Morphology and phylogenetic relationships of *Micractinium* (Chlorellaceae, Trebouxiophyceae) taxa, including three new species from Antarctica

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Three new species of the genus *Micractinium* were collected from five localities on the South Shetland Islands in maritime Antarctica, and their morphological and molecular characteristics were investigated. The vegetative cells are spherical to ellipsoidal and a single chloroplast is parietal with a pyrenoid. Because of their simple morphology, no conspicuous morphological characters of new species were recognized under light microscopy. However, molecular phylogenetic relationships were inferred from the concatenated small subunit rDNA, and internal transcribed spacer (ITS) sequence data indicated that the Antarctic microalgal strains are strongly allied to the well-supported genus *Micractinium*, including *M. pusillum*, the type species of the genus, and three other species in the genus. The secondary structure of ITS2 and compensatory base changes were used to identify and describe six Antarctic *Micractinium* strains. Based on their morphological and molecular characteristics, we characterized three new species of *Micractinium*: *M. simplicissimum* sp. nov., *M. singularis* sp. nov., and *M. variabile* sp. nov.

**Key Words:** Antarctica; ITS2 secondary structure; *Micractinium*; morphology; nuclear rDNA; phylogenetic relationships

**INTRODUCTION**

Antarctica is an important location for exploring living organisms adapted to its harsh environment with relatively low anthropogenic impact. However, limited accessibility to Antarctica makes it difficult to collect samples and data. Nevertheless, several studies have examined Antarctic terrestrial life (Wall 2005, Convey et al. 2011, Dennis et al. 2019). The South Shetland Islands have diverse environments and valuable biological resources and are located 120 km north of the Antarctic Peninsula. Although several studies examined the diversity, ecology, and distribution of their terrestrial microalgae, taxonomic information on Antarctic microalgae remains limited (Fermani et al. 2007, Llames and Vinocur 2007, Zidarova 2008, Hamsher et al. 2016).

Members of the genus *Micractinium* Fresenius (family Chlorellaceae) are characterized by spherical or oval cells mostly arranged in colonies (Krienitz et al. 2004). The shape, size, and arrangement of cells and bristles are considered species-specific characters. However, the type species of the genus, *Micractinium pusillum* growing in dense culture as solitary spherical cells without bristles, is surprisingly morphologically similar to *Chlorella vulgaris*. Its morphological traits return to the original form while exposed to grazing zooplankton (Luo et
This morphological similarity between the two genera causes difficulties when identifying taxa and defining them using morphology alone.

Recent phylogenetic studies of the secondary structure of internal transcribed spacer 2 (ITS2) have demonstrated characteristic differences among species with the typical morphology of Micractinium (Pröschold et al. 2010, 2011, Hoshina and Fujiwara 2013). The secondary structure of ITS2 differentiates species of Micractinium and Chlorella (Luo et al. 2010). Compensatory base changes (CBCs) and hemi-CBCs in the secondary structure of ITS2 can be useful for distinguishing individual species (Krienitz et al. 2004, 2010, Müller et al. 2007).

Because of their morphological simplicity and phenotypic plasticity, some algal collections tend to keep unclassified Antarctic coccoid green microalgal strains at species level. We investigated morphological and molecular characteristics of six Antarctic microalgal strains from the Korea Polar Research Institute (KOPRI) Culture Collections for Polar Microorganisms (KCCPM) in order to define morphologically ambiguous specimens at the species level. We have characterized three new species.

