

## The complex case of the calcareous sponge Leucosolenia complicata (Porifera: Calcarea): hidden diversity in Boreal and Arctic regions with description of a new species

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#### The complex case of the calcareous sponge Leucosolenia complicata (Porifera: Calcarea): hidden diversity in Boreal and Arctic regions with description of a new species

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| Abstract:        | In this study, we present the first integrative revision of the Boreal and<br>Arctic calcareous sponges of the genus Leucosolenia with a specific focus<br>on its biodiversity in the White Sea. The material for this work included a<br>combination of newly collected specimens from different regions of the<br>North-East Atlantic and the White Sea and historical museum collections.<br>An integrative analysis was implemented based on vast morphological<br>data (light microscopy, scanning and transmission electron microscopy),<br>microbiome observations, ecological data, accompanied by molecular<br>phylogenetic and species delimitation analyses based on three nuclear<br>markers (28S rRNA, 18S rRNA and histone H3). We demonstrate that<br>Leucosolenia complicata, previously reported from Arctic waters, is<br>restricted to the North-East Atlantic, while in the Arctic, the Leucosolenia<br>diversity is represented by at least four species: Leucosolenia<br>corallorrhiza, Leucosolenia variabilis and two new species, one of which<br>is described herein under the name Leucosolenia creepae sp. nov. The<br>molecular phylogeny analysis supports the species identity of these<br>species. In addition to conventional morphological characters, new<br>informative fine morphological characters (skeleton and oscular crown<br>organisation; cytological structure including morphotypes of symbiotic<br>bacteria) were found, providing a baseline for further revision of this<br>group in other regions. |  |
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#### 1 Abstract

In this study, we present the first integrative revision of the Boreal and Arctic calcareous sponges of the genus *Leucosolenia* with a specific focus on its biodiversity in the White Sea. The material for this work included a combination of newly collected specimens from different regions of the North-East Atlantic and the White Sea and historical museum collections. An integrative analysis was implemented based on vast morphological data (light microscopy, scanning and transmission electron microscopy), microbiome observations, ecological data, accompanied by molecular phylogenetic and species delimitation analyses based on three nuclear markers (28S rRNA, 18S rRNA and histone H3). We demonstrate that *Leucosolenia complicata*, previously reported from Arctic waters, is restricted to the North-East Atlantic, while in the Arctic, the Leucosolenia diversity is represented by at least four species: Leucosolenia corallorrhiza, Leucosolenia variabilis and two new species, one of which is described herein under the name Leucosolenia *creepae* sp. nov. The molecular phylogeny analysis supports the species identity of these species. In addition to conventional morphological characters, new informative fine morphological characters (skeleton and oscular crown organisation; cytological structure including morphotypes of symbiotic bacteria) were found, providing a baseline for further revision of this group in other regions. 

Keywords: biogeography, molecular phylogeny, phylogenetic systematics, species delineation,
species boundaries, North Atlantic, ultrastructure, scanning electron microscopy

#### 23 Introduction

The calcareous sponges (Class Calcarea) represent rather a small group in terms of biodiversity across all sponge taxa (<5% of diversity) (Manuel et al., 2004), but are characterized by a unique mineral skeleton of calcium carbonate spicules and display great diversity in their body organisation (Borojevic *et al.*, 1990). The classification of this group is currently facing many challenges and rearrangements at all taxonomical levels due to high levels of homoplasy and convergent evolution (Manuel et al., 2004; Dohrnmann et al., 2006; Voigt et al., 2012; Voigt & Wörheide, 2016; Alvizu et al., 2018). In the course of the last two decades, the integrative taxonomy has became the most popular approach to define and describe taxa on different taxonomic levels and produce a reliable phylogeny-based classification of living organisms (Dayrat, 2005; Padial et al., 2010; Schlich-Steiner et al., 2010). In many cases, such studies can resolve existing taxonomic disputes strictly and definitively (Goulding & Dayrat, 2016). Molecular phylogenetics augments the ability of a researcher to find species boundaries and thus taxonomically important morphological characters. The integrative studies of calcareous sponges gave a new insight into understanding the actual diversity and evolutionary history of this group, with numerous new taxa being described during the last years (Azevedo et al., 2017; Riesgo et al., 2018; van Soest & de Voogd 2018; Alvizu et al., 2019; Cóndor-Luján et al., 2019; Sanamyan et al., 2019; Chu et al., 2020; Klautau et al., 2020 and many others). 

The Russian Arctic represents a poorly studied region with an unknown diversity of calcareous sponges. At the same time, extensive studies of Calcarea in the western part of the Arctic were carried out, thanks to which new species were described and a deep revision of the species described by that time was carried out (Rapp et al., 2001; Rapp, 2006, 2015; Alvizu et al., 2019). In other regions, like the Mediterranean, Caribbean and some others, the diversity of calcareous sponges has been extensively studied using molecular methods (Cavalcanti et al., 2014; Klautau et al., 2016, 2021; Fontana et al., 2018; van Soest & de Voogd, 2018). However, contemporary studies of the Russian Arctic tend to lend heavily to traditional morphology with 

limited regard for newer methods and trends in taxonomy and biogeography (Breitfuss, 1898a.b.c. Koltun, 1952; Ereskovsky, 1994a). Despite limited taxonomic studies, calcareous sponges from the White and Barents Seas are widely involved in various ecological (Ereskovsky, 1994b,c; 1995a.b), physiological and embryological (Anakina, 1997; Anakina & Korotkova, 1989; Anakina & Drozdov, 2000, 2001) researches. Two of the focus-sponges, Leucosolenia complicata (Montagu, 1814) and Sycon ciliatum (Fabricius, 1780) have become rising model species in evolutionary developmental studies (Leininger et al., 2014; Fortunato et al., 2014, 2015, 2016; Ereskovsky et al., 2017a; Lavrov et al., 2018, 2022; Lavrov & Ereskovsky, 2022; Melnikov et al., 2022). 

The genus Leucosolenia Bowerbank, 1864 (phylum Porifera, class Calcarea, subclass Calcaronea) includes more than 40 distinct species distributed worldwide, being more specious in the northern regions (Burton, 1963; Borojevic *et al.*, 2000). The history of systematics of this group is complicated by the different species conceptions proposed by various researchers since Haeckel's monograph on calcareous sponges (Haeckel, 1872). Haeckel established 21 new genera, seven of which represent an asconoid type of organisation. However, he did not consider any generic names for calcareous sponges used before him; hence, this system has undergone various modifications (Minchin, 1904). At the species-level, many of his names and his 'connection varieties' for leucosolenoid sponges were further considered synonyms of three widely distributed species: Leucosolenia botrvoides (Ellis & Solander, 1786), L. complicata, Leucosolenia variabilis Haeckel, 1870 (Minchin, 1904, 1905). Further research highlighted the complications of taxonomical studies due to the absence of informative characters. As a result, some North Atlantic species, i.e., L. complicata, L. variabilis and L. botryoides were believed to have a wide cosmopolitan distribution ranging from the Arctic to South Africa, New Zealand and the Antarctic (Burton, 1963). Sarà (1956) found a connection between spicular characters in these species and suggested they may hybridise, which caused even more simplification of the system with a single species, L. botrvoides represented by different 'forms' of uncertain taxonomical status (Burton,

1963). With the advent of new microscopic techniques and molecular studies, the systematics of the genus *Leucosolenia* has received much attention during the last twenty years. The valid status of some species was reconsidered, e.g., North-eastern Atlantic Leucosolenia corallorrhiza (Haeckel, 1872) (Rapp, 2015) and Leucosolenia somesii (Bowerbank, 1874) (van Soest et al., 2007). Also, several new species were described based on studies using scanning electron microscopy that allowed the identification of fine spines on spicules (van Soest, 2017; Chu et al., 2020). Recent molecular phylogenetic analysis of North Atlantic *Leucosolenia* species showed the paraphyly of the genus (Alvizu et al., 2018) and the high rate of hidden biodiversity within it. Nevertheless, no integrative revision has been conducted yet, and the taxonomical status of most North-Atlantic species remains unverified.

The main goal of our study is to revise the diversity and taxonomy of the genus *Leucosolenia* from the White Sea using an integrative approach, which includes vast molecular, morphological and cytological data. Additionally, the specific aims of the study are to identify phylogenetically significant morphological characters and to propose optimal sets of molecular markers for further taxonomical and phylogenetic research on calcareous sponges.

