isopolar,  
171 unbranched. Trichomes slightly constricted at the cross-walls, 0,8-1,12 µm diam. Short  
172 cells 1,58 - 4,28 µm long, 2,5 times longer than large. Thylakoids arranged parietally.  
173 Apical cells slightly rounded.

174 Holotype: Brazil, Rio Grande do Norte state, Nísia Floresta city, Pirambúzios beach,  
175 06°00'44.62"S x 35°06'26.48"W. Coll. T.A.Caires 640. 15 April 2014 (ALCB 125750).

176 *Habit and habitat:* epipsammic or epiphytic in protected areas in mediolittoral zone.  
177 This species occurred epiphyting an algae of genus *Bostrichia*.

178 Ultrastructural notes: thylakoids are arranged parietally to the cell wall. Cells slightly  
179 constricted at the cross-wall.

180 Phylogenetic Analyses

181 The 16S rRNA sequence obtained in this study was aligned with 63 other sequences of  
182 representatives of Cyanobacteria orders obtained from GenBank (Table 2). Our  
183 phylogenetic results show that our new genus based on ALCB 125750 strain forms an  
184 independent clade in the Maximum Likelihood (ML) and Bayesian posterior probability  
185 (BP) analyses, with 88% and 100% of support values, respectively, emerging as  
186 monophyletic group with high supports (Figure 2).

## 187 DISCUSSION

188 Concerning to results of the ultrastructural analysis, it showed that our strain has  
189 parietal arrangement of thylakoids, such as described for Synechococcales by Komárek  
190 *et al.* (2014). Alone, this characteristic is not very helpful to identify in which order our  
191 new genus ALCB 125750 must be included, because other genera also possess this  
192 thylakoidal arrangement, *e. g.* *Spirulina*, *Gloeothece* and *Synechococcus*, all three  
193 belonging to different orders, the first two belongs to the orders Spirulinales, and  
194 *Synechococcus* belongs to Synechococcales. (Komárek and Kaštovsk, 2003; Komárek  
195 *et al.* 2014). However, the members of the order Spirulinales are known to have  
196 trichomes in coiled form, while our new genus has straight trichomes. Thus, the form of  
197 the trichome combined with the parietal arrangement of thylakoids put our new genus  
198 ALCB 125750 into the order Synechococcales.

199 The new genus ALCB 125750 shows some morphological features similar to the ones  
200 found inside the family Pseudanabaenaceae (Anagnostidis *et* Komárek), such as the  
201 solitary or in colony trichomes, forming mats, straight e without sheaths and the cells  
202 are longer than wider, and cylindrical (Table 1). These similar characters make the  
203 identification of *Pseudanabaena*-like morphotypes very puzzling, demonstrating that

204 only morphological characters are not trustworthy for generic and specific  
205 identification. We ran an identification key, based on Komárek & Anagnostidis (2005)  
206 and the result shows that ALCB 125750 present characteristics that are quite similar to  
207 species that belongs to the genus *Pseudanabaena*, notably *P. catenata* (Lauterborn), and  
208 *P. minima* (G., S., An) Anagnostidis. However, ALCB 125750 presents morphometric  
209 characteristics that none of the known *Pseudanabaena* species possess (Table 1).

210 The family Pseudanabaenaceae is a clade that, according with Komárek & Anagnostidis  
211 (2005), is characterized by parietal thylakoids, trichomes without sheaths or only singly  
212 in one sheath, small trichomes (typically less than 4  $\mu\text{m}$ ). Within this family, individual  
213 genera are typically circumscribed based on a single morphological character  
214 (Perkerson III, 2010).

215 In relation to the ecological characteristics, our new species ALCB 125750 was  
216 collected in a marine benthic environment as an epiphyte on *Bostrychia* sp. red algae,  
217 while *Pseudanabaena catenata* is found in freshwater, as an epipelagic, sapropelic or  
218 periphytic in freshwater. *Pseudanabaena minima* is found in ponds, moist soils, swamps  
219 or salt fields, as benthic and epipelagic. None of these two species has yet been cited for  
220 the Brazilian marine benthic environment, which demonstrates the importance of  
221 ecology on the characterization of this new genus and species.

## 222 Phylogenetic Analyses

223 The genus ALCB 125750 formed a supported clade in the phylogenetic analyses,  
224 backing up the establishment of this new taxon (Fig 2). This clade (colored in green)  
225 showed high values of bootstrap and Posterior Probability (88% and 100%,  
226 respectively). The clade of the new genus is composed by the species: ALCB 125750



227 and a “Filamentous cyanobacterium” sequence (GenBank GQ243430) from the USA.  
228 The “Filamentous” cyanobacterium” was not analyzed because the type material has not  
229 been published.

230 Bayesian and Maximum Likelihood analysis of 64 OTUs clearly exhibited the  
231 separation of ALCB 125750 from other genera in the Synechococcales order (colored in  
232 gray), such as *Leptolyngbya* (Anagnostidis & Komárek), *Haloleptolyngbya*  
233 (P.K.Dadheech, H.Mahmoud, K.Kotut & L.Krienitz), *Halomicronema* (Abed, Garcia-  
234 Pichel & Hernández-Mariné) and *Nodosilinea* (R.B.Perkerson & D.A.Casamatta).