**MATERIALS AND METHODS**

**Collections**

Freshwater samples were collected from five sites on the South Shetland Islands, Antarctica in January 2014 (Table 1). Strains were isolated using sterile Pasteur pipettes. The cells were rinsed repeatedly and grown in artificial freshwater solution B5282 (Sigma-Aldrich, Saint Louis, MO, USA). They were cultivated at approximately 2°C under 25 μmol photons m⁻² s⁻¹ (16 : 8 h light : dark cycle) in a culture room at KCCPM.

| Species                          | Strain     | Collection information                                      | Accession No. |
|---------------------------------|------------|-------------------------------------------------------------|---------------|
| *Micractinium simplicissimum* sp. nov. | KSF0100    | Nelson Island, South Shetland Islands, Antarctica (62°18’17.53” S, 59°11’55.77” W; Jan 19, 2014) | MN414472      |
|                                 | KSF0112    | King George Island, South Shetland Islands, Antarctica (62°14’24.49” S, 58°43’29.18” W; Jan 11, 2014) | MN414470      |
|                                 | KSF0114    | Deception Island, South Shetland Islands, Antarctica (62°58’36.50” S, 60°40’31.90” W; Jan 24, 2014) | MN414467      |
|                                 | KSF0127    | Robert Island, South Shetland Islands, Antarctica (62°23’02.80” S, 59°41’31.50” W; Jan 28, 2014) | MN414471      |
| *Micractinium singularis* sp. nov.     | KSF0094    | Deception Island, South Shetland Islands, Antarctica (62°58’36.50” S, 60°40’31.90” W; Jan 24, 2014) | MN414469      |
| *Micractinium variabile* sp. nov.     | KSF0085    | Ardley Island, South Shetland Islands, Antarctica (62°12’39.84” S, 58°55’19.68” W; Jan 16, 2014) | MN414468      |

**Microscopy**

Morphological identification was performed using an Axio Imager A2 (Carl Zeiss Inc., Hallbergmoos, Germany) equipped with differential interference contrast optics. Photomicrographs were taken with an AxioCam HRC camera (Carl Zeiss Inc.).

**DNA extraction, amplification, and sequencing**

DNA was extracted using the i-genomic Plant DNA Extraction Kit (iNtRON Biotechnology, Seongnam, Korea) according to the manufacturer’s instructions. We performed polymerase chain reaction (PCR) to amplify small subunit (SSU) rDNA, internal transcribed spacer 1 (ITS1), 5.8S rDNA, and ITS2 with the primer pairs NS1/NS4, NS5/NS8, and ITS1/ITS4 (White et al. 1990). The PCR products were purified with the MG PCR Product Purification Kit (Macrogen, Seoul, Korea) following the manufacturer’s recommendations. The DNA was sequenced by Macrogen.

**Phylogenetic analyses**

Phylogenetic analyses were performed on a concatenated dataset of the SSU, ITS1, 5.8S, and ITS2 rDNA sequences. The aligned sequences were checked for possible misaligned positions in BioEdit 7.0.5.3 (Hall 1999). The alignment was constructed by taking sequences of Micractinium species from previous studies (Krienitz et al. 2010, Luo et al. 2010, Hoshina and Fujiwara 2013) and comparing them to the newly obtained ones.

Appropriate evolutionary model was assessed using jModelTest 2 (Darriba et al. 2012) under the Akaike Information Criterion. Selected model has the following parameters: GTR + I + G with gamma distribution = 0.5220;
**RESULTS**

**Morphological observations by light microscopy**

The Antarctic strains KSF0085, KSF0094, KSF0100, KSF0112, KSF0114, and KSF0127 possessed the typical morphological characteristics of *Micractinium* species in terms of cell shape, chloroplasts, and pyrenoids (Table 2). However, bristles and colony formation were observed only for KSF0085. The vegetative cells of KSF0085 were spherical or ellipsoidal, and 5.0-5.4 × 8.2-8.6 μm in diameter with or without bristles in culture conditions (Fig. 1A & B). The cells contained a cup- to girdle-shaped chloroplast with a spherical to ellipsoidal pyrenoid. The cells of KSF0094 were spherical or ellipsoidal, and 4.5-4.7 × 7.2-7.4 μm in diameter (Fig. 1C). A single cup- or girdle-shaped chloroplast with a pyrenoid was present. KSF0112 was spherical to ellipsoidal, 3.3-3.9 × 5.5-5.7 μm, and contained cup- or girdle-shaped chloroplasts with a pronounced pyrenoid (Fig. 1D). KSF0100, KSF0114, and KSF0127 were morphologically similar to KSF0112.