#### Material and methods

#### *Material*

The representatives of the genus Leucosolenia were collected in the White Sea at N.A. Pertsov White Sea Biological Station MSU (66°34'N, 33°08'E) during 2016–2018 at the upper subtidal zone and by SCUBA diving (Table S1). The identification of the specimens was based on external and internal (skeletal) characteristics, such as spicule types and their spatial arrangement in different regions of the sponge body. Additionally, we studied specimens of Leucosolenia complicata collected in the English Channel (Roscoff, France) and a specimen of Leucosolenia somesii from the collection of the Zoological Museum of Amsterdam (ZMA). Several spicule slides from the collection of the British Museum of Natural History (BMNH) of L. complicata, L. *variabilis* and *L. somesii*, including the type material for the latter two species, were also examined for comparison with species from the White Sea (Table S1). Collection data, voucher numbers and GB accession numbers are summarised in Table S1. All new samples were fixed in 96% ethanol. Voucher specimens are deposited in the collections of the Zoological Museum of Moscow State University, White Sea Branch (WS). 

## 106 Taxon sampling

Our molecular sampling included 279 individuals from the White Sea, the Netherlands and Roscoff (Table S1). For species delimitation, the 28S C-region was sequenced for all specimens available for study (to make sure that the morphological identification is correct). In several cases, 28S did not give enough resolution to undoubtedly support the genetic distinctness of species; therefore, for a large number of samples from the White Sea, we additionally obtained a novel histone H3 marker (see below for details). Finally, 18S was obtained only for several specimens for the concatenated phylogenetic analysis to test the monophyly of the genus *Leucosolenia*.

Seventeen *Leucosolenia* specimens from the White Sea, Greenland, Norway and the NorthEast Atlantic (unspecified), for which only GenBank sequences were available, were also included
in the analyses (Table S1). The final datasets for each gene included as many specimens from

different localities as possible to improve the resolution of phylogenetic reconstructions. *Plectroninia novaecaledoniense* Vacelet, 1981 and *Clathrina arnesenae* (Rapp, 2006) were
chosen as outgroups based on recent papers on the calcaronean phylogeny (Voigt & Wörheide,
2016; Alvizu *et al.*, 2018). To test the identity of high-level taxonomic groups, specimens included
in the most recent Calcaronea phylogenetic study (Alvizu *et al.*, 2018) were used in the
concatenated analysis (Table S1).

#### 123 DNA extraction, amplification and sequencing

DNA was extracted from small pieces of tissue using PALL AcroPrep 96-well plates (PALL Corp., USA) (Ivanova *et al.*, 2006) and the Diatom DNA Prep100 kit (Isogen Lab, Russia) according to manufacturer protocol with minor modifications: an increased lysis stage time for better dissolving of tissues and a reduced volume of extraction reagents at the final step for increasing the final DNA concentration. Extracted DNA was used as a template for the amplification of partial 28S rRNA (C-region, LSU) and 18S rRNA (SSU). To support the results obtained with ribosomal markers, we also included the nuclear protein-coding gene histone H3 (H3) in the analysis. Although this marker is commonly used in systematic and phylogenetic studies in other invertebrate groups, sequences for the calcarean sponges are absent in GenBank. Primers were modified from standard H3 metazoan primers (H3AF: 5'-ATG GCT CGT ACC AAG CAG ACV GC-3' H3AR: 5'-ATA TCC TTR GGC ATR ATR GTG AC-3' see Colgan, 1998), the annotated transcriptome of *S. ciliatum* (Fortunato *et al.*, 2014) was used for a primer design (Table 1). Polymerase chain reactions were carried out in a 25-µL reaction volume, which included 5 µL of 5x Taq Red Buffer (Eurogen Lab, Russia), 0.5 µL of HS-Taq Polymerase (Eurogen Lab, Russia), 0.5 µL of dNTP (50 µM stock), 0.3 µL of each primer (10 µM stock), 1 µL of genomic DNA and 17.7 µL of sterile water. The PCR conditions for the corresponding primers are given in Table 1. Sequencing for both strands proceeded with the Big Dye Terminator v3.1 sequencing kit (Applied Biosystems, USA), the same primers as for PCR were used. 

142 Sequencing reactions were analysed using the ABI 3500 Genetic Analyzer (Applied Biosystems,

143 USA). All new sequences were deposited in GenBank (Table S1).

#### 144 Phylogenetic reconstruction

Raw reads for each gene were assembled and checked for improper base-calling using GeneiousPro 4.8.5 (Biomatters, New Zealand). Original data and publicly available sequences were aligned with the MUSCLE (Edgar, 2004) algorithm in MEGA7 (Kumar et al., 2016). Protein-coding sequences were translated into amino acids to verify the coding sequences. The resulting alignments were 353 bp for 28S, 1493 bp for 18S and 359 bp for H3. Phylogenetic analyses were conducted for each marker individually and for the concatenated dataset. Sequences were concatenated by a simple biopython script (Chaban et al., 2019). The best-fitting nucleotide evolution model was tested in MEGA7 based on the Bayesian information criterion (BIC) for each partition. The best-fitting model for the 28S partition was HKY+G, for the 18S partition – K2, and for the H3 partition – K2P+G. Phylogenetic reconstruction of the concatenated dataset was performed applying evolutionary models for partitions separately. The Bayesian estimation of posterior probability for all datasets was performed in MrBayes 3.2 (Ronquist & Huelsenbeck, 2003). Markov chains were sampled at intervals of 500 generations. The analysis was started with random starting trees and 10<sup>7</sup> generations. Maximum likelihood-based phylogeny inference for all datasets was performed in the HPC-PTHREADS-AVX version of RaxML (Stamatakis, 2014) with ultrafast bootstrapping (UFBoot approximation approach) (Minh et al., 2013) in 1000 pseudoreplicates under the GTRCAT model of nucleotide evolution. Bootstrap values were placed on the best tree found with SumTrees 3.3.1 from DendroPy Phylogenetic Computing Library Version 3.12.0 (Sukumaran & Holder, 2010). The final phylogenetic tree images were rendered in FigTree 1.4.0.

#### 165 Species delimitation

Uncorrected inter- and intra-specific *p*-distances were calculated in MEGA7 (Kumar *et al.*,
2016). Assemble Species by Automatic Partitioning (ASAP) (Puillandre *et al.*, 2021), calculation

of uncorrected *p*-distances and single-gene trees were applied to assist in the species delimitation analysis. 28S and H3 sequences of Leucosolenia species (excluding outgroups) used in the phylogenetic analysis, were aligned for ASAP analysis with the MUSCLE algorithm (Edgar, 2004) in MEGA7 (Kumar et al., 2016). 18S sequences were not used in the analysis due to low substitution rates. The analysis was run on the online version of the programme (https://bioinfo.mnhn.fr/abi/public/asap/asapweb.html) with the default setting and three proposed models: Jukes-Cantor (JC), Kimura (K80) and Simple distance. Uncorrected *p*-distances were calculated for the same alignments used for ASAP in MEGA7 (Kumar et al., 2016). Single-gene trees were calculated using the Maximum Likelihood approach in HPC-PTHREADS-AVX version of RaxML (Stamatakis, 2014) with ultrafast bootstrapping (UFBoot approximation approach) (Minh et al., 2013) with 1000 pseudoreplicates under the GTRCAT model of nucleotide evolution. 

A visualisation of character heterogeneity was assessed using PopART 1.7 (http://popart.otago.ac.nz) (Leigh & Bryant, 2015) with the TCS network algorithm (Clement *et al.*, 2002) and a connection limit of 5%. The resulting networks were edited in Adobe Illustrator CS 2015 to highlight certain features.

#### 184 Morphological studies

The external morphology of each species was studied under Leica M165FC stereomicroscope (Leica, Germany) equipped with a digital camera Leica DFC420 (Leica, Germany). Extraction of spicules and slide preparation for a total of 34 specimens were made according to standard protocols (Klautau & Valentine, 2008). The general skeleton morphology of both the oscular rim and cormus tubes was studied under Leica M165C stereomicroscope (Leica, Germany). For detailed studies of skeleton morphology, parts of the oscular rim and cormus tubes were treated with Murray's Clear according to the standard protocol to clarify soft tissues (Miller *et al.*, 2005). Slides of dissociated spicules and skeletons were studied under Zeiss Axioplan 2 (Carl Zeiss, Germany) and Leica DM2500 (Leica, Germany) with the digital cameras

AxioCam HRm (Carl Zeiss, Germany) and Leica DFC420C (Leica, Germany), respectively. Scanning electron microscopy (SEM) analysis of the spicules was performed under Carl Zeiss EVO-40 (Carl Zeiss, Germany), Hitachi S-405A (Hitachi, Japan) and CamScan S2 (Clinton Electronics Corp., UK) scanning electron microscopes. For this purpose, isolated spicules in 96% ethanol were transferred to cover slips, which were mounted onto stubs with nail polish, dried and sputter-coated. Figures of spicules under SEM are montages of the most typical spicule morphology obtained from a number of specimens of each studied species.

The measurements of the spicules (length and basal width at the base of the actines) were
made for every spicule category on SEM images using ImageJ v. 1.48 software (National Institute
of Health, USA). Strait Line tool was used for straight spicules and rays, Segmented Line tool was
used for curved and undulating spicules and rays.