235 The presence of members of the genera *Leptolyngbya* and *Pseudanabaena* in distinct  
236 clades throughout the tree, including as the sister group (colored in brown) of ALCB  
237 125750, corroborate the polyphyletic condition of those groups (Castenholz, 2001;  
238 Komárek and Anagnostidis, 2005), showing that unrelated lineages possess highly  
239 comparable morphological features, which can be the result of convergent evolution,  
240 and not common descent (Engene et al. 2011). According to Perkerson III et al.  
241 (2010), the systematic and taxonomic problems within cyanobacteria occur due to  
242 convergent evolution and the lack of genera-level autapomorphies. Moreover, Komárek  
243 et al. (2014) put a light over both groups regarding their polyphyletic condition and the  
244 lack of revision of them.

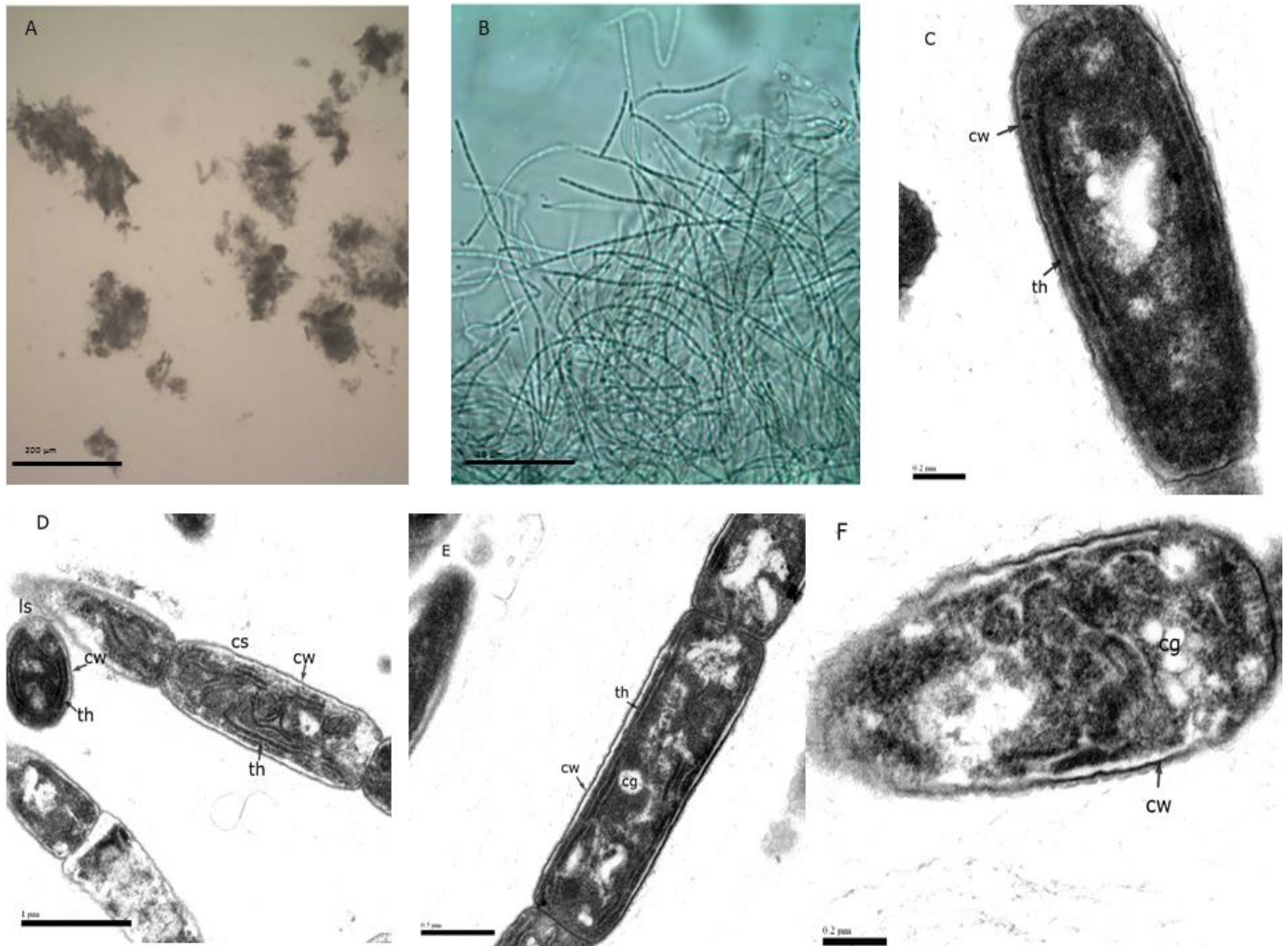
## 245 CONCLUSION

246 The combination of genetic, ultrastructural, and morphological features, it’s a very  
247 useful way to provide different evidence that allowed us to highlight differences infra  
248 and interspecific. The use of polyphasic approach allowed us to recognize and describe  
249 a well-supported genus and species ALCB 125750, with two species. The species

250 ALCB 125750 is characterized by its small cell size, its marine benthic habit and the  
251 disposition of thylakoids. We also corroborate the polyphyly of the genus *Leptolyngbya*,  
252 that is present throughout the phylogenetic tree, except in the clade in brown.