**ITS2 secondary structure**

To locate hemi-CBCs and CBCs, the ITS2 secondary structures were constructed using Mfold (Zuker 2003) and 4SALE (Seibel et al. 2006, 2008).
Fig. 2. A maximum likelihood (ML) tree constructed from small subunit, internal transcribed spacer 1, 5.8S, and internal transcribed spacer 2 ribosomal DNA sequences. Numbers at each node are the ML (>50%, left) and Bayesian values (>0.95, right). Antarctic strains are in bold, and the type species of the genus Micractinium is highlighted with a star.

Table 2. A comparison of Micractinium simplicissimum, M. singularis, and M. variabile with other species of Micractinium

| Species                   | Cell shape                        | Cell size (µm) | Form colony | Bristle size (µm) | Reference                  |
|---------------------------|-----------------------------------|----------------|-------------|-------------------|----------------------------|
| M. simplicissimum sp. nov.| Spherical to ellipsoidal          | 3.3-3.9 × 5.5-5.7 | -           | NA                | Present study              |
| M. singularis sp. nov.    | Spherical to ellipsoidal          | 4.5-4.7 × 7.2-7.4 | -           | NA                | Present study              |
| M. variabile sp. nov.     | Spherical to ellipsoidal          | 5.0-5.4 × 8.2-8.6 | +, -        | 10-30             |                            |
| M. pusillum               | Spherical                        | (3-) 7-13       | +, -        | 20-35             | Kim and Kim (2012)         |
| M. appendiculatum         | Oval, ovoid                       | 8              | +           | 70                | Korshikov (1953)           |
| M. belenophorum           | Spherical to ellipsoid            | (3.3-) 4.5-5.5 × (5-) 8-10 | -   | 30 (-55)         | Pröschold et al. (2010)    |
| M. bornhemiense           | Spherical                        | 6-8            | +           | 30-60 (-90)       | Korshikov (1953)           |
| M. conductrix             | Spherical to slightly ellipsoid   | 7-12           | -           | NA                | Pröschold et al. (2011)    |
| M. crassisetum            | Globose                           | 5.5-6.5        | +           | 24.7-33.8         | Horthogáyi (1973)          |
| M. extremum               | Spherical                        | 5.2-6.4        | +           | 30-50             | Horthogáyi (1979)          |
| M. inermum                | Spherical to ellipsoidal          | 3.2-3.7 × 5.0-5.4 | -          | NA                | Hoshina and Fujisawa (2013)|
| M. quadrisetum            | Spherical to obovoid              | 3.5-7 × 5-10   | +           | 20-50             | Kim and Kim (2012)         |

NA, not available.
Sequencing results and phylogenetic analyses

SSU sequence-based approach of the members of Chlorellaceae, did not provide enough information to distinguish taxa from public database as well as from this study at species level. Further analyses were performed using the combined SSU and ITS rDNA dataset. The genus *Micractinium* was highly supported in the ML and BI analyses (Fig. 2). The tree revealed 10 distinct lineages within the genus, including four lineages known as *Micractinium* species: *M. inermum*, *M. conductrix*, *M. pusillum*, and *M. belenophorum*. The six Antarctic strains (KSF0085, KSF0094, KSF0100, KSF0112, KSF0114, and KSF0127) clustered with non-bristle forming *M. inermum* in both analyses. The six strains were subdivided into two clades; one clade contained four strains (KSF0100, KSF0112, KSF0114, and KSF0127), and the other two strains formed two subclades. The strains (KSF0100, KSF0112, KSF0114, and KSF0127) formed first clade were collected from four sites of the South Shetland Islands and were identical to *Micractinium* sp. (KNUA029) (Hong et al. 2015). KSF0085 belonged to the same clade as *Micractinium* sp. (KNUA034), which is an Antarctic strain without bristles (Hong et al. 2015). KSF0085 was sister to KSF0094 with strong to moderate support in the analyses.