#### 205 Cytological studies

For semithin sections and transmission electron microscopy (TEM), sponges were fixed with 2.5% glutaraldehyde (Ted Pella, USA) on 0.2M Millonig's phosphate buffer (Millonig, 1964), and postfixed with 1% OsO<sub>4</sub> (Spi Supplies, USA) according to standard protocol (Lavrov et al., 2016a, 2018; Lavrov & Ereskovsky, 2022). Specimens were embedded in SPI-Pon 812/Araldite 6005 epoxy embedding media (Spi Supplies, USA) according to the manufacturer's instructions. Semi-thin sections (1 µm in thickness) were cut on a Reichert Jung ultramicrotome (Reichert, USA) equipped with a "Micro Star" 45° diamond knife before being stained with toluidine blue and observed under a WILD M20 microscope (Wild-Leitz, Germany). Digital photographs were taken with a Leica DMLB microscope (Leica Microsystems, Germany) using the Evolution LC colour photo-capture system (Media Cybernetics, USA). Ultrathin sections (60-80 nm) were cut with a Leica UCT6 (Leica, Germany) and PowerTome XL ultramicrotomes equipped with a Drukkert 45° diamond knife. Ultrathin sections, contrasted with uranyl acetate, were observed under Zeiss-1000 TEM (Carl Zeiss, Germany) and Tecnai G2 20 TWIN (FEI Company, USA). The detailed cytological studies were conducted only for *Leucosolenia corallorrhiza*, as it represents the most

 common species in the White Sea. For other species, only the general morphology of the cell types

221 was described. Figures of cell types are montages of the most typical cell morphology obtained

222 from a number of specimens of each studied species.

#### 223 Nomenclatural acts

The electronic edition of this article conforms to the requirements of the amended International Code of Zoological Nomenclature (ICZN), and hence the new name contained herein is available under that Code from the electronic edition of this article. The LSID for this publication is: urn:lsid:zoobank.org:pub:BA13614B-2884-4E02-9D45-0EE44CBD01A5.

#### **Results**

#### Phylogenetic reconstruction

We obtained 486 new sequences of different Leucosolenia species (Table S1). The sequence alignment of concatenated 28S, 18S and H3 loci includes 2160 positions. Single-gene trees based on 28S and H3 loci give good resolution at the species level while, 18S single-gene tree is unresolved due to low substitution rates (Data S1).

The topology of the resulting concatenated trees from Bayesian Inference (BI) and Maximum Likelihood (ML) analyses are congruent and well-supported in most cases (Figs 1, S1, S2). The genus Leucosolenia is recovered monophyletic with moderate support (posterior probabilities from BI (PP) = 0.9; bootstrap support from Maximum Likelihood (ML) = 92). Within *Leucosolenia* clade three species groups are monophyletic and highly supported: Clade I, including most specimens from the White Sea (Leucosolenia sp. 1, Leucosolenia sp. 3 and Leucosolenia sp. 4) and several specimens from GenBank initially identified as Leucosolenia cf. variabilis, *Leucosolenia* cf. *corallorrhiza* and *Leucosolenia* sp. 1 (PP = 1; ML = 99); Clade II with specimens from the White Sea (Leucosolenias sp. 2), Leucosolenia somesii from Netherlands, Leucosolenia sp. from GenBank (accession number AF100945) and *Leucosolenia botryoides* from GenBank (voucher number SA60) (PP = 1; ML = 93); and Clade III, which comprised only *Leucosolenia* complicata specimens from Roscoff (ws11881-11883) and from GenBank (PP = 1; ML = 100). The Clade III (L. complicata) is sister to the Clade II (PP = 1; ML = 93). Within the Clade I at least six monophyletic units are found: (1) Leucosolenia sp. 1, clustering with Leucosolenia cf. variabilis FB33 and Leucosolenia cf. variabilis FB12 from GenBank (PP = 0.99; ML = 96), (2) Leucosolenia cf. variabilis FB58 and Leucosolenia cf. variabilis FB60 (PP = 1; ML = 100), (3) *Leucosolenia* sp. 3 and *Leucosolenia* cf. *variabilis* SA62 (PP = 1; ML = 98), (4) *Leucosolenia* sp. 4 and *Leucosolenia* cf. *corallorrhiza* FB14, FB20, SA43 (PP = 0.81; ML = 100), (5) Leucosolenia cf. *corallorrhiza* FB59 and SA44 (PP = 1; ML = 100) and (6) *Leucosolenia* sp. 1 FB73 and FB81 from GenBank (PP = 0.61; ML = 100). Representatives of (1), (2), (4), (5) clades form a compact 

monophyletic group (PP = 0.99; ML = 96) with very low genetic distances within it (Table 2). *Leucosolenia* sp. 3 is recovered sister to it (PP = 0.92; ML = 98), and *Leucosolenia* sp. 1 from
GenBank shows sister relationships to other representatives of Clade I.

Within Clade II, *L. somesii* specimens from the Netherlands form a single clade with *Leucosolenia* sp. from GenBank. *Leucosolenia* sp. 2 also forms a well-supported clade (PP = 1;
ML = 93). It represents a monophyletic group with *L. botryoides* and *L. somesii* (PP = 1; ML =
93), but relationships within this clade are unsupported.

262 Species delimitation

ASAP analysis of 271 LSU sequences recovers a different number of operational taxonomical units (OTUs) depending on the ASAP score (Data S2). The lowest ASAP score is found for threshold distances of 2.16%, 2.29%, 4.30% or 6.68% in this case five OTUs are recovered: four of them correspond to species from Clades II and III (Leucosolenia complicata, L. botryoides, L. somesii and Leucosolenia sp. 2) and the fifth group includes all specimens of Clade I. A scenario revealing the same 10 species-level units as in the phylogenetic analysis receives a relatively high ASAP score (5.00) with a threshold distance of 0.69%. ASAP analysis of the H3 dataset (170 sequences) contains only five candidate species (Leucosolenia sp. 1, Leucosolenia sp. 2, Leucosolenia sp. 3, Leucosolenia sp. 4, L. complicata) due to the absence of H3 sequences for calcareous sponges in GenBank. The lowest ASAP score (2.00) receives a scenario with all five candidate species as distinct (the threshold distance is 1.38%). Scenarios with an ASAP score of 2.5-4.0 reveal 2-4 candidate species; the identities of Leucosolenia sp. 1, Leucosolenia sp. 3, and Leucosolenia sp. 4 are not supported in this case. 

A visualisation of character heterogeneity using the Medium Parsimony Network (TCS algorithm) reveals similar results in the 28S and H3 datasets (Fig. 2). On the 28S network, each candidate species either forms a distinct group (*Leucosolenia* sp. 1, *Leucosolenia* sp. 2, *Leucosolenia* sp. 3, *L. complicata*) or has a unique genotype (*Leucosolenia* sp. 4, *L. somesii, L. botryoides* and three candidate species from GenBank) (Fig. 2A). These groups, or genotypes,

differ from each other by three substitutions among species of Clade I and by 8–11 substitutions
among species of Clade II. The intraspecific differences do not exceed two substitutions. There
are 15–22 substitutions between representatives of Clades I, II, III.

The same results are observed in the H3 dataset, but the overall nucleotide diversity is higher (Fig. 2B). There are 1-2 substitutions within each candidate species except *Leucosolenia* sp. 3, where genotypes differ by 1–4 substitutions. Differences of 6–23 substitutions are found between candidate species.

Uncorrected *p*-distance values of 28S and H3 markers are presented in Tables 2–3 (data from GenBank are not included). Overall, intraspecific *p*-distances of 28S within Clade I show an overlapping range with interspecific distances (0–0.4% intraspecific, 0.4–1.3% intraspecific distances). The distances between the three large clades vary from 5.1 to 10.5%. The H3 marker shows a higher diversity: intraspecific distances vary from 0 to 1.7%, while interspecific distances vary from 2.5 to 5% within larger clades and are of 7.9–10.9% between the clades.

#### 294 Comparison of morphological and molecular data

Three recovered monophyletic lineages corresponding to the genus *Leucosolenia*, have distinct morphological traits. They differ in the external appearance, the general skeleton composition, the spicular set and the cellular composition (Figs 3-24; Table 4-9). Definitions and technical terms used in this study can be found elsewhere (Boury-Esnault & Rützler, 1997; Ereskovsky & Lavrov, 2021; Łukowiak *et al.*, 2022).

Clade III is represented by a single species, *Leucosolenia complicata*, which was found only in European waters (Roscoff and Norway in our material and public data). Although the type material for this species is not known if it ever existed, morphologically our specimens perfectly fit the descriptions made by previous authors with few exceptions (Haeckel, 1872; Minchin, 1904; Rapp, 2015). The main diagnostic traits in spicular characters are (Fig. 4): (1) two populations of diactines, large lanceolate smooth diactines and small trichoxeas with irregular spines; (2)

306 parasagittal tri- and tetractines, with the unpaired actine is commonly longer than the paired ones;

307 (3) tetractines are commonly found in both the oscular region and the cormus.