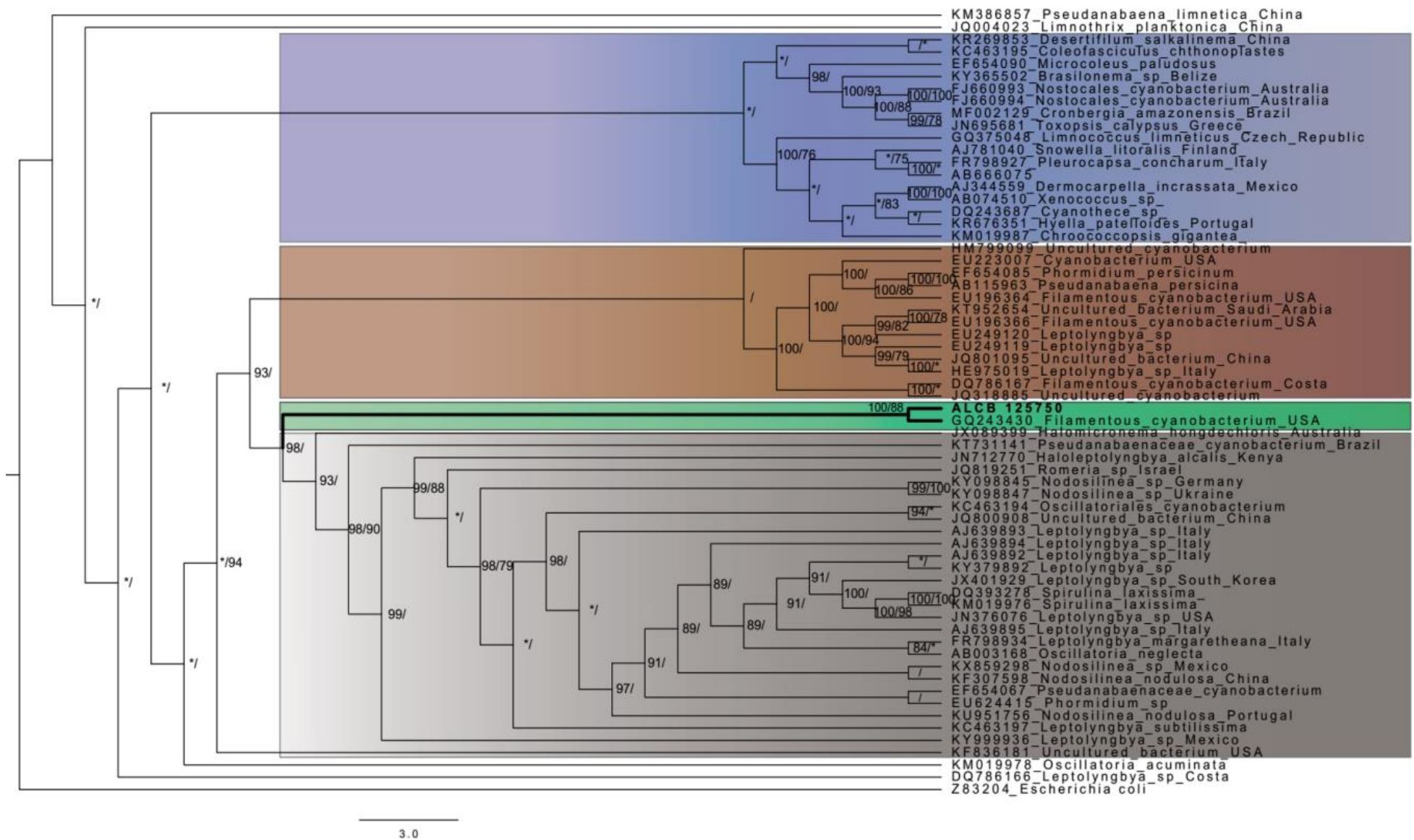
253 Molecular, ecological and ultrastructural data about the homocytous cyanobacteria are  
254 not abundant due the lack of studies to unveil the biodiversity of this cyanobacterial  
255 group. This setting makes the taxonomy of cyanobacteria very challenging.

256 Acknowledgments To CNPq/CAPES (National Council for Scientific and  
257 Technological Development/Coordination of Personal Enhancing in Superior  
258 Education) for the scholarship granted to the first author.



259

260 **Fig 1.** Characterization of the new genus and species ALCB 125750. **A.**  
 261 General aspects of the mat. **B.** Detail of the trichomes. **C-F.** Ultrastructural  
 262 sections (TEM). **C.** arrangement of the thylakoids (th), parietally to the cross-  
 263 wall (cw). **D.** Cross section (cs) and longitudinal sections (ls) of filaments. **E.**  
 264 Detail of the arrangement of the thylakoids (th), parietally to the cross-wall (cw).  
 265 **F.** The arrow indicates the cyanophycin granules (cg). Cell walls (cw);  
 266 thylakoids (th); cyanophycin granules (cg). Scale bars: A= 200 µm; B= 10 µm;  
 267 C=0,2 µm; D= 1 µm; E=0.5 µm; F= 0,2 µm



268 **Fig 2.** Maximum likelihood (ML) phylogeny based on the 16S rRNA gene sequences of Cyanobacteria (ntax = 64). Values are ML bootstrap values  
 269 (right) and Bayesian posterior probabilities (PP) converted to percentages (left). Values under 75% are not shown. New sequence obtained in this study  
 270 is indicated in bold; other sequences from Genbank, in which the access numbers are followed by names. \* = clade not supported. Blue= Clade formed  
 271 by representants of the order Nostocales. Brown= Sister group of the genus ALCB 125750. Green= Genus generated in this study. Gray =  
 272 Representants of the order Synechococcales.

273 **Table 1.** Comparison between ALCB 125750 and *Pseudanabaena* species who related to it (Komárek & Anagnostidis, 2005).