**ITS2 secondary structure**

Fig. 3 shows the secondary structures of ITS2 of the Antarctic strains and other species of *Micractinium*. All structures possess motifs conserved within the green algae (Mai and Coleman 1997), including a four-fingered hand (helices I-IV), pyrimidine–pyrimidine bulge on helix II, and a conserved sequence TGGT (UGGU) on the 5′-side of helix III. *Micractinium* species have C-G pairing at the tip of helix III (Luo et al. 2006, 2010). We found distinct characteristics in the ITS2 secondary structures of the six Antarctic strains, with three types of ITS2 secondary structure (Fig. 3). These structures and base changes differed from the closely related species *M. inermum*. Compared to those of *M. inermum*, four hemi-CBC sites of ITS2 secondary structures were revealed in KSF0085 and KSF0094, and two CBC sites were revealed in KSF0012. KSF0085 and KSF0094 showed differences in helix I structure, and no CBC or hemi-CBC was found in these strains. KSF0012 showed a base change from G:U to U:A in helix IV and three hemi-CBCs in helices II and III, compared to KSF0085 and KSF0094.

**DISCUSSION**

The family Chlorellaceae, containing small, round, non-flagellate microalgae, has been divided into several groups using various characteristics, such as colony formation, bristle, mucilage, and pyrenoid (Krienitz et al. 2004). They recognized *Micractinium* has spherical to ellipsoidal cells with bristle formation in a colony. However, not all *Micractinium* species are capable of bristle producing (Luo et al. 2010, Hoshina and Fujiwara 2013), and the differentiation of *Micractinium* species based on the length and number of bristles is unreliable (Luo et al. 2006). Luo et al. (2005) observed single-celled and bristle-free forms when *M. pusillum*, the type species of the genus, was cultivated without zooplankton. However, culturing *Micractinium* strain with zooplankton does not always produce bristles and colonies (Pröschold et al. 2011). *M. conductrix*, *M. inermum*, and other Antarctic strains (KNUA029 and KNUA034) of the genus are non-bristle and similar to *Chlorella* morphologically (Pröschold et al. 2011, Hoshina and Fujiwara 2013, Hong et al. 2015). Species delimitation in *Micractinium* has always been problematic because there are few diagnostic morphological characters and they are not easily observed. Just one out of six Antarctic strains, KSF0085 produced various colony types and bristle formations when observed in the light microscope, whereas the other strains did not show these typical characteristics of *Micractinium* (Fig. 1).

Phylogenetic analyses of the SSU and ITS sequences of the nuclear rDNA gene revealed that the Antarctic *Micractinium* strains were clearly separated from the genus *Chlorella* and these strains were closely related to *M. inermum* (Fig. 2). Hong et al. (2015) reported that KNUA029 and KNUA034 were collected from two sites in West Antarctica. Morphological and molecular characteristics of KNUA029 and KNUA034 are identical with four Antarctic strains (KSF0100, KSF0112, KSF0114, and KSF0127) and those of KNUA034 are same with strain KSF0085.

Recent molecular studies have demonstrated the usefulness of ITS2 sequences for defining species of green coccolid algae. The ITS2 region is highly conserved within species (Müller et al. 2005), whereas it is highly divergent between species (Hoshina et al. 2010). This strongly encourages a species concept based on ITS2 sequence differences (Mai and Coleman 1997). In addition, many taxonomic studies have used CBC in the ITS2 secondary structure for species delimitation (Krienitz et al. 2004, Müller et al. 2007, Hoshina and Fujiwara 2013). The distinctive base pairing of helix III of ITS2 clearly
Fig. 3. Predicted internal transcribed spacer 2 secondary structure diagrams for KSF0085, KSF0094, KSF0112, and other species of Micractinium. The synapomorphic signature of the genus Micractinium (Luo et al. 2010) is highlighted in black boxes. Single base changes are circled and the compensatory base changes are highlighted in boxes. The models are modified from Hoshina and Fujiwara (2013).
distinguishes *Micractinium* from other genera within the *Chlorella* clade (Luo et al. 2010). As our study showed, the six investigated strains belong to the genus *Micractinium* based on the C-G pairing CBC at the tip of ITS2 helix III (Fig. 3). The six Antarctic strains are considered to be three separate species, with differences in the secondary structure of ITS2, CBCs, and hemi-CBCs. KSF0085 and KSF0094 showed the greatest difference in helix I structure among the ITS2 secondary structures in the Antarctic strains. KSF0112 showed differences in one CBC and three hemi-CBCs with KSF0085 and KSF0094. The morphological and molecular characteristics of KSF0100, KSF0112, KSF0114, and KSF0127 were identical.