The studied representatives of Clade II (Leucosolenia sp. 2 and Leucosolenia somesii) have an echinate external appearance due to the high number of diactines protruding through the external surface (Fig. 19). Diactines are non-lanceolate, with rows of spines on the outer tip (commonly only two rows are visible, but there are more) or smooth (Fig. 20A, B). In these species, tri- and tetractines are thinner than in representatives of Clade C (see below), and abnormal spicules are common (Fig. 20C-E; Table 8). At the same time, Leucosolenia sp. 2 differs from Leucosolenia somesii by having only a single population of spined diactines, while Leucosolenia somesii possesses a second type of smooth non-lanceolate diactines (Fig. 24A-C).

Species of Clade I have several common traits: all of them have unique cells with inclusions
(see below) (Figs 10C-E, 15C, E-F; Table S2); tri- and tetractines are predominantly T-shaped, the
angle between paired actines is commonly 130-150° (Figs 8C-D, 12B, D, 18D, F; Table 5-7);
lanceolate diactines are always present, with or without spines (Figs 8A-B, 12A, 13, 18A-C).

Within this clade, *Leucosolenia* sp. 1 has the largest angle between the paired actines (mean 140°) and the unpaired actine is much shorter than the paired ones in tri- and tetractines (Fig. 8C-D; Table 5). Also, it has only lanceolate diactines bearing short spines on the lanceolate tip in some cases (Fig. 8A-B). These characters perfectly fit the description and illustrations of *L. corallorrhiza* (Haeckel, 1872), but not *L. variabilis*.

*Leucosolenia* sp. 3 commonly forms a large voluminous rounded cormus with a very large oscular tube (there may be more than one oscular tube, but one of them is always enlarged) (Fig. 11A). Other representatives of Clade A commonly have cormus spreading along substrate with numerous oscular tubes of more or less equal size (Figs 7A, 17A). In spicular morphology, this species has a unique type of diactines: extremely long and thin trichoxeas, covered with irregularly placed spines (Fig. 13). Lanceolate diactines lack spines (Fig. 12A). The length of unpaired actines in tri- and tetractines is commonly the same as that of paired ones (Fig. 12B-D; Table 6). The

shape and measurements of spicules, as well as overall body shape mostly resemble those states
described for *L. variabilis* (Haeckel, 1872).

Leucosolenia sp. 4 is very similar to Leucosolenia sp. 1 in external appearance. In spicular characters, this species contains very few tetractines in both the oscular region and the cormus (Fig. 17B-D). Triactines are usually with undulated paired actines (Fig. 18F). We also detected a high amount of abnormal tri- and tetractines (Fig. 18E), while in other species their amount is lower. Although these differences seem to be valuable to support the species distinctness, the limited studied material (only three specimens available) does not allow us to test for possible intraspecific variation. Also, this species demonstrates low genetic divergence from *Leucosolenia* sp. 1.

To sum up, our integrative approach indicates that the species *Leucosolenia complicata* is restricted to European waters, while in the White Sea, the genus *Leucosolenia* is represented by a complex of four species: *Leucosolenia corallorrhiza* (= *Leucosolenia* sp. 1), *Leucosolenia variabilis* (= *Leucosolenia* sp. 3), and two undescribed species. *Leucosolenia* sp. 2 is described herein under the name *Leucosolenia creepae* sp. nov., while in the case of *Leucosolenia* sp. 4 (further named *Leucosolenia* sp. A), more material is needed to confirm or reject its identity as a separate species from the closely related *Leucosolenia corallorrhiza*.

- 350 Systematic descriptions
- 351 Subclass Calcaronea Bidder, 1898
- 352Order Leucosolenida Hartman, 1958
- 353 Family Leucosoleniidae Minchin, 1990
- *Leucosolenia* Bowerbank, 1864
- *Type species: Spongia botryoides* (Ellis & Solander, 1786) (by original designation).
- *Type locality:* Harbour near Emsworth, between Sussex and Hampshire, the English Channel.

*Diagnosis* (based on Hooper et al., 2002): Leucosoleniidae, in which the skeleton can consist of

358 diactines, triactines and/or tetractines. There is no reinforced external layer on the tubes.

- *Leucosolenia complicata* (Montagu, 1814)
- 361 (Figs 3-6; Table 4)
- *Type material:* Not known, probably lost.
- *Type locality:* British Isles, Devon coast (Montagu, 1818).
- *Material studied:* 3 specimens. Molecular data 3 specimens (WS11661, WS11662, WS11663),
- seternal morphology 3 specimens (WS11661, WS11662, WS11663), skeleton organisation 1

**366** specimen (WS11662), spicules (SEM) – 3 specimens (WS11661, WS11662, WS11663), cytology

367 (TEM) – 3 specimens (WS11661, WS11662, WS11663) (Table S1).

*External morphology:* Cormus more or less spherical, bearing multiple erect oscular tubes with 369 short lateral diverticula in basal part (Fig. 3A). Prominent perioscular spicular crown absent (Fig.

- 370 3B). Surface minutely hispid. Colouration of living and preserved specimens grayish white (Fig.
- 371 3A).

*Spicules:* Diactines (Fig. 4A-C). Two populations: (1) Curved lanceolate diactines (Fig. 4A), mean
length 263.7 μm, mean width 9.5 μm (Table 4). Slightly curved, smooth, with lanceolate outer tip,
variable in length. (2) Trichoxeas (Fig. 4B), mean length 127.3 μm, mean width 2.4 μm (Table 4).
Thin, straight, narrowing toward outer end, both ends pointed, not lance-shaped. Numerous
irregularly distributed spines, number and size of spines decrease toward inner end (Fig. 4C).

377 Triactines (Fig. 4D). Predominantly parasagittal V-shaped (mean angle 125.7°), with 378 unpaired actine usually longer than paired (mean length: 113.5  $\mu$ m – unpaired, 94.9  $\mu$ m – paired) 379 (Table 4), but equal and shorter unpaired actines also occur. Unpaired actine usually slightly 380 slender than paired (mean width: 6.3  $\mu$ m – unpaired, 6.8  $\mu$ m – paired) (Table 4). T-shaped sagittal 381 triactines absent.

Tetractines (Fig. 4E). Predominantly parasagittal V-shaped (mean angle 123.5°). Unpaired actine usually longer than paired, rarely equal (mean length: 109.3  $\mu$ m – unpaired, 93.9  $\mu$ m – paired, 23.8  $\mu$ m – apical) (Table 4). Paired and unpaired actines equal in width, apical actine more slender (mean width: 6.7  $\mu$ m – unpaired, 6.9  $\mu$ m – paired, 5.3  $\mu$ m – apical) (Table 4). Apical actine curved and smooth.

*Skeleton:* Skeleton of both oscular and cormus tubes predominantly formed by tetractines, triactines quite rare (Fig. 3C, D). In oscular tubes spicules constitute organized array with their unpaired actines directed toward cormus and oriented more or less in parallel to proximo-distal axis of oscular tube (Fig. 3C). In cormus tubes spicule array less organized (Fig. 3D). Lanceolate diactines cover tubes surface, orienting in different directions and extending outside by lanceshaped tip. Trichoxeas sparsely distributed on outer surface. No prominent spicular crown on oscular rim (Fig. 3B).

*Cytology:* Body wall, 6-9 μm thick, three layers: exopinacoderm, loose mesohyl, choanoderm
395 (Fig. 5A, B; Table S2). Flat endopinacocytes located in only distal part of oscular tube (oscular
396 ring) replacing choanocytes. Inhalant pores scattered throughout exopinacoderm, except the
397 oscular ring area.

Exopinacocytes non-flagellated T-shaped, rarely flat (Fig. 5C). External surface covered by
glycocalyx. Cell body (height 4.8 μm, width 2.8 μm), containing nucleus (diameter 2.2 μm),
submersed in mesohyl (Fig. 5C). Cytoplasm with specific spherical electron-dense inclusions (0.30.4 μm diameter) (Fig. 5C).

402 Endopinacocytes non-flagellated flat cells, size 20-30  $\mu$ m × 2-2.5  $\mu$ m (Fig. 5D). External 403 surface covered by glycocalyx. Nucleus (2.4x1.8  $\mu$ m) oval with or without nucleolus. Cytoplasm 404 without specific inclusions (Fig. 5D).

405 Choanocytes flagellated trapeziform or prismatic (height 6 μm, width 3.7 μm) (Fig. 5E).
406 Flagellum surrounded by collar of microvilli. Characteristic pyriform nucleus (diameter 2.3 μm)
407 in apical position. Cytoplasm with phagosomes and small vacuoles (Fig. 5E).