274 NI=Not informed

275	Characteristics	<i>P. catenata</i> Lauterborn 1915	<i>P. minima</i> (G.S. An.) Anagnostidis 2005	ALCB 125750 sp. nov
276	<b>Filament</b>	Trichome solitary or aggregated in small mats	Trichome solitary, or crowded in clusters, straight or almost straight,	
277				
278	<b>Filament width (µm)</b>	1,2-2 (2.2) wide	1.3-2,5 wide	0,8-1,12 wide
279	<b>Sheath</b>		NI	
280	<b>Trichom</b>	Strongly constricted; ungranulated;	Intensely constrict	Slightley constrict
281	<b>Trichom width (µm)</b>	40-200 long	1.5-4 long	1,58-4,28 long
282	<b>Cells: form and lenght (µm)</b>	Cylindrical, 1.5x3xlonger than wide; rarely isodiametric; (1.5) 2-6 (8- 10?)	1.5x longer than wide,	2,5x longer than wide
283				
284	<b>Apicall cell</b>	<i>Rounded or slightly conic</i>	Widely rounded	Slightly rounded
285	<b>Environment</b>	Freshwater, benthic,	Moist soils, benthic and epipelic in ponds, springs, swamps and salty fields	Marine, benthic

286 **Table 2.** Taxa included in the analysis, with locality and/or environment information, reference publication, and GenBank  
 287 accession numbers. The taxa sequenced in this study is presented in bold.

Taxon	Locality &/or Environment	Voucher/stra in	Reference publication	GenBank accessions
<i>Outgroup</i>				
<i>Escherichia coli</i>		562/	Ridell <i>et al.</i> (1995)	Z83204
<i>Ingroup</i>				
<b>640</b>	<b>Pirabúzios beach, Nísia Floresta city, Rio Gr&amp;e do Norte state, BRazil</b>	<b>ALCB 125750</b>	<b>This Study</b>	AB666075
<i>Snowella litoralis</i>	Finland	371757/0tu3 7s04	Rajaniemi- Wacklin,P. <i>et al.</i> (2006)	AJ781040
(Häyrén) Komárek & Hindák				

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<i>Limnococcus</i> <i>limneticus</i>	Czech Republic: Svet	665911/Svet 06	Komarkova, J. <i>et al.</i> (2010)	GQ375048
(Lemmermann) Komárková, Jezberová, O.Komárek & Zapomelová				
<i>Microcoleus</i> <i>paludosus</i>		450472/SA G 1449-1a	Siegesmund, M.A. <i>et al.</i> (2008)	EF654090
Gomont				
<i>Chroococcopsis</i> <i>gigantea</i>		1521635/SA G 12.99/	Friedl, T. Unpublished	KM019987
Geitler				
<i>Coleofasciculus</i> <i>chthonoplastes</i>		1352644/Ec FYyyy500	Dojani, S. <i>et al.</i> (2013)	KC463195

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(Thuret ex  
Gomont)  
M.Siegesmund,  
J.R.Johansen &  
T.Fiedl

*Toxopsis  
calypsus*

Greece: Peloponnese

1108047/PL  
F

Lamprinou, V. *et  
al.* (2012)

JN695681

Lamprinou,  
Skaraki,  
Kotoulas,  
Economou-Amili  
& Pantazidou

*Cronbergia  
amazonensis*

Brazil

1983329/C  
MAA1598

Genuario, D., B.  
(2018)

MF002129

Genuário,  
Sant'Anna &  
I.S.Melo

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<i>Hyella patelloides</i>	Portugal: Moledo do Minho beach	945734/LEG E 07179	Brito, A. <i>et al.</i> (2017)	KR676351
Ramos, Brito <i>et</i> Kaštovsky				
<i>Dermocarpella incrassata</i>	Mexico	54303/SAG 29.84; PCC 7326	Fewer, D. <i>et al.</i> (2002)	AJ344559
Lemmermann				
<i>Haloleptolyngbya álcalis</i>	Kenya: Lake Nakuru	1169407/KR 2005/106	Dadheech, P. K. <i>et al.</i> (2012)	JN712770
P.K.Dadheech, H.Mahmoud, K.Kotut & L.Krienitz				
<i>Brasilonema</i> sp.	Belize	1980447/BZ	Johansen, J., R. <i>et</i>	KY365502

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		-HDL-007	<i>al</i> , (2017)	
<i>Desertifilum salkalinema</i>	China	1736058/CH AB7200	Cai, F., -F. <i>et al.</i> (2017)	KR269853
F.-F.Cai & Li, nom. inval (holotype = <i>Desertifilum tharensis</i> Dadheech & Krienitz).				
<i>Pleurocapsa concharum</i>	Italy:Firenze, Villa La Pietra	980868/VP3 -02	Cuzman, O., A. <i>et al</i> , (2010)	FR798927
Hansgirg				
<i>Cyanothece</i> sp.		354330/	Zhang, Y. Unpublished	DQ243687

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<i>Xenococcus</i> sp.		179406/PC C 7307	Seo,P. & Yokota,A.  Unpublished	AB074510
<i>Limnothrix planctonica</i> (Woloszynska) Meffert	China	1131003/CH AB753	Zhu,M.L. <i>et al.</i> (2012)	JQ004023
<i>Pseudanabaena limnetica</i>  (Lemmermann) Komárek	West Lake, Hangzhou	1572394/CH AB774	Yu,G. Unpublished	KM386857
<i>Pseudanabaena ceae</i> cyanobacterium		450516/isola te=DPG1- KK5	Siegesmund, M. A., <i>et al.</i> 2008	EF654067

