Dictyostelid Cellular Slime Molds from Christmas Island, Indian Ocean

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ABSTRACT Christmas Island (10°30’S, 105°40’E) is an Australian external territory located in the Indian Ocean, approximately 350 km south of Java and Sumatra and about 1,550 km northwest of the closest point on the Australian mainland. In May 2017, 20 samples of soil/humus were collected on Christmas Island and processed for dictyostelid cellular slime molds. Four species were recovered. Two of these (Dictyostelium purpureum and Cavenderia aureostipes) are common and widely distributed throughout the world, but two other species (Dictyostelium insulativaitis sp. nov. and Dictyostelium barbarae sp. nov.) were found to be new to science and are described here.

IMPORTANCE Reported here are the results of a study for dictyostelids carried out on Christmas Island, Indian Ocean. Six isolates representing four species of dictyostelid cellular slime molds were obtained from two of the four localities from which samples were collected on the island. Two of the species (Dictyostelium insulativaitis and D. barbarae) belong to the Dictyosteliaceae, genus Dictyostelium, and are new to science. These are described based on both morphology and phylogeny. The diversity and abundance of dictyostelids on Christmas Island appear to be low, which might in part be due to the abundance of land crabs, which considerably reduce the extent of the litter layer on the forest floor.

KEYWORDS Amoebozoa, Cavenderia, Dictyostelium, phylogeny, taxonomy

Dictyostelid cellular slime molds (dictyostelids) are single-celled, eukaryotic, phagotrophic bacteriovores usually present and sometimes abundant in terrestrial ecosystems. These organisms represent a normal component of the microbiota in soils and apparently play a role in maintaining the natural balance that exists between bacteria and other microorganisms in the soil environment. The primary microhabitat for dictyostelids is the leaf litter decomposition zone of forest soils, but they also occur in other types of soils (1).

Dictyostelids have been reported from numerous localities throughout the world, but there are relatively few reports of the group from small isolated islands (2, 3). During the course of a survey for myxomycetes carried out on Christmas Island (S. L. Stephenson and B. C. Stephenson, submitted for publication) in May 2017, 20 samples of soil/humus for isolation of dictyostelids were collected by the senior author and sent to the first author for processing. Four species of dictyostelids were recovered from these samples, and two of these are species new to science. The primary objective of this paper is to describe these two new species.

Christmas Island (10°30’S, 105°40’E) (Fig. 1A) is an Australian external territory located in the Indian Ocean, approximately 350 km south of Java and Sumatra and about 1,550 km northwest of the closest point on the Australian mainland (Fig. 1B). It has a total area of 135 km², and the highest point on the island is 361 m. Most of the island consists of a gently undulating plateau that represents the summit of an

Citation Liu P, Zou Y, Li W, Li Y, Li X, Che S, Stephenson SL. 2019. Dictyostelid cellular slime molds from Christmas Island, Indian Ocean. mSphere 4:e00133-19. https://doi.org/10.1128/mSphere.00133-19.

Editor Aaron P. Mitchell, Carnegie Mellon University

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Received 24 February 2019
Accepted 19 March 2019
Published 10 April 2019
underwater mountain more than 4,500 m high. Christmas Island receives an average of 214 cm of rainfall each year, with the majority falling between November and June. Very little surface water exists on the island. Temperatures vary little throughout the year. The highest temperature is usually around 29°C in March and April, whereas the lowest temperature is about 23°C and occurs in August.

Much of the island is covered by tropical forests, with two types which are strongly correlated with soil depth. The predominant type is an evergreen tall forest that is found in areas characterized by deep soils. This includes most of the central plateau of the island (Fig. 1C). A semideciduous closed forest occurs in areas with shallow soils. A total of 213 species of native plants are known from Christmas Island, 17 of which are endemic to the island. However, approximately 250 species of exotic plants also have been introduced to the island (4). One of the more prominent of these is the coconut palm (*Cocos nucifera* L.).

The most conspicuous animals in the forests on Christmas Island are indigenous large land crabs (*Gecarcoidea natalis* Pocock), which are often exceedingly abundant. Densities of >50 crabs per 100 m² are not uncommon. The crabs (Fig. 1D) are omnivorous scavengers and feed upon fallen leaves, fruits, seeds, various types of plant debris, tree seedlings, and dead animals. Their presence on the forest floor has a

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**FIG 1** Christmas Island. (A) Map showing the collecting localities on the island. (B) Location of the island with respect to mainland Australia. (C) Typical tropical forest on the central plateau of the island. (D) Example of the large land crabs which are often exceedingly abundant in the forests on the island. (Maps courtesy of Carlos Rojas, reproduced with permission.)
considerable ecological impact by greatly reducing the extent of the litter layer and limiting the establishment of tree seedlings and other plants.