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408 Porocytes tubular cylindrical (height 4.5  $\mu$ m, width 2  $\mu$ m), connecting external milieu with 409 choanocyte tube (Fig. 5B, F). Nucleus oval to spherical, diameter 1.8  $\mu$ m, some time with 410 nucleolus. Cytoplasm with phagosomes and small vacuoles (Fig. 5F).

411 Sclerocytes amoeboid, size 4  $\mu$ m × 2  $\mu$ m (Fig. 6A). Nucleus usually oval or pear-shaped 412 (diameter 1.6  $\mu$ m), containing single nucleolus. Well-developed Golgi apparatus and rough 413 endoplasmic reticulum. Cytoplasm usually with phagosomes and/or lysosomes (Fig. 6A).

414 Amoebocytes of different shape (from oval to amoeboid) without special inclusions, size 5.8
 415 μm × 3.4 μm (Fig. 6B). Nucleus spherical (diameter 2.2 μm), sometimes with nucleolus.

Two morphotypes of bacterial symbionts in mesohyl. Morphotype 1 most abundant. Bacteria
large rod-shaped slightly curved, diameter 0.3 μm, length 2.2 μm (Fig. 6C). Double cell wall,
cytoplasm transparent, nucleoid region filamentous.

419 Morphotype 2 rare. Bacteria rod-shaped, diameter 0.18 μm, length 1.2 μm (Fig. 6D). Double
 420 cell wall, cytoplasm transparent, nucleoid region filamentous.

421 *Distribution:* Boreal species. Molecular species identity confirmed for specimens from France
422 (Roscoff). Live in low intertidal and subtidal zones up to 20 m depth, on rocks and kelps (Borojevic
423 et al., 1968).

# 424 *Reproduction:* The specimens collected in February 2017 in Roscoff contain oocytes at the early 425 stages of development.

426 Remarks: Leucosolenia complicata was one of the most undoubted species described in the XIX 427 century. According to our data, it shows stable internal characters and easily diagnosable external 428 features, *i.e.*, erect multiply oscular tubes extending from the small cormus. The species identity 429 and validity of L. complicata are strongly supported by our molecular data as well. It represents a 430 distinct monophyletic lineage on all phylogenetic trees, and p-distance values to other 431 Leucosolenia species are very high (more than 5% in LSU and 3.8% in SSU). Extensively studied 432 morphology allows clarifying the species diagnosis, which varied from author to author (Haeckel, 433 1872; Minchin, 1904; Jones, 1954; Rapp, 2015): small cormus, erect multiple oscular tubes, two

populations of diactines (curved lanceolate diactines and small trichoxea), parasagittal tri- and tetractines with predominately longer unpaired actines, skeleton of tubes predominately formed by tetractines. Leucosolenia complicata is easily differentiated from other Leucosolenia species (Leucosolenia variabilis, L. somesii and others) in these traits. In addition, the mesohyl cell composition of L. complicata is very poor compared to other studied Leucosolenia species: the mesohyl contains only sclerocytes and amoebocytes (Table S2). The composition of symbiotic bacteria (two morphotypes of rod-shaped bacteria) differs in L. complicata from L. corallorrhiza and L. variabilis (Table S2).

Although the type material of this species is not available, if it ever existed, we studied spicule slides from Minchin type collections (BMNH 1910.1.1.415a and BNMH 1910.1.1.435.Aa). They are listed as the type material of *L. complicata* in the BMNH collection. These slides contain hand-written information on the corresponding paragraphs in Minchin (1904) work with relevant collection information (slides nos. 1, 2; Minchin, 1904:372). Accordingly, both slides appeared from Canon Normans's Collection. The specimen BMNH 1910.1.1.415a was collected at Scarborough (the North Sea) by Bean and sent to Haeckel for examination. The specimen BNMH 1910.1.1.435.Aa was collected at Guernsey Islands (the English Channel) by J. Bowerbank and likely represents a syntype of *Ascandra contorta* (Bowerbank, 1866). According to Minchin (1904), this slide contains an admixture of *L. complicata* spicules with *A. contorta*. All this indicates that slides BMNH 1910.1.1.415a and BNMH 1910.1.1.435.Aa are not the type material of *L. complicata*, and the label 'type' likely refers to the Minchin's type collection, which contained most typical specimens. Since no type material exists, the designation of neotype is needed once the material from the type locality (British Isles, Devon coast) becomes available for the molecular study.

457 Ascandra pinus Haeckel, 1872 and Leucosolenia fabricii Schmidt, 1869 are regarded herein
458 as minor synonyms. Ascandra pinus lacks small trichoxeas, which were most probably overlooked
459 by Haeckel (Minchin, 1904), and in Leucosolenia fabricii, the skeleton is formed mostly by

triactines, which was considered intraspecific variation by many authors (Minchin, 1904; Burton,

1963; Rapp, 2015). However, our data shows that these characters may be regarded as diagnostic interspecific features, as shown for the L. variabilis species complex (see below); therefore, both of these species names should be taken into account for future research on European Leucosolenia. Our data also suggest the absence of *L. complicata* in the White Sea. In works by Breitfuss (1898a), three Leucosolenia species were found at different localities in the White Sea and described under the names Ascandra variabilis Haeckel, 1872, Ascandra contorta (Bowerbank, 1866) and Ascandra fabricii (Schmidt, 1869). Minchin (1904) later considered the latter two species *sensu* Breitfuss (1898a) as minor synonyms of *Leucosolenia complicata* due to external morphological characters, while spicular characters were ignored in most cases. However, Ascandra contorta sensu Breitfuss (1898a) possesses tri- and tetractines with short unpaired actines, which is most likely a diagnostic feature for L. variabilis. Due to the absence of L. *complicata* in our material from the White Sea and uncertainties in previous research, more material is required from different localities in the White and Barents Seas to clarify the distribution ranges of this species in Arctic waters. Leucosolenia corallorrhiza (Haeckel, 1872)

- =Ascortis corallorrhiza Haeckel, 1872
- *=Sycorrhiza corallorrhiza*, Haeckel, 1870
- =Auloplegma corallorrhiza Haeckel, 1872
- =Leucosolenia cf. variabilis (Alvizu et al., 2018; Lavrov et al., 2018).
- =Leucosolenia variabilis (Lavrov & Ereskovsky, 2022; Lavrov et al., 2022; Melnikov et al.,
- 2022).
  - =Leucosolenia complicata (Ereskovsky et al., 2017a).
- *Type material:* Type material is not known.

- *Type locality:* Haeckel based his description on one specimen from Norway and one from
  487 Greenland, without designating the type material (Rapp, 2015).
- *Material studied:* 177 specimens. Molecular data 177 specimens, external morphology 177
- 489 specimens, skeleton organisation 2 specimens (WS11650, WS11653), spicules (SEM) 5

11 490 specimens (WS11649, WS116450, WS11653, WS11657, WS11658), cytology (TEM) - 6

<sup>13</sup> 14 491 specimens (WS11631, WS11632, WS11634, WS11635, WS11636, WS11637) (Table S1).

- External morphology: Cormus formed by basal reticulation of tubes, from which erect oscular tubes and long diverticula arising. Sponge bear from one to multiple slightly curved oscular tubes with or without short lateral diverticula in basal part. Oscular tubes gradually narrows to oscular rim, possessing short spicular crown (Fig. 7A, B). Surface minutely hispid or echinate. Colouration of living and preserved specimens gravish white (Fig. 7A).
- 497 Spicules: Diactines (Fig. 8A, B). Curved lanceolate diactines, mean length 179 μm, mean width 6
   498 μm (Table 5). Slightly curved with lanceolate outer tip, variable in size, smooth or with few small
   499 spines at lanceolate tip (Fig. 8B).
  - 500 Triactines (Fig. 8D). T-shaped sagittal (mean angle 142.9°), unpaired actine usually shorter 501 than paired (mean length: 70.5  $\mu$ m – unpaired, 82.7  $\mu$ m – paired) (Table 5), rarely equal. Actines 502 equal in width (mean width: 6.5  $\mu$ m – unpaired, 6.5  $\mu$ m – paired) (Table 5).
- Tetractines (Fig. 8C). T-shaped sagittal (mean angle 151.4°), unpaired actine shorter than paired or equal (mean length: 68.8 µm – unpaired, 80.7 µm – paired, 22.9 µm – apical) (Table 5). All actines equal in width (mean width: 5.6  $\mu$ m – unpaired, 5.8  $\mu$ m – paired, 5.5  $\mu$ m – apical) (Table 5). Apical actine curved and smooth.
- Skeleton: Skeleton of oscular tubes predominantly formed by both tri- and tetractines, while in cormus tubes tetractines rare (Fig. 7C, D). In oscular tubes, spicules constitute organized array with their unpaired actines directed toward cormus and oriented more or less in parallel to proximo-distal axis of oscular tube (Fig. 7C). In cormus tubes, spicule network appears completely disordered (Fig. 7D). Both populations of diactines forming small oscular crown up to 60 µm and

512 cover tubes surface, orienting in different directions and extending outside by lance-shaped tip513 (Fig. 7B).