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<i>Nodosilinea nodulosa</i>	China: Hong Kong, South China Sea	416010/UTE X B 2910	Muhlsteinova, R. <i>et al.</i> (2014)	KF307598
(Z.Li & J.Brand) Perkerson & Casamatta				
<i>Leptolyngbya</i> sp.	Italy:Emilia-Romagna, Imola, Bubano Basin	271630/0BB 30S02"	Castiglioni, B. <i>et al.</i> (2005)	AJ639892
Uncultured bacterium	Saudi Arabia	77133/clone =AG2_B08	Miyake, S. <i>et al.</i> (2016)	KT952654
Filamentous cyanobacterium	USA: NN Dry Rocks Reef, Northern Florida Keys	492282/73-2	Myers, J., L. & Richardson, L., L. (2009)	EU196366
Uncultured cyanobacterium	Puerto Rico Trench	1211/clone= PRTBB8722	Eloe, E., A., <i>et al.</i> (2011)	HM799099
Filamentous	Costa Rica	397329/LLi7	Finsinger, K. &	DQ786167

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cyanobacterium		1	Hess,W., R. Unpublished	
<i>Leptolyngbya</i> sp.	South Korea	1229172/KI OST-1	Kim, J., H., Unpublished	JX401929
<i>Leptolyngbya</i> sp.	Italy:Formentera	1221563/CC AP 1442/1	Gachon, C., M., M. Unpublished	HE975019
Filamentous cyanobacterium	USA: Great Salt Lake, Utah	658598/GSL 035	Beer, L., L. Unpublished	GQ243430
<i>Leptolyngbya</i> sp.	Mexico: Cuatrociénegas	1983256/CC M 4"	Barrera-Rojas, J. Unpublished	KY999936
Nostocales cyanobacterium	Australia: Heron Isl&	994809/HI1 5	Dominguez- Escobar, J., (2011)	FJ660993
<i>Leptolyngbya</i> <i>subtilissima</i>		1352645/Ec FYyyy700	Dojani, S. <i>et al.</i> (2013)	KC463197

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(Hansgirg) Komárek				
Uncultured bacterium	China: Yellow River Delta	77133/clone =BJGMM- 3s-467	Jia,F. <i>et al.</i> Unpublished	JQ801095
<i>Phormidium</i> sp.		545217/SA G 61.90	Siegesmund, M., A. <i>et al</i> , (2008)	EU624415
<i>Leptolyngbya</i> sp.	Bahamian marine stromatolite	487779/HB C1	Foster, J., S. <i>et al.</i> (2009)	EU249120
<i>Leptolyngbya</i> sp.	Italy:Emilia-Romagna, Imola, Bubano Basin	271632/0BB 32S02	Castiglioni, B. <i>et</i> <i>al</i> , (2005)	AJ639894
Uncultured cyanobacterium		1211/clone= T-Pc3	Jin, H. <i>et al.</i> Unpublished	JQ318885

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Pseudanabaena ceae cyanobacterium	Brazil: Bertioga/SP	1716595/CE NA320	Alvarenga, D., O., <i>et al.</i> , (2016)	KT731141
Filamentous cyanobacterium	USA: Horseshoe Reef, Northern Florida Keys	492280/FLK 9	Myers, J., L. <i>et al.</i> (2009)	EU196364
<i>Nodosilinea</i> sp.	USA: California, Imperial Valley	1115757/BL 0902	Taton, A. <i>et al.</i> (2012)	JN376076
Cyanobacterium	USA: Florida, Northern Florida Keys, Conch  Shallow Reef	481519/62-2	Myers, J., L. <i>et al.</i> (2009)	EU223007
<i>Nodosilinea</i> sp.		1963213/isol ate=Prim-5- 5	Mikhailyuk, T. <i>et</i> <i>al.</i> Unpublished	KY098847
<i>Leptolyngbya</i>		1930147/RK	Albrecht, M., <i>et al.</i>	KY379892

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sp.		_2_3	(2017)	
<i>Nodosilinea</i> sp.	Mexico	1902582/PC 471	Vazquez-Martinez, J. Unpublished	KX859298
Uncultured bacterium	China: Yellow River Delta	77133/clone =BJGMM- 3s-118	Jia, F. <i>et al.</i> Unpublished	JQ800908
<i>Leptolyngbya</i> sp.	Italy:Emilia-Romagna, Imola, Bubano Basin	271633/0BB 19S12	Castiglioni, B. <i>et</i> <i>al.</i> (2005)	AJ639895
Uncultured bacterium	USA	77133/ clone=INDI_ S_SPR_11H _(2)	Reihle, J. Unpublished	KF836181
<i>Spirulina</i> <i>laxíssima</i>		1236702/ SAG 256.80	Friedl, T. Unpublished	KM019976
G.S.West				

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<i>Halomicronema hongdechloris</i>	Australia: Shark Bay	1641165/C2 206	Chen, M. (2012)	JX089399
Chen, Li, Birch & Willows				
<i>Leptolyngbya</i> sp.		487783/HB C8	Foster, J., S. <i>et al.</i> (2009)	EU249119
<i>Pseudanabaena persicina</i> (Reinke ex Gomont) Anagnostidis		402770/SA G 80.79	Siegesmund, M. A., <i>et al.</i> (2008)	EF654085
<i>Romeria</i> sp.	Israel: Hula Nature Reserve	1192302/ KLL-H-201	Alster, A. Unpublished	JQ819251
Nostocales cyanobacterium	Australia: Heron Isl&	994810/ HI14	Dominguez-Escobar, J. <i>et al.</i> (2011)	FJ660994