RESULTS

Six isolates representing four species of dictyostelids were recovered from samples collected at four localities on Christmas Island (Fig. 1A). Two isolates (Dictyostelium insulinativitatis and Dictyostelium barbarae) (Fig. 2 and 3) from one of the localities (L1) are new to science. Four isolates from a second locality (L2) were Dictyostelium purpureum Olive (isolate 2-1) and Cavenderia aureostipes (Cavender) S. Baldauf, S. Sheikh & Thulin [isolates 2-2, 2-3(1), and 2-3(2)] (Fig. 4). Two other localities (L3 and L4) did not yield any isolates. Phylogenetic studies of the ribosomal small subunit (SSU) further support the taxonomic placement of all four of the species recorded from Christmas Island (Fig. 5). Isolates of all of these species were deposited in the mycological herbarium of Jilin Agricultural University.

Taxonomy and molecular phylogeny. Dictyostelium insulinativitatis S. L. Stephen- son, P. Liu, Y. Li et Y. Zou, sp. nov. (Fig. 2). MycoBank accession number MB829168. GenBank accession number MK322958. When cultured at 23°C on nonnutrient agar
with *Escherichia coli*, sorocarps white, usually erect, gregarious, unbranched, commonly 0.45 to 1.26 mm high. Sorophore white, tips obtuse with one or two tiers of cells (6.99 to 9.03 μm in diameter), bases clavate with one or two tiers of cells (10.89 to 24.70 μm in diameter). Sori white, globose, commonly 0.11 to 0.15 mm in diameter. Spores hyaline, oblong to oval, 6.23 to 10.74 by 4.70 to 6.71 μm or 5.96 to 8.31 by 4.73 to 6.80 μm, without obvious polar granules. Aggregations mound-like, without radiate streams. Pseudoplasmodia migrate with stalk formation, producing clustered sorogens.

(i) Etymology. Referring to the locality (Christmas Island) where this species was isolated.

**FIG 3** Morphological features of *Dictyostelium barbarae*. (A and B) Sorocarps. (C) Aggregations. (D) Pseudoplasmodia. (E to G) Sorophore tips. (H and I) Sorophore bases. (J) Spores. Bars: A to D, 1 mm; E to J, 20 μm.
(ii) Holotype. HMJAU MR300 (strain 1-2) isolated from a soil/humus sample collected in an evergreen tall forest (L1, 10°28’24”S, 105°35’28”E), West White Beach Walking Track, off North West Point Road, 17 May 2017, Christmas Island, Indian Ocean.

(iii) Comments. This species belongs to dictyostelid group 4 (5) in an SSU ribosomal DNA (rDNA) phylogeny (Fig. 5). It forms a clade together with *D. barbarae* and *Dictyostelium macrocephalum* H. Hagiw. (6, 7). However, it differs morphologically from *D. barbarae* in the width of the sorophore. The sorophore of *D. barbarae* is composed of several tiers of cells, whereas that of *D. insulinativitatis* is made up of only one or two tiers of cells. The sorophore tip of *D. barbarae* is clavate or obtuse and wider than in *D. insulinativitatis*. Moreover, the base of the sorophore for *D. barbarae* is composed of several tiers of cells as opposed to only one or two tiers of cells in *D. insulinativitatis*. The spore size for *D. insulinativitatis* is larger than that of *D. barbarae*. The ratio of spore length and width in *D. barbarae* (1.5 to 1.63 or 1.05 to 1.38) is larger than in *D. insulinativitatis* (1.32 to 1.59 or 0.96 to 1.26). The habits of the sorocarps are clustered in *D. barbarae* and gregarious in *D. insulinativitatis*. The aggregation type of *D. insulinativitatis* is mound-like, but the aggregation type of *D. barbarae* is radiate with streams.