*Cytology:* Body wall, 8.4-12 μm thick, three layers: exopinacoderm, loose mesohyl, choanoderm
515 (Fig. 9A, B; Table S2). Flat endopinacocytes located only in the distal part of oscular tube (oscular
516 ring) replacing choanocytes. Inhalant pores scattered throughout exopinacoderm, except the
517 oscular ring area.

518 Exopinacocytes non-flagellated T-shaped, rarely flat (Fig. 9C). External surface covered by
519 glycocalyx. Cell body (height 7-10.5 μm, width 4.3-5.5 μm), containing spherical to oval nucleus
520 (diameter 3.1 μm), submersed in mesohyl. Cytoplasm with specific spherical electron-dense
521 inclusions (0.2-0.4 μm diameter) (Fig. 9C).

522 Endopinacocytes non-flagellated flat cells, size 16.8  $\mu$ m × 2.2  $\mu$ m (Fig. 9D). External surface 523 covered by glycocalyx. Nucleus (2.1  $\mu$ m × 1.6  $\mu$ m) spherical to oval with nucleolus. Cytoplasm 524 with specific spherical electron-dense inclusions (0.2-0.5  $\mu$ m diameter) (Fig. 9D).

Choanocytes flagellated trapeziform or prismatic (height 8.2 µm, width 4.1 µm) (Fig. 9E). Flagellum surrounded by collar of microvilli. Characteristic pyriform nucleus (2.6  $\mu$ m × 4.1  $\mu$ m) in apical position. Cytoplasm with phagosomes and small vacuoles. Choanocytes united by specialized intercellular contacts similar to septate junctions, but has no basal membrane (Fig. 9E). Porocytes tubular cylindrical (height 5.5 µm, width 4.2 µm), connecting external milieu with choanocyte tube (Fig. 9F). Nucleus pyriform (diameter 3.1 µm), containing nucleolus. Cytoplasm with phagosomes, small vacuoles and spherical electron-dense inclusions identical with inclusions of exopinacocytes. 

533 Sclerocytes amoeboid, size 8.7  $\mu$ m × 3.5  $\mu$ m (Fig. 10A). Nucleus usually oval or pear-shaped 534 (diameter 2.5  $\mu$ m), containing single nucleolus. Well-developed Golgi apparatus and rough 535 endoplasmic reticulum. Cytoplasm usually with phagosomes and/or lysosomes. During spicules 536 secretion, sclerocytes form groups of three to six cells, connected by septate junctions (Fig. 10A). 537 Amoebocytes of different shape (from oval to amoeboid) without special inclusions, size 5.7

 $\mu$ m × 4.7  $\mu$ m (Fig. 10B). Nucleus spherical (diameter 2.9  $\mu$ m), sometimes with nucleolus.

Granular cells oval, size 9  $\mu$ m × 5.5  $\mu$ m. Regularly distributed numerous cells, usually located under choanocytes (Fig. 10C-E). Nucleus in peripheral position, spherical (diameter 2.5  $\mu$ m). Cytoplasm with oval electron dense inclusions (size 0.9-2.7  $\mu$ m × 1.1-3.7  $\mu$ m) (Fig. 10E). Inclusion content homogenous or granulated. Often found in stage of degradation, cytoplasm completely filled with 2-4 large oval inclusions with highly osmiophilic granulated content (Fig. 10F).

545 Myocytes are fusiform cells, size 22  $\mu$ m × 2.7  $\mu$ m; distributes in the mesohyl mostly in the 546 oscular ring. Nucleus usually oval (2.9  $\mu$ m × 1.6  $\mu$ m), without nucleolus (Fig. 10G). Cytoplasm 547 includes mitochondria, ribosomes, small vesicles and, most importantly, the presence of 548 cytoplasmic myofilaments of 19-12 nm in diameter (Fig. 10G). Myofilaments form bundles (0.37-549 0.16  $\mu$ m diameter) that located along the long axis of the cell.

One morphotype of bacterial symbionts in mesohyl. Bacteria numerous, rod-shaped with
double cell wall, diameter 0.3-0.33 µm, length 3.0-5.6 µm (Fig. 10H). Nucleoid region electrondense with irregular network of filaments.

*Distribution:* Boreal-Arctic species. Molecular identity confirmed for Greenland and the White Sea (Alvizu *et al.*, 2018). In the White Sea it is the most abundant species, inhabiting kelps and hard substrates in low intertidal and subtidal zones up to 15-20 m depth.

*Reproduction:* In the White Sea specimens collected in late October contained early oocytes,
specimens collected in January-February contained fully-developed larvae.

*Remarks:* In the White Sea, this species was initially identified as *Leucosolenia variabilis*, based on its external morphology (Lavrov *et al.*, 2018). In addition, most of our sequences for this species were identical to LSU and SSU sequences downloaded from the GenBank under the name *L. variabilis*. Regarding morphology, the spicular characters of our specimens were different from the original description of *L. variabilis* (Haeckel, 1872), but partly fit the description given in Minchin (1904). The main differences relate to diactine morphology: in our specimens, there is a

single type of curved lanceolate diactines. In L. variabilis sensu Haeckel (1872), two diactine populations were found: the first is small strait trichoxea, and the second is normal curved lanceolate diactines. Minchin (1904) found connectivity in size among small and long diactines and suggested they represented a single type of diactines, which was overlooked by Haeckel. Since our specimens possess only a single diactine population, it might support Minchin's conclusions. However, Leucosolenia variabilis sensu Minchin (1904) is a species complex since he designated Leucosolenia somesii a junior synonym of L. variabilis, while morphological and molecular data supported its identity as a distinct species (see below, see also van Soest et al., 2007). Therefore, the diagnosis provided by Minchin (1904) should not be taken into consideration. 

To address the possible ontogenetic variation of diactines, we studied the type material L. *variabilis* from the collection of BMNH (syntype BMNH-1910.1.1.421). The spicular characters of this specimen perfectly fit the original description made by Haeckel (1872), with two diactine types, tri- and tetractines of equal abundance and unpaired actine in tri- and tetractines always shorter than paired ones. On the other hand, specimens in our material possess only a single diactine type, and tetractines are rare. Therefore, the species from the White Sea is not L. variabilis, despite its molecular similarity to specimens, placed in the GenBank under the name L. variabilis. Another species, that is characterised by a single diactine type and short unpaired actines in tri- and tetractines is *Leucosolenia corallorrhiza*, which was designated a valid species in the most recent morphology-based revision of Greenland calcareous sponges (Rapp, 2015). Haeckel (1872) described this species under the name Ascortis corallorrhiza, addressing a small proportion or absence of tetractines, small and thick triactines with short unpaired actine. Diactines are curved lance-shaped (Haeckel, 1872:74). This feature is characteristic of samples from the White Sea, although in our specimens, some diactines bear small spines on their lance-shaped tips. These spines are hardly visible with light microscopy and may be overlooked even during SEM studies. Since we could not study the morphology of specimens whose sequences were obtained from GenBank, and morphological data for those specimens are absent in the respective paper (Alvizu

et al., 2018), we designate our specimens from the White Sea as Leucosolenia corallorrhiza, until both morphological and molecular conformation for specimens from the type localities would be available. Also, neotype designation for this species is necessary to establish the type material, specimens for this purpose should be collected in the type locality. It should be mentioned, that our specimens demonstrate minor differences in colouration from the original description (L. corallorrhiza is brown according to Haeckel, 1872). Also, actines in tri- and tetractines are thicker in initial description (widths  $\sim 15 \text{ }\mu\text{m}$  in Haeckel, 1872; up to 12.5  $\mu\text{m}$  in our material (Table 5); up to 10.7 µm in Rapp, 2015), but this difference may be associated either with ontogenetic or intraspecific variation, or different measurement procedures and equipment. 

From *Leucosolenia variabilis* this species differs by spicular characters: in *L. variabilis*, there are two types of diactine, while there is only one type of diactines in *L. corallorrhiza*. *Leucosolenia corallorrhiza* never forms a large, massive cormus. *Leucosolenia corallorrhiza* also differs from other species in cytological characteristics (Table S2): in contrast to *L. complicata*, the mesohyl of *L. corallorrhiza* includes not only amoeboid cells but also numerous granular cells, regularly distributed in the body wall; in contrast to *L. variabilis*, *L. corallorrhiza* has larger granular cells, no spherulous cells and only one morphotype of rod-shaped symbiotic bacteria.