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<i>Leptolyngbya</i> sp.	Costa Rica	397324/LLi1 8	Finsinger, K. & Hess, W.R. Unpublished	DQ786166
<i>Oscillatoriales</i> <i>cyanobacterium</i>		1352668/Ec FYyyy400	Dojani, S. <i>et al.</i> (2013)	KC463194
<i>Leptolyngbya</i> <i>margaretheana</i>  (G.Schmid) Anagnostidis & Komárek	Italy:Firenze, Piazza SS. Annunziata	980873/1T1 2	Cuzman, O., A. <i>et al.</i> (2010)	FR798934
<i>Nodosilinea</i> sp.	Germany: Warnemuende, Mecklenburg- Vorpommern, coast of Baltic Sea	1963211/isol ate=WD-4-2	Mikhailyuk, T. <i>et al.</i> Unpublished	KY098845
<i>Leptolyngbya</i>	Italy:Emilia-Romagna,	271631/0BB	Castiglioni, B. <i>et</i>	AJ639893

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sp.	Imola, Bubano Basin	24S04	<i>al.</i> (2005)	
<i>Oscillatoria neglecta</i>		454132/ IAM M-82	Ishida, T. <i>et al.</i> Unpublished	AB003168
Lemmermann				
<i>Nodosilinea nodulosa</i>	Portugal: Praia da Luz	1828761/LE GE 06101	Ramos, V. <i>et al.</i> Unpublished	KU951756
(Z.Li & J.Brand)				
Perkerson & Casamatta				
<i>Spirulina laxíssima</i>		1828761/SA G 256.80	Choi, G.,-G. <i>et al.</i> Unpublished	DQ393278
G.S.West				
<i>Oscillatoria acuminata</i> PCC		56110/SAG 1449-3	Friedl, T. <i>et al.</i>	KM019978

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6304		Unpublished	
<i>Pseudanabaena</i>	945773/SA	Chaw, S. <i>et al.</i>	AB115963
<i>persicina</i>	G 80.79	Unpublished	
(Reinke ex Gomont) Anagnostidis			

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## 5 – ANEXOS

### Normas da Revista ALGAE

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INTRODUCTION, MATERIALS AND METHODS, RESULTS (or OBSERVATIONS), DISCUSSION, ACKNOWLEDGEMENTS, and REFERENCES, although this may not be appropriate for some articles (such as some taxonomic papers). To avoid unnecessary errors, authors are strongly advised to use the "spell-check" and "grammar-check" functions of their word processor. Do your very best to use correct English grammar, spelling and punctuation; if you are not a native English speaker, you should have the text edited by someone, as the editors cannot always be expected to carry out major linguistic revision.

Research Articles should be prepared according to the following format:

**1. The title page** should contain a concise title, the name(s) and address(es) of the author(s), any necessary footnotes, and a short running title suitable for page headings.

- **Title.** Titles are often used in information-retrieval systems. Avoid abbreviations and formulae where possible. The title should have not more than 100 characters (ca. 15 words, 2 lines in print), and 150 characters at most. If the name of an organism is used in the title, an indication of its taxonomic position must be given. Nomenclatural authorities should only be used in titles when nomenclatural changes are being proposed.
- **Author names and affiliations.** Where the family name may be ambiguous (e.g., a double name), please indicate this clearly. Present the authors' affiliation

addresses (where the actual work was done) below the names. Indicate all affiliations with a lower-case superscript number immediately after the author's name and in front of the appropriate address for identification. Provide a valid address with ZIP or postal number for each affiliation, including the country name.

- **Corresponding author.** Clearly indicate who will handle correspondence at all stages of refereeing and publication, as well as post-publication. Ensure that telephone and fax numbers (with country and area code) are provided, in addition to the e-mail address and the complete postal address.

- **Present/permanent address.** If an author has moved since the work described in the article was done, or was visiting at the time, a "Present address" (or "Permanent address") may be indicated as a footnote to that author's name. The address at which the author actually did the work must be retained as the main affiliation address. Superscript letters are used for such footnotes.

- **Running title.** Provide a condensed running head with 3 to 6 words.

**2. The ABSTRACT** should not be more than 250 words and should be concise, informative and intelligible without reference to the main text. It should indicate the objectives, main results and conclusions of the paper. Do not repeat information in the title or make reference to the literature. Authorities for species names should be included in the abstract only for primarily taxonomic papers. Begin the abstract by stating the scientific question of concern. Explain the methods used to tackle the question. The results should be outlined briefly and put into a concise, broad perspective. Immediately after the abstract, list 5 to 8 **Key Words** (arranged alphabetically and separated by semicolons), using American spelling and avoiding general and plural terms and multiple concepts (avoid, for example, "and" or "of"). Include a section of **Abbreviations** after the keywords. List the abbreviation, followed by its meaning written out in full (e.g., DIN, dissolved inorganic nitrogen). Common abbreviations (e.g., DNA) do not need to be listed. The journal accepts standard abbreviations from the *Journal of Biological Chemistry*. All non-standard abbreviations should be listed alphabetically. The abbreviation is spelled out at first mention in the main text, and thereafter, only the abbreviation/acronym is used.

**3. The INTRODUCTION** must define the problem within the context of existing knowledge, state the objectives of the work and provide an adequate background; avoid a detailed literature survey or a summary of the results. It should not be a general review of the field, but it should provide essential background for those who are not experts in the particular area.