In addition, the sorocarps of *D. macrocephalum* (0.25 to 2.25 [−9.0]) are larger than this new species. The base of the sorophore in *D. macrocephalum* is conical, whereas
FIG 5  Phylogeny of four species obtained in this study along with other species of dictyostelids based on SSU rRNA, indicating the phylogenetic position of the new species Dictyostelium insulinativitatis and D. barbara. Names with an asterisk are the sequences obtained in this study.
the base of the sorophore in *D. insulinativitatis* is clavate. The ratio of spore length and width in *D. macrocephalum* (1.57 to 1.98) is larger than in the latter species (1.32 to 1.59 or 0.96 to 1.26). The aggregation of *D. macrocephalum* is radiate with streams, but aggregation in *D. insulinativitatis* is the mound-like type.

**Dictyostelium barbarae** S. L. Stephenson, P. Liu, Y. Li et Y. Zou, sp. nov. (Fig. 3). MycoBank accession number MB829169. GenBank accession number MK322959.

When cultured at 23°C on nonnutrient agar with *E. coli*, sorocarps white, usually erect, clustered, unbranched, commonly 0.22 to 1.59 mm high. Sorophore white with several tiers of cells, tips clavate or obtuse with one or two tiers of cells (9.87 to 14.83 μm in diameter), bases clavate or round with several tiers of cells (13.98 to 31.29 μm in diameter). Sori white, globose, commonly 0.120 to 0.202 mm in diameter. Spores hyaline, oblong or oval, 5.83 to 8.34 by 3.90 to 5.11 μm or 4.50 to 5.46 by 3.99 to 5.25 μm, without obvious polar granules. Aggregations radiate with streams. Pseudoplasmodia migrate with stalk formation.

(i) **Etymology.** Named in honor of the wife (Barbara Stephenson) of one of the coauthors of this paper.

(ii) **Holotype.** HMJAU MR301 (strain 1-5), isolated from a soil/humus sample collected in an evergreen tall forest (L1, 10°28′24″S, 105°35′28″E), West White Beach Walking Track, off North West Point Road, 17 May 2017, Christmas Island, Indian Ocean.

(iii) **Comments.** Molecular data from the ribosomal small subunit (SSU) support the placement of this species in the genus *Dictyostelium* (5) (Fig. 5). This species forms a clade with *D. insulinativitatis* and *D. macrocephalum*. The differences between *D. barbarae* and *D. insulinativitatis* were discussed in the comments provided for *D. insulinativitatis*. The sorocarps of *D. macrocephalum* (0.25 to 2.25 μm in diameter) are larger than in *D. barbarae*. The base of the sorophore in *D. barbarae* is clavate or round, whereas it is conical in *D. macrocephalum*. The ratio of spore length and width in *D. macrocephalum* (1.57 to 1.98) is larger than that in *D. barbarae* (1.5 to 1.63 or 1.05 to 1.38).

**DISCUSSION**

Although long-distance dispersal by wind undoubtedly accounts for the presence of many of the microorganisms found on an isolated land mass such as Christmas Island, this is not necessarily true for all groups. For example, the spores produced by dictyostelids are embedded in a mucilaginous matrix that dries and hardens. As such, the spores stand little chance of being dispersed by wind (3, 8). However, as noted in the introductory section of this paper, numerous exotic plants have been introduced to Christmas Island, and it is certainly possible that some of these could have served as vectors for dictyostelids. Two of the four species (*Dictyostelium purpureum* and Cavendia aureostipes) recovered in the present study are widely distributed throughout the world. The presence of the two species new to science is more problematic. Their occurrence may or may not be restricted to Christmas Island, since it is possible they can be found elsewhere but have not yet been reported. Conversely, these two species could be restricted to the island and thus provide evidence of the speciation process in dictyostelids. At this point, there is no way of knowing just which of the two situations exists.

In general, the overall diversity and abundance of dictyostelids on Christmas Island appear to be low. As noted above, two (L3 and L4) of the four localities yielded no isolates, and a total of only six clones were recovered from all 20 samples. For mainland Australia, the closest landmass for which comparable data exist, extensive sampling over a number of years (>300 samples from localities throughout the country) yielded an average of approximately 2.5 clones per sample (J. C. Landolt, unpublished data). The land crabs which are rather abundant in the forests on Christmas Island have a considerable ecological impact by reducing the extent of the litter layer. In some areas, very little litter exists. Since the soil-litter interface zone in forests represents the primary microhabitat for dictyostelids, any reduction in the amount of litter present on the forest floor might be expected to reduce the extent of this microhabitat and by extension the number of dictyostelids present. Interestingly, the most productive set of
samples was obtained from the second locality (L2), a forest dominated by coconut palms. As noted earlier, coconut palm is an exotic plant that was introduced to Christmas Island.