Leucosolenia variabilis Haeckel, 1870

607 (Figs 11-16; Table 6)

*Type material:* Syntype BMNH-1910.1.1.421. Other type material is not known.

*Type locality:* Norway, Bergen.

*Material studied:* 40 specimens. Molecular data – 40 specimens, external morphology – 40

611 specimens, skeleton organisation – 3 specimens (WS11643, WS11708, WS11735), spicules (light

612 microscopy, SEM) – 7 specimens (WS11707, WS11714, WS11731, WS11732, WS14637,

**613** WS14671, WS14681), cytology (TEM) – 3 specimens (WS11643, WS11644, WS11645) (Table

<sup>′</sup> 614

S1).

External morphology: Length of cormus up to 5 cm. Cormus massive, often spherical, otherwise formed by basal reticulation of tubes. Cormus built as reticulation around one or several largest central tubes. Outline of cormus formed by numerous short diverticula. Largest tubes of cormus always end with oscula. Main oscular tubes large prominent erect, bearing many small diverticula, spreading to 2/3 of tubes length. Oscular tube gradually narrows to oscular rim, possessing short spicular crown (Fig. 11A, B). In addition to main oscula on largest tubes of cormus, smaller oscular tubes usually scattered all over the cormus. Surface minutely hispid. Colouration of living specimens gravish white. Colouration of preserved specimens from gravish white to ochre (Fig. 11A). 

*Spicules:* Diactines (Figs 12A, 13, 16A). Two populations: (1) Curved smooth lanceolate diactines 625 (Fig. 12A), mean length 306.7  $\mu$ m, mean width 9.8  $\mu$ m, (Table 6). Slightly curved, smooth, with 626 lanceolate outer tip, variable in length. (2) Trichoxeas (Fig. 13). Thin (mean width 0.9  $\mu$ m) (Table 627 6), with numerous irregularly distributed spines (Fig. 13C), long, but usually represented by 628 fragments of variable length (up to 362.4  $\mu$ m long) (Table 6).

Triactines (Figs 12B, C, 16A). Predominantly T-shaped sagittal (mean angle 138.5°), unpaired actine variable in length: most frequently equal to paired actines, commonly shorter or rarely longer than paired (mean length: 122.3  $\mu$ m – unpaired, 128.0  $\mu$ m – paired) (Table 6). Abnormal triactines with one of paired actines undulated also common (Fig. 12C). Actines equal in width (mean width: 8.1  $\mu$ m – unpaired, 8.5  $\mu$ m – paired) (Table 6).

Tetractines (Fig. 12D). Predominantly T-shaped sagittal (mean angle 142.2°), unpaired actine variable in size: equal to, shorter or longer than paired actines (mean length: 147.6 µm -unpaired, 143.0 µm – paired, 22.8 µm – apical) (Table 6). Unpaired actine usually slightly slender than paired (mean width: 8.5  $\mu$ m – unpaired, 9.1  $\mu$ m – paired) (Table 6). Apical actine curved, smooth and slender (mean width  $5.9 \,\mu\text{m}$ ) (Table 6). 

57 639 *Skeleton:* Skeleton of both oscular and cormus tubes formed by dense net of tetractines and
 58 640 triactines (Fig. 11C, D). In oscular tubes spicules constitute organized array with their unpaired

actines directed toward cormus and oriented more or less in parallel to proximo-distal axis of oscular tube (Fig. 11C). In cormus tubes spicule array completely disordered (Fig. 11D). Diactines form small oscular crown up to 100  $\mu$ m (Fig. 11B) and cover tubes surface, orienting in different directions and extending outside by lance-shaped tip.

*Cytology:* Body wall, 9-13.8 μm thick, three layers: exopinacoderm, loose mesohyl, choanoderm
646 (Fig. 14A, B; Table S2). Flat endopinacocytes located only in distal part of oscular tube (oscular
647 ring) replacing choanocytes. Inhalant pores scattered throughout exopinacoderm, except the
648 oscular ring area.

Exopinacocytes non-flagellated T-shaped, rarely flat (Fig. 14C). External surface covered
by glycocalyx. Cell body (height 6.3 μm, width 3.7 μm), containing spherical to oval nucleus
(diameter 2.7 μm), submersed in mesohyl. Cytoplasm with specific spherical electron-dense
inclusions (0.2-0.35 μm diameter) (Fig. 14C).

Endopinacocytes non-flagellated flat cells, size 16  $\mu$ m × 2.8  $\mu$ m. External surface covered by glycocalyx. Nucleus (3.2  $\mu$ m × 2.3  $\mu$ m) spherical to oval with nucleolus. Cytoplasm without specific inclusions (Fig. 14F).

656 Choanocytes flagellated trapeziform or prismatic (height 10.7 μm, width 4.1 μm) (Fig. 14D).
657 Flagellum surrounded by collar of microvilli. Characteristic pyriform nucleus (diameter 2.5 μm)
658 in apical position. Cytoplasm with phagosomes and small vacuoles (Fig. 14D).

659 Porocytes tubular cylindrical (height 2.5-4.7 μm, width 4.3-5 μm), connecting external 660 milieu with choanocyte tube (Fig. 14E). Nucleus spherical (diameter 2.7 μm), containing 661 nucleolus. Cytoplasm with spherical electron-dense inclusions identical with inclusions of 662 exopinacocytes (Fig. 14E).

663 Sclerocytes amoeboid, size  $6 \ \mu m \times 3.1 \ \mu m$  (Fig. 15G). Nucleus usually oval or pear-shaped 664 (diameter 2.2  $\ \mu m$ ), containing single nucleolus. Well-developed Golgi apparatus and rough 665 endoplasmic reticulum. Cytoplasm usually with phagosomes and/or lysosomes (Fig. 15G).

666 Amoebocytes of different shape (from oval to amoeboid) without special inclusions, size 3

 $\mu m \times 4-7.5 \ \mu m$  (Fig. 15A). Nucleus spherical (diameter 2.7  $\mu m$ ), sometimes with nucleolus.

Large amoeboid cells of different shape (from elongate to amoeboid), size  $20 \ \mu m \times 4.2 \ \mu m$ (Fig. 15B). Rare cells located under choanoderm. Nucleus oval (size  $4.8 \ \mu m \times 1.7 \ \mu m$ ). Cytoplasm with numerous large heterophagosomes (diameter 1.1-3.2 \ \mu m), well-developed Golgi apparatus (Fig. 15B).

Granular cells small oval, size 4  $\mu$ m × 3.3  $\mu$ m (Fig. 15C). Rare cell type, located under the choanoderm. Nucleus in peripheral position, spherical (diameter 1.7  $\mu$ m) with large amounts of heterochromatin, associated with nucleus membrane Cytoplasm with electron dense oval inclusions (size 0.7-6  $\mu$ m × 0.4-1.1  $\mu$ m) and rare spherical electron-transparent vacuoles (diameter 1.2  $\mu$ m) (Fig. 15C).

577 Spherulous cells with irregular shape from amoeboid to crescent, size 2.7-9.2 μm × 4.7-5.3 578 μm (Fig. 15E-G). Regularly distributed numerous cells, usually located under choanocytes. 579 Distance between cells 2-9 μm (Fig. 15F). Nucleus deformed (size 2.4 μm × 1.7 μm). Cytoplasm 580 mostly occupied by large crescent or irregular electron-dense homogenous inclusions (diameter 581 1.8-4.5 μm) and less electron-dense fine-granular inclusions (diameter 0.7-2.6 μm). Granular or 582 foamy material fills cytoplasm spaces between inclusions (Fig. 15E, G).

683 Myocytes rare fusiform cells, size  $18 \ \mu m \times 2.7 \ \mu m$ , located in mesohyl (Fig. 15D). Nucleus 684 oval (3.5  $\mu m \times 2.7 \ \mu m$ ), with nucleolus. Cytoplasm with mitochondria, ribosomes, small vesicles 685 and cytoplasmic myofilaments. Myofilaments grouped in bundles (diameter 0.07-0.2  $\mu m$ ) located 686 along long axis of myocyte (Fig. 15D).

687 Three morphotypes of bacterial symbionts in mesohyl (Fig. 15H-J). Morphotype 1 numerous
688 (Fig. 15H). Bacteria large spiral-shaped, diameter 0.2 μm, length 2.5-3.9 μm. Spiral turns regular
689 and compact. Single membrane cell wall, cytoplasm granular, nucleoid region tubular (Fig. 15H).
690 Morphotype 2 rare (Fig. 15I). Bacteria small spiral-shaped, diameter 0.3 μm, length 1.5-1.8
691 μm. Spiral turns irregular and sparse. Cytoplasm transparent, nucleoid region tubular (Fig. 15I).

692 Morphotype 3 rare (Fig. 15J). Bacteria small rod-shaped bacteria, diameter 0.23 μm, length

693 0.8 μm. Double membrane cell wall, cytoplasm with dark filamentous materials, no distinction
694 between cytoplasm and nucleoid region (Fig. 15J).