#### 4. MATERIALS AND METHODS

Describe the methodology used in the study in sufficient detail to allow the work to be reproduced by another scientist. Methods already published should be indicated by a reference, and only relevant modifications should be described. Whenever possible, give sources of materials in detail. If cultures are used, indicate the strain or clone number and the source. If study sites are mentioned, provide latitudes/longitudes. For materials and supplies (including software), indicate the source (company name, city, state, country) on first reference. In addition, list the model number for equipment used, as appropriate. Use metric units (*Système International d'Unités*, SI) and SI style (e.g.,  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ ,  $\mu\text{g L}^{-1}$ ). For description of laboratory procedures, the terms liter (abbreviated to L) and milliliter (mL) may be used. Do not use dots or full-stops between parts of the term. For more information on this see below.

#### 5. RESULTS

Only results pertinent to the subject may be included. Data must not be repeated in figures and tables. Rationale for undertaking certain aspects of the investigation, methods, techniques, and so on, must be excluded. As far as possible, cite each figure and table in the text in order of presentation (e.g., Figure 1 before Figure 2). When tables or figures are presented, cite only the significant results in the text. Tables of specimens studied or gene sequences used should be placed in supplementary materials. Photographs should only be included if necessary to illustrate results. Include statistical analyses or other indicators to enable assessment of the variance of replicates of the experiments. Names of new taxa must be followed immediately by the Latin description or, preferably, diagnosis (using only essential characters), with citation of holotype. For new combinations, basionyms should be cited in full.

For further guidance on taxonomic papers, please see the web site ([http://www.algae.kr/layout\\_taxonomy](http://www.algae.kr/layout_taxonomy)).

#### 6. DISCUSSION

This should explore the significance of the results in relation to the problem outlined in the introduction - not repeat them. Reference to illustrative material should be minimal and should be provided only when necessary to emphasize a specific interpretation. A **CONCLUSION** should be added if results and discussion are combined.

#### 7. ACKNOWLEDGEMENTS

This is to be used as necessary to acknowledge the source of financial grants in completing the study. The contribution of colleagues or institutions should also be acknowledged. Thanks to anonymous reviewers are not allowed.

#### 8. REFERENCES

In the text, refer to the author's name (without initial) and year of publication. When reference is made to a work by two authors, both names should be given using "and" (e.g., Lee and Kang 1986); for three or more author names, give the first author followed by "et al." and the year (e.g., Kim et al. 2010). Multiple references must be arranged in chronological order (e.g., Lee and Kang 1986, Kim 1990, Kim et al. 2000a, 2000b). Only cite articles or books already published or in press, not unpublished work "in preparation". The author is responsible for verifying the accuracy of unpublished citations.

In the list at the end of the paper, references in the **REFERENCES** section should be typed, double-spaced and in alphabetical order, with multiple references by the same author(s) arranged chronologically. If an author's name in the list is also mentioned with co-authors, the following order should be used: publications of the single author, arranged according to publication dates; publications of the same author with one co-author; publications of the author with more than one co-author. Publications by the same author(s) in the same year should be listed as 2000a, 2000b, etc. For Volume (Vol.) and Bulletin (Bull.), Arabic numerals should be used; the full number of pages should be given in the form of pp. 123-128. Type references flush left with an extra space between citations; they will

be formatted properly at the press. The name of the journal should be abbreviated according to the *World List of Scientific Periodicals* and the selection according to *Abbreviated Titles of Biological Journals* issued by the Biological Council or the *Biosis Serial Sources*, which is published each December and is available in most libraries (or go to <http://www.library.uq.edu.au/faqs/endnote/biosciences.txt>). If in doubt about any journal abbreviation, write out the journal title in full. The number of a fascicle in brackets after the volume number should be given only if the volume is not paginated consecutively. The titles of publications in non-Latin alphabets should be transliterated, and a notation such as "(in Korean)" or "(in Greek, with English abstract)" should be added. Work accepted for publication but not yet published should be referred to as "in press". References concerning unpublished data and "personal communications" should not be cited in the reference list but may be mentioned in the text. All publications cited in the text should be presented in a list of references following the text of the manuscript. The manuscript should be carefully checked to ensure that the spelling of author's names and dates are exactly the same in the text as in the reference list.

**The style to be used for references is as follows:**

#### Journal articles

Kim, K. H. & Lee, I. K. 2000. Mixed-phase reproduction in *Antithamnion sparsum* Tokida (Ceramiaceae, Rhodophyta) from Korea. *Algae* 15:183-193.

Mann, H., Mann, S. & Fyfe, W. S. 1987. Aragonite crystals in *Spirogyra* sp. (Chlorophyta). *J. Phycol.* 23:506-509.

#### Articles from books, conference reports, symposium proceedings, etc.

Give the title of the chapter, the editor(s) and title of the volume, the publisher and place of the publisher (not the location where the conference was held), and the pages of the chapter. The date cited must be the year of publication (not the year in which the conference was held).

#### Example :

Sheath, R. G. & Hambrook, J. A. 1990. Freshwater ecology. In Cole, K. M. & Sheath, R. G. (Eds.) *Biology of the Red Algae*. Cambridge University Press, Cambridge, pp. 423-453.

Conover, J. T. & Sieburth, J. McN. 1966. Effect of tannins excreted from Phaeophyta on planktonic animal survival in tide pools. In Young, E. G. & McLachlan, J. L. (Eds.) *Proc. 5th Int. Seaweed Symp.*, Pergamon Press, Oxford, pp. 99-100.