There are members of the genus *Micractinium* that are distributed worldwide, but no species has been reported yet in Antarctica (Guiry and Guiry 2019). The Antarctic strains show tolerance to lower temperature and produced polyunsaturated fatty acids in cold condition (Hong et al. 2015). It suggests an evolutionary evidence of their adaptation to the Antarctic environment. The six Antarctic strains are similar to other *Micractinium* species morphologically (Table 2). However, these Antarctic strains are clearly discriminated from other *Micractinium* species by molecular characteristics such as their phylogenetic relationships and ITS2 secondary structures, therefore we propose them as three new species.

**Taxonomic conclusion**

*Micractinium simplicissimum* H. Chae, H. -G. Choi & J. H. Kim sp. nov. (Fig. 1D)

Cells solitary, planktonic, spherical to ellipsoidal shaped, 3.3-3.9 × 5.5-5.7 μm, without bristles. Mucilage absent. Chloroplast single, parietal, cup-, girdle-shaped, with a broadly ellipsoidal to spherical pyrenoid. Reproduction by autospores, zoospores not observed. Differs from other species of this genus by the order of nucleotides in ITS2 and the barcoding signatures.

**Holotype.** KSF0112 collected on Jan 11, 2014 in King George Island (South Shetland Islands, Antarctica; 62°14′24.49″ S, 58°43′29.18″ W) by S. Lim and stored at KCCPM, Incheon, Korea.

**Type locality.** King George Island, South Shetland Islands, Antarctica.

**Etymology.** Referring to the solitary habit.

**Iconotype.** Fig. 1D.

**Distribution.** King George Island, South Shetland Islands, Antarctica.

**GenBank accession number.** MN414469.

*Micractinium variabile* H. Chae, H. -G. Choi & J. H. Kim sp. nov. (Fig. 1A & B)

Cells solitary or colonial, planktonic, spherical to ellipsoidal shaped, 5.0-5.4 × 8.2-8.6 μm, with or without bristles. Mucilage absent. Chloroplast single, parietal, cup-, girdle-shaped, with a broadly ellipsoidal to spherical pyrenoid. Reproduction by autospores, zoospores not observed. Differs from other species of this genus by the order of nucleotides in ITS2 and the barcoding signatures.

**Holotype.** KSF0085 collected on Jan 16, 2014 in Ardley Island (South Shetland Islands, Antarctica; 62°12′39.84″ S, 58°55′19.68″ W) by S. Lim and stored at KCCPM, Incheon, Korea.

**Type locality.** Ardley Island, South Shetland Islands, Antarctica.

**Etymology.** Referring to the changeable morphology with bristles or without them.

**Iconotype.** Fig. 1A & B.

**Distribution.** Ardley Island, South Shetland Islands, Antarctica.

**GenBank accession number.** MN414468.

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REFERENCES

Convey, P., Hopkins, D. W., Roberts, S. J. & Tyler, A. N. 2011. Global southern limit of flowering plants and moss peat accumulation. Polar Res. 30:8929.

Darriba, D., Taboada, G. L., Doallo, R. & Posada, D. 2012. jModelTest 2: more models, new heuristics and parallel computing. Nat. Methods 9:772.