For those species of dictyostelids present on Christmas Island, it seems likely that the land crabs have the potential of serving as vectors of their spores, since there is little question that as they move about, these animals invariably come into direct contact with the soil/litter microhabitat on the forest floor. The role of invertebrates in the dispersal of spores for dictyostelids is well established (9, 10).

There are few data relating to dictyostelids on isolated islands, but the total number of species (four) recorded in the present study is higher than the totals (three and one, respectively) reported for Ascension Island in the mid-Atlantic Ocean and Macquarie Island in the Subantarctic. However, these differences are not appreciable enough to be significant. Clearly, the dictyostelids of other small isolated oceanic islands need to be investigated in more detail.

MATERIALS AND METHODS

Sampling, isolation, and cultivation. Twenty samples of soil/humus were collected from four localities on Christmas Island (Fig. 1A) in May 2017. Two of the localities (L1 and L4) were located in evergreen tall forests, one (L3) in a semideciduous closed forest, and one (L2) in a forest dominated by coconut palms. Each sample consisted of 10 to 20 g of soil/humus and was placed in a sterile Whirl-Pak plastic bag. The isolation methods used in the present study followed those described by Cavender and Raper (11) and are briefly summarized here. Each sample was weighed, and enough ddH2O water was added for an initial dilution of 1:10. This dilution was shaken to disperse the material and to suspend the amoebae, microcysts, and spores of dictyostelids. Afterward, an 0.5-ml aliquot of this dilution was added to each of five duplicate culture plates prepared with hay infusion agar (1). Approximately 0.4 ml of a heavy suspension of the bacterium E. coli was added to each culture plate as a food source. The plates were incubated at temperatures of 17 and 23°C with a 12-h light and dark cycle. Each plate was examined at least once per day for 2 weeks after the appearance of initial aggregations. Each isolate recovered from one of the plates was purified and cultivated for taxonomic studies and preservation on nonnutrient water agar plates with E. coli pregrown for 12 to 24 h. Spores from these plates were frozen in HL 5 medium (12) and stored at −80°C in the herbarium of the Mycological Institute of Jilin Agricultural University (HMJAU), Changchun, China.

Observations of morphological features. Six isolates were identified with the use of the descriptions provided by Raper (1) and molecular characteristics proposed by Sheikh et al. (5). First, the location of each early aggregating clone and sorocarp that developed in a plate was marked. The characteristic stages in the life cycle, including cell aggregation and the formation of pseudoplasmodia and sorocarps, were observed under a Zeiss dissecting microscope (Axio Zoom V16) with a 1.5× objective and a 10× ocular. Slides with sorocarps were prepared with water as the mounting medium. Features of spores, sorophores, and sorocarps were observed and measured on the slides by using a Zeiss light microscope (Axio Imager A2), with 10× ocular and 10, 40, and 100× (oil) objectives. Photographs were taken with a Zeiss AxioCam 506 color microscope camera.

DNA isolation, PCR amplification, and sequencing. The spores of all six isolates being studied were collected with a sterile tip and mixed with the lysis buffer of the MiniBEST universal genomic DNA extraction kit ver.5.0 (TaKaRa, Japan) according to the manufacturer's protocol. The genomic DNA solution was used directly for the SSU PCR amplification using the primers 18SF-A (AACCTGGTTGATCC TGCCAG) and 18SR-B (TGATCCTTGTGCGAGGTTCAC) (13) along with DS42F (ACAATTGGAGGGCAAGTCTG) and D1340R (TCCAGGTTCTGTCGCGTATC) (14). PCR products were sent to Sangon Biotech Co., Ltd. (Shanghai, China), for sequencing. Sequences obtained were deposited in the GenBank database.

Phylogenetic analysis. The six newly generated sequences were checked and then submitted to GenBank. The SSU sequences were aligned and compared using the program MUSCLE v.3.6 (15, 16) and then manually adjusted in MEGA 7.0 (17). Maximum likelihood (ML) analyses were performed using RAxML v7.18 (18). In the ML analyses, the best-fit substitution models were estimated using the GTR submission model and a gamma correction for rate variation among sites (GTR+Γ) using the CIPRES server. The statistical support of clades was assessed with 1,000 rapid-bootstrap (BS) replications.