#### Book

Write the title of the book in lower case, and give the publisher, place of publication and pages. In the case of a book series, give the series editor as well.

#### Example :

van den Hoek, C., Mann, D. G. & Jahns, H. M. 1995. *Algae: an introduction to phycology*. Cambridge University Press, Cambridge, 623 pp.

#### Dissertations

Write the title in lower case, 'MS / PhD thesis / dissertation', and give the university and its location.

#### Example :

Boo, S. M. 1985. A systematic study on six tribes of eramiaceae (Rhodophyta, Ceramiales) in Korea. Ph.D. dissertation, Seoul National University, Seoul, Korea, 446 pp.

#### 9. Tables

Authors should take notice of the limitations set by the size and layout of the journal. Large tables should be avoided. Reversing columns and rows will often reduce the dimensions of a table. If many data are to be presented, an attempt should be made to divide them over two or more tables. Tables should be created using the Table function in the word processor (rather than using tabs). Tables should have a descriptive title at the top of each table. The title and table of contents must be double-spaced throughout, in 12-point font, and on a separate numbered page. Tables should be numbered consecutively with Arabic numerals according to their sequence in the text. The text should include references to all tables. Tables should not include vertical lines or shading; if either is essential, the material must be submitted as a figure for direct reproduction. Column headings should be brief, with units in parentheses. Tables and their captions should be self-explanatory; e.g., abbreviations and acronyms must be defined again. Any explanation essential to understanding the table should be given as a footnote at the bottom of the table, not in the heading. Include in the footnotes all non-standard abbreviations used and enough information for the table to be understood without undue recourse to the text. For table footnotes, use superscripted lower case letters (<sup>a</sup>, <sup>b</sup>, <sup>c</sup>) rather than symbols or numbers; asterisks (\*, \*\*, \*\*\*) can be used to indicate statistical significance. These can be applied in the case of figures. Statistical measures such as SD or SEM should be identified in the headings. If a table

provides data on biological species, its legend should begin with the full Latin name of that species.

#### 10. Figures

Line drawings, diagrams and photographs should be planned so that after reduction they will fit within either the width of one column (8 cm) or two columns (17 cm), and be no more than 25 cm in length. Normally, only previously unpublished illustrations are acceptable. Figures should be numbered in Arabic numerals consecutively as they are mentioned in the text (Fig. 1), (Figs 2 & 3), (Figs 1-4), etc. Each figure should be submitted as a separate Tagged Image Format (TIFF), Encapsulated PostScript (EPS) or MS Office files created at a resolution of 300 dpi (or 600 dpi for combination images). In the case of very large files, Joint Photographic Experts Group (JPEG) or Graphics Interchange Format (GIF) files may be submitted initially. Do not import the figures into the text file, but, instead, indicate their approximate locations directly on the manuscript. Combination images, or images that contain both line artwork (vector graphics) and halftones or photographs (bitmap graphics), must be supplied at a higher resolution to prevent image quality loss in the bitmap graphics, such as in the symbol keys. Please do not attempt to increase the resolution of a lower resolution figure by resaving it at 300 dpi or higher. The result may be pixilated or grainy images and poor text quality. Photographs (halftones) will not be reduced or enlarged; thus, they must be printed to fit in one column or two columns in width and be no more than 18–20 cm (8 inches) in length so that the legend will fit on the same page as the illustration. Plan line drawings and graphs to fit these dimensions after reduction, and with all lines, symbols and lettering bold enough to permit 1/2 to 2/3 reduction in size; many line drawings are reduced to the width of a single column. All terms, abbreviations, and symbols should correspond to those used in the text of the paper. It is preferable to include a symbol key on line drawings. Component figures within a plate should be labeled with letters, not numbers and should not be encircled (e.g., a, b, c, not 1, 2, 3, ①, ②, ③). Only use the following fonts in your illustrations: Arial, Courier, Times, Symbol. Do not use two shades of gray in histograms.

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## NOMENCLATURE AND UNITS

### Genus and Species Names

Genus and species names must be in italics. Generic names should only be abbreviated when immediately preceded in the text by the mention of the same species or another of the same genus. When referring to a species, do not use the genus name alone, unless you have previously defined it that way; be precise when using 'sp.' (singular) and 'spp.' (plural). All Latin binomials should be followed by the authorities in full, e.g. (J.E. Smith) Setchell & N.L. Gardner, when first used in the title or text (but not in the abstract), unless a large number of names with authorities are grouped in a table. To find the taxonomic author(s), check the Algaebase Web site (<http://www.algaebase.org>).

### Equations and Units

Use standard SI units. Compound units are given with the proper exponent without a period (e.g.,  $\text{gO}_2 \text{g}^{-1} \text{dw h}^{-1}$ ). Relations or concentrations (e.g., mg per l) must be given as ' $\text{mg l}^{-1}$ ' (not mg/l). Variables are usually italicized (except for Greek letters). Italicization should be consistent in normal, superscript and subscripted text. Leave one blank space on either side of '=', '>', ' $\pm$ ' etc. where these denote equalities or inequalities. SI (metric) units must be used. Leave a space between numerals and their units (e.g., 10 mm). Example: ' $p < 0.05$ ,  $r^2 = 0.879$ ' (not ' $p < 0.05$ ,  $r^2 = 0.879$ '); but: 'we studied organisms of size  $< 0.5 \mu\text{m}$ '.

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