Dennis, P. G., Newsham, K. K., Rushton, S. P., O’Donnell, A. G. & Hopkins, D. W. 2019. Soil bacterial diversity is positively associated with air temperature in the maritime Antarctic. Sci. Rep. 9:2686.

Ferrmani, P., Mataloni, G. & Van de Vijver, B. 2007. Soil micro-algal communities on an antarctic active volcano (Deception Island, South Shetlands). Polar Biol. 30:1381-1393.

Guindon, S., Dufayard, J. F., Lefort, V., Anisimova, M., Hordijk, W. & Gascuel, O. 2010. New algorithms and methods to estimate maximum-likelihood phylogenies: assessing the performance of PhyML 3.0. Syst. Biol. 59:307-321.

Guiry, M. D. & Guiry, G. M. 2019. AlgaeBase. World-wide electronic publication, National University of Ireland, Galway. Available from: http://www.algaebase.org. Accessed Sep 17, 2019.

Hall, T. A. 1999. BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. Nucleic Acids Symp. Ser. 41:95-98.

Hamsher, S., Kopalova, K., Kociolek, J. P., Zidarova, R. & Van de Vijver, B. 2016. The genus Nitzschia on the South Shetland Islands and James Ross Island. Fottea 16:79-102.

Hong, J. W., Jo, S. -W., Cho, H. -W., Nam, S. W., Shin, W., Park, K. M., Lee, K. I. & Yoon, H. -S. 2015. Phylogeny, morphology, and physiology of Micractinium strains isolated from shallow ephemeral freshwater in Antarctica. Phycol. Res. 63:212-218.

Hortobágyi, T. 1973. Neue Chlorococcales aus den Absetz- und Grundwasseranreicherungsbecken der Budapester Wasserwerke. Acta Bot. Acad. Sci. Hung. 18:119-130.

Hortobágyi, T. 1979. New Chlorococcales species in the Danube hybrid algae? Acta Biol. Szeged 25:7-12.

Hoshina, R. & Fujiwara, Y. 2013. Molecular characterization of Chlorella cultures of the National Institute for Environmental Studies culture collection with description of Micractinium inermum sp. nov., Didymogenes sphaerica sp. nov., and Didymogenes soliella sp. nov. (Chlorococcales, Trebouxiophyceae). Phycol. Res. 61:124-132.

Hoshina, R., Iwataki, M. & Imamura, N. 2010. Chlorocella variabilis and Micractinium reisseri sp. nov. (Chlorococcales, Trebouxiophyceae): redescriptions of the endosymbiotic green algae of Paramecium bursaria (Peniculina, Oligohymenophorea) in the 120th year. Phycol. Res. 58:188-201.

Kim, Y. G. & Kim, H. S. 2012. Algal flora of Korea. Vol. 6. No. 2. Freshwater green algae: Chlorophyta: Chlorococcales I: Micractiniaceae, Botryococcaceae, Characiaceae, Hydrodictyaceae. National Institute of Biological Resources, Incheon, 117 pp.

Korshikov, A. A. 1953. The freshwater algae of the Ukrainian SSR. Vol. 5. Sub-class Protococccineae. Vacuolales and Protococcales. Akademii Nauk URSR, Kiev, pp. 1-439.

Krienitz, L., Bock, C., Luo, W. & Pröschold, T. 2010. Polyphyletic origin of the Dictyosphaerium morphotype within Chlorococcales (Trebouxiophyceae). J. Phycol. 46:559-563.

Krienitz, L., Hegewald, E. H., Hepperle, D., Huss, V. A. R., Rohr, T. & Wolf, M. 2004. Phylogenetic relationship of Chlorella and Parachlorella gen. nov. (Chlorophyta, Trebouxiophyceae). Phycologia 43:529-542.

Llamas, M. E. & Vinocur, A. 2007. Phytoplankton structure and dynamics in a volcanic lake in Deception Island (South Shetland Islands, Antarctica). Polar Biol. 30:849-857.