Nomenclature. According to the International Code of Nomenclature used for algae, fungi, and plants, the electronic version of this article in portable document format (PDF) will represent a published work. In addition, new names contained in this study were submitted to MycoBank and allocated a unique MycoBank number which is accessible through MycoBank, Index Fungorum, GBIF, and other international biodiversity initiatives, where they are available to the Global Names Index.

Data availability. The isolates and the NCBI GenBank accession numbers of SSU DNA sequences considered in the present study are listed in Table 1. Sequence data are available in GenBank (accession numbers MK322958, MK322959, MK322960, MK322961, MK322962, and MK322963). The nomenclature of the new species in the present study is available in MycoBank (MB829168 and MB829169).
**TABLE 1** NCBI GenBank accession information for SSU sequences of all isolates included in the phylogenetic analysis

| Taxon                          | Isolate no. | Accession no. |
|-------------------------------|-------------|---------------|
| Acytostelium amazonicum       | Landolt X   | HQ141510.1    |
| Acytostelium anastomosans     | PP1         | AM168115.1    |
| Acytostelium digitatum        | OHS17       | AM168114.1    |
| Acytostelium leptosomum       | FG12        | AM168111.1    |
| Acytostelium magnisorum       | 08A         | HQ141513.1    |
| Acytostelium serpentinum      | SAB3A       | AM168113.1    |
| Acytostelium singularare      | FDI1        | HQ141514.1    |
| Acytostelium subglobosum      | LB1         | AM168110.1    |
| Cavenderia amphissora         | BM9A        | HQ141521.1    |
| Cavenderia aureostipes        | YA6         | AM168083.1    |
| Cavenderia aureostipes        | 2-2         | MK322961      |
| Cavenderia aureostipes        | 2-3(1)      | MK322962      |
| Cavenderia aureostipes        | 2-3(2)      | MK322963      |
| Cavenderia bifurcata          | UK5         | AM168084.1    |
| Cavenderia delicata           | TNS-C-226   | AM168093.1    |
| Cavenderia deminutiva         | MexM19A     | AM168092.1    |
| Cavenderia exigua              | TNS-C-199   | AM168085.1    |
| Cavenderia granulophora       | CHII-4      | AM168072.1    |
| Cavenderia macrocarpa         | MGE2        | HQ141519.1    |
| Cavenderia medusoides         | OH592       | AM168088.1    |
| Cavenderia mexicana           | MexTF4B1    | AM168089.1    |
| Cavenderia multistipes        | UK26b       | AM168070.1    |
| Cavenderia myxobasis          | NT2A        | HQ141522.1    |
| Cavenderia parvispora         | OS126       | AM168091.1    |
| Coremiosistem polystemhalum   | MY1-1       | AM168056.1    |
| Dicystostelium australindulium|             |              |
| Dicystostelium aureostephalum | TNS-C-180   | AM167876.1    |
| Dicystostelium aureum         | SL1         | AM168028.1    |
| Dicystostelium barbara        | 1-5         | MK322959      |
| Dicystostelium barbiliaus     | Sweden-4R   | JX173878.1    |
| Dicystostelium brefeldianum   | TNS-C-115   | AM168030.1    |
| Dicystostelium brunneum       | WS700       | AM168031.1    |
| Dicystostelium capitatum      | 91HO-50     | AM168032.1    |
| Dicystostelium chordate       |             | GQ496159.1    |
| Dicystostelium citrimum       | OH494       | AM168033.1    |
| Dicystostelium clavatum       | TNS-C-220   | AM168035.1    |
| Dicystostelium crassicaule    | 93HO-33     | AM168037.1    |
| Dicystostelium dimigraformum  | AR5b        | AM168038.1    |
| Dicystostelium discoideum     | V34         | AM168039.1    |
| Dicystostelium firmibasis     | TNS-C-14    | AM168041.1    |
| Dicystostelium garrantum      |             | GQ496161.1    |
| Dicystostelium giganteum      | WS589       | AM168042.1    |
| Dicystostelium implicatum     | 93HO-1      | AM168043.1    |
| Dicystostelium insulinativatis| 1-2         | MK322958      |
| Dicystostelium intermedium    | PJ11        | AM168044.1    |
| Dicystostelium leptosomum     | NZN49A      | HQ141480.1    |
| Dicystostelium longosporum    | TNS-C-109   | AM168048.1    |
| Dicystostelium macrocephalum  | B33         | AM168049.1    |
| Dicystostelium medium         | TNS-C-205   | AM168050.1    |
| Dicystostelium mucoroides     | Sweden 20   | HQ141482.1    |
| Dicystostelium purpureum      | C143        | AM168060.1    |
| Dicystostelium pseudobrefeldianum | 91HO8     | AM168059.1    |
| Dicystostelium quercibrachium | NZ201B      | HQ141479.1    |
| Dicystostelium robustum       | TNS-C-219   | AM168064.1    |
| Dicystostelium rosarium       | M45         | AM168065.1    |
| Dicystostelium septentrionale | IY49        | AM168066.1    |
| Dicystostelium sphaerocephalum| GR11        | AM168068.1    |
| Dicystostelium valdivianum    |             |              |
| Hagiwaraea coerulespites      | CRLC53B     | AM168036.1    |
| Hagiwaraea lavandula          | B15         | AM168047.1    |
| Hagiwaraea radiculata         | MLSA        | HQ141494.1    |
| Hagiwaraea rhizophodium       | AusKY-4     | AM168063.1    |
| Hagiwaraea vinaceofusca       | CC4         | AM168062.1    |