Luo, W., Krienitz, L., Pflugmacher, S. & Walz, N. 2005. Genus and species concept in Chlorella and Micractinium (Chlorophyta, Chlorococcales): genotype versus phenotypic variability under ecosystem conditions. Verh. Int. Ver. Limnol. 29:170-173.

Luo, W., Pflugmacher, S., Pröschold, T., Walz, N. & Krienitz, L. 2006. Genotype versus phenotype variability in Chlorella and Micractinium (Chlorophyta, Trebouxiophyceae). Protilist 157:315-333.

Luo, W., Pröschold, T., Bock, C. & Krienitz, L. 2010. Generic concept in Chlorella-related coccoid green algae (Chlorophyta, Trebouxiophyceae). Plant Biol. 12:545-553.

Mai, J. C. & Coleman, A. W. 1997. The internal transcribed spacer 2 exhibits a common secondary structure in green algae and flowering plants. J. Mol. Evol. 44:258-271.

Müller, J., Friedl, T., Hepperle, D., Lorenz, M. & Day, J. G. 2005. Distinction between multiple isolates of Chlorella vulgaris (Chlorophyta, Trebouxiophyceae) and testing for conspecificity using amplified fragment length polymorphism and ITS rDNA sequences. J. Phycol. 41:1236-1241.

Hoshina, R., Iwataki, M. & Imamura, N. 2010. Chlorocella variabilis and Micractinium reisseri sp. nov. (Chlorococcales, Trebouxiophyceae): redescriptions of the endosymbiotic golden algae of Paramecium bursaria (Peniculina, Oligohymenophorea) in the 120th year. Phycol. Res. 58:188-201.
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1247.
Müller, T., Philippi, N., Dandekar, T., Schultz, J. & Wolf, M. 2007. Distinguishing species. RNA 13:1469-1472.
Pröschold, T., Bock, C., Luo, W. & Krienitz, L. 2010. Polymorphic distribution of bristle formation in Chlorellaceae: *Micractinium*, *Diacanthos*, *Didymogenes* and *Hegewaldia* gen. nov. (Treouxiophyceae, Chlorophyta). Phycologia 58:1-8.
Pröschold, T., Darienko, T., Silva, P. C., Reisser, W. & Krienitz, L. 2011. The systematics of *Zoochlorella* revisited employing an integrative approach. Environ. Microbiol. 13:350-364.
Ronquist, F., Teslenko, M., van der Mark, P., Ayres, D. L., Darling, A., Höhna, S., Larget, B., Liu, L., Suchard, M. A. & Huelsenbeck, J. P. 2012. MrBayes 3.2: efficient Bayesian phylogenetic inference and model choice across a large model space. Syst. Biol. 61:539-542.
Seibel, P. N., Müller, T., Dandekar, T., Schultz, J. & Wolf, M. 2006. 4SALE: a tool for synchronous RNA sequence and secondary structure alignment and editing. BMC Bioinformatics 7:498.
Seibel, P. N., Müller, T., Dandekar, T. & Wolf, M. 2008. Synchronous visual analysis and editing of RNA sequence and secondary structure alignments using 4SALE. BMC Res. Notes 1:91.
Wall, D. H. 2005. Biodiversity and ecosystem functioning in terrestrial habitats of Antarctica. Antarct. Sci. 17:523-531.
White, T. J., Bruns, T., Lee, S. & Taylor, J. 1990. Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. In Innis, M. A., Gelfand, D. H., Sninsky, J. J. & White, T. J. (Eds.) *PCR Protocols: A Guide to Methods and Applications*. Academic Press, San Diego, CA, pp. 315-322.
Zidarova, R. P. 2008. Algae from Livingston Island (S Shetland Islands): a checklist. Phytol. Balcan 14:19-35.
Zuker, M. 2003. MFold web server for nucleic acid folding and hybridization prediction. Nucleic Acids Res. 32:3406-3415.