(Continued on next page)
TABLE 1 (Continued)

| Taxon                        | Isolate no. | Accession no. |
|------------------------------|-------------|---------------|
| Heterostelium anisocaule     | NZ478       | AM168096.1    |
| Heterostelium asymmetricum   | OH567       | AM168097.1    |
| Heterostelium australicum    | NB1AP       | HQ141508.1    |
| Heterostelium candidum       |             | AY040337.1    |
| Heterostelium colligatum     | HN13C1      | HQ141505.1    |
| Heterostelium equisetoides   | B7JB        | AM168099.1    |
| Heterostelium filamentosum   | SU-1        | AM168100.1    |
| Heterostelium flexuosum      | AU4B        | HQ141500.1    |
| Heterostelium gloeosporum    | TCK52       | AM168074.1    |
| Heterostelium luridum        | LR-2        | AM168101.1    |
| Heterostelium multicystogenum| A52         | HQ141506.1    |
| Heterostelium oculare        |             | HQ141497.1    |
| Heterostelium pallidium      | TNS-C-98    | AM168103.1    |
| Heterostelium pseudocandidum | TNS-C-91    | AM168107.1    |
| Heterostelium rotatum        | QC2C        | HQ141501.1    |
| Heterostelium stolonicoideum | K12A        | HQ141507.1    |
| Heterostelium tenuissimum    | TNS-C-97    | AM168105.1    |
| Heterostelium tikalense      | HH1C1       | HQ141509.1    |
| Heterostelium tikalense      | OH595       | AM168106.1    |
| Physarum polycephalum        | CL          | X13160.1      |
| Polysphondylium fuscans      | Sweden-11D  | JX173877.1    |
| Polysphondylium laterosorum  | AE4         | AM168046.1    |
| Polysphondylium patagonicum  |             | Q496156.1     |
| Polysphondylium violaceum    | 209         | HQ141486.1    |
| Raperostelium australe       | NZ808B      | AM168029.1    |
| Raperostelium gracile        | TNS-C-1B3   | AM168078.1    |
| Raperostelium ibericum       | 214riBu     | HQ141495.1    |
| Raperostelium minutum        | 71-2        | AM168051.1    |
| Raperostelium monochasoioides| HAG653      | AM168052.1    |
| Raperostelium ohiense        | Okla4C      | HQ141493.1    |
| Raperostelium potamoideus    | FP1A        | AM168069.1    |
| Raperostelium tenue          | PR4         | AM168075.1    |
| Rostrostelium ellipticum     | AE2         | AM168112.1    |
| Synstelium polycarpum        | VE1b        | AM168057.1    |
| Tieghemostelium lacteum      |             | AM168045.1    |

*A new sequences are indicated in bold.*

ACKNOWLEDGMENTS

Special thanks are extended to Steven L. Stephenson’s wife (Barbara Stephenson) for assisting with the field collecting. We express our appreciations to two anonymous reviewers for their valuable comments relating to the manuscript. Carlos Rojas supplied the map used in Fig. 1.

The project reported here was supported by a grant (W463-16) from the National Geographic Society to S.L.S., who collected the samples on Christmas Island and prepared the manuscript, whereas the laboratory research was supported by funding from the National Natural Science Foundation of China (grants 31870015 and 31300016) and the Science and Technology Development Program of Jilin Province (No. 20180101273JC) to P.L., who carried out the lab work and helped prepare the manuscript.

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