*Distribution:* Boreal-Arctic species, described from Norway. Molecular identity confirmed for the

696 White Sea and Greenland (Alvizu et al., 2018). In the White Sea occurs in low intertidal and

697 subtidal zones up to 40-45 m depth, on rocks and kelps.

*Reproduction:* No data about reproduction time for this species.

*Remarks:* We studied three type specimens (slides with spicules) of *Leucosolenia variabilis* from

the British Museum of Natural History (BMNH): BMNH-1910.1.1.421, BMNH-1906.12.1.40 and

701 BMNH-1906.12.1.50. Spicules are similar morphologically across these specimens (Fig. 16B-D),

which supports the idea that they belong to the same species. At the same time, their type status should be reconsidered due to the data represented in the revision by Minchin (1904). Slide labels contain specific information (exact page and number), allowing an unambiguous comparison with the collection data of these samples given in Minchin (1904). Accordingly, BMNH-1906.12.1.50 was collected from Bantry Bay, Ireland, by C. Norman and identified by him as Leucosolenia botryoides; this label was endorsed by Haeckel 'Ascandra variabilis' (slide no. 1; Minchin, 1904:385). BMNH-1906.12.1.40 was received by Haeckel for reexamination from Bowerbank and collected from Guernsey (slide no. 4; Minchin, 1904:385). Finally, BMNH-1910.1.1.421 was collected by Haeckel in Bergen, Norway, the type locality of this species, and contained a printed label 'Ascandra variabilis H.' (slide no. 3; Minchin, 1904:385). Therefore, the slide BMNH-1910.1.1.421 could be designated as a syntype.

The analysis of *L. variabilis* syntype BMNH-1910.1.1.421 indicated two diactine types (lanceolate diactines and trichoxeas), and V- and T-shaped tri- and tetractines with shorter unpaired actines (Fig. 16D). Although Haeckel's description lacks long trichoxeas, it should be mentioned that such spicules are easily broken during preparation. It may also be suggested that the second type of diactines without lanceolate tips described by Haeckel (1872) is in fact broken

718 long trichoxeas. Direct comparison of spicule slides of specimens from the White Sea with *L*.

*variabilis* syntype BMNH-1910.1.1.421 shows strong correspondence between them.

Leucosolenia variabilis has a large, massive, sometimes spherical cormus, which could be a good distinctive trait, since all other sympatrically living species (Leucosolenia complicata, L. corallorrhiza, Leucosolenia sp. A) are represented by basal reticulation of tubes with extended oscular tubes. In spicular characters, L. variabilis differs from L. somesii by the presence of lanceolate spined diactines; and from L. complicata and Leucosolenia sp. A by the presence of extremely long and highly spined trichoxeas. Leucosolenia variabilis also has the highest diversity of mesohyl cells and symbiotic bacteria among the studied *Leucosolenia* species (Table S2). In addition to the usual amoebocytes, L. variabilis also has rare large amoebocytes and small granular cells, as well as numerous unusual spherulous cells of different shapes regularly distributed in the body wall. The composition of symbiotic bacteria of L. variabilis includes three morphotypes: one typical rod-shaped and two unusual spiral-shaped. 

*Leucosolenia* sp. A

733 (Figs 17-18; Table 7)

Material studied: Three specimens. Molecular data – 3 specimens (WS11692, WS11752,
WS11770), external morphology – 3 specimens (WS11692, WS11752, WS11770), skeleton
organisation – 2 specimens (WS11752, WS11770), spicules (SEM) – 2 specimens (WS11692,
WS11770) (Table S1).

*External morphology:* Studied specimens small. Length of cormus up to 1 cm. Cormus represented
by compact reticulation of tubes, from which several oscular tubes arising. Oscular tubes erect and
almost straight. Surface minutely hispid. Colouration of living and preserved specimens grayish
white (Fig. 17A).

*Spicules:* Diactines (Fig. 18A-C). Two populations: (1) Curved spiny lanceolate diactines (Fig.
743 18B), mean length 189.1 μm, mean width 7.2 μm (Table 7). Small, from almost straight to slightly

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curved and undulating, with lanceolate and spiny outer tip, spines in distinct rows (Fig. 18C). (2)
Curved smooth diactines (Fig. 18A), mean length 515.0 µm, mean width 11.6 µm (Table 7). Rare,
long, slightly curved, without spines and lanceolate tips (Fig. 18C).

Triactines (Fig. 18D, E). Predominantly T-shaped sagittal (mean angle 146.5°) (Table 7). Unpaired actine variable in size: equal to, shorter or longer than paired actines, but shorter unpaired actine most common (mean length: 118.5  $\mu$ m – unpaired, 125.1  $\mu$ m – paired) (Table 7). Both straight and bent paired actines common. Abnormal triactines in high number (Fig. 18E), sometimes with undulated actines. Unpaired actine usually slightly slender than paired (mean width: 11.1  $\mu$ m – unpaired, 11.6  $\mu$ m – paired) (Table 7).

Tetractines (Fig. 18F). Quite rare. Predominantly T-shaped (mean angle 140.8°) (Table 7), variable in size. Unpaired actine equal to paired ones (mean length: 114.3  $\mu$ m – unpaired, 113.2  $\mu$ m – paired, 30.0  $\mu$ m – apical) (Table 7). Unpaired actine straight, paired actines straight or undulating, apical actine curved or undulating, smooth. Paired and unpaired actines equal in width, apical actine more slender (mean width: 8.6  $\mu$ m – unpaired, 8.5  $\mu$ m – paired, 7.1  $\mu$ m – apical) (Table 7).

*Skeleton:* Skeleton of both oscular rim and cormus tubes predominantly formed by triactines,
tetractines rare (Fig. 17C, D). In oscular tubes spicules constitute organized array with their
unpaired actines directed toward cormus and oriented more or less in parallel to proximo-distal
axis of oscular tube (Fig. 17C). In cormus tubes spicule array completely disordered (Fig. 17D).
Prominent oscular crown absent (Fig. 17B). Both populations of diactines cover tubes surface,
orienting in different directions and extending outside.

*Cytology:* No material was available for cytological studies.

*Distribution:* Arctic species. Molecular identity confirmed only for the White Sea and Greenland.

Found subtidal up to 15 m on rocks and red algae.

*Reproduction:* No data about reproduction time for this species.

Remarks: Although both our species delimitation analysis based on the H3 dataset and morphological data suggest that this species represents a distinct species-level unit, we avoid describing a new species as this case requires additional studies for several reasons. *Leucosolenia* sp. shares some features with *Leucosolenia corallorrhiza*: (1) the external appearance is similar, (2) the angle between unpaired actines in tri- and tetractines is similar (the mean angle is 142.9° in *L. corallorrhiza* and 146.5° in *Leucosolenia* sp. A), (3) the unpaired actine in tri- and tetractines is commonly shorter than the unpaired ones. However, these two species show several differences. Firstly, Leucosolenia sp. A has two populations of diactines, the smaller with lanceolate tips and the rare large curved smooth non-lanceolate diactines, whereas in L. corallorrhiza only the first type is present. Also, the tetractines are rare in both the cormus and oscular regions of *Leucosolenia* sp. A, while they are commonly present in the osculum of *L. corallorrhiza. Leucosolenia* sp. A commonly has triactines with bent unpaired actines, which are straight in *L. corallorrhiza*. Finally, the mean length of actines in tri- and tetractines of L. corallorrhiza is shorter than those of Leucosolenia sp. A (L. corallorrhiza: 70.5 µm – unpaired actine mean length, 82.7 µm – paired actine mean length; Leucosolenia sp. A: 118.5 µm – unpaired mean length, 125.1 µm – paired mean length). At the same time, the limited material of *Leucosolenia* sp. A (only three specimens were collected and studied) does not allow us to study the possible interspecific variation and ontogenetic variation. Therefore, we avoid the designation of this species as a distinct one, until more material would be available for study.

*Leucosolenia creepae* sp. nov.

790 (Figs 19-22; Table 8)

ZooBank LSID: urn:lsid:zoobank.org:act:D60461BE-F215-4BE2-AFB5-25AF542FC4B9 *Type material: Holotype:* WS11702, White Sea, Kandalaksha Bay, Velikaya Salma Strait, vicinity
of the N.A. Pertsov White Sea Biological Station, 0-2 m depth, 28.viii.2018, coll. A.I. Lavrov. *Paratypes:* WS11703, 1 specimen, White Sea, Kandalaksha Bay, Velikaya Salma Strait, vicinity

of the N.A. Pertsov White Sea Biological Station, 0-2 m depth, 28.viii.2018, coll. A.I. Lavrov. WS11728 paratype agrees in locality, date and collector with holotype WS11702 and paratype WS11703. WS11655 was collected in 30.viii.2017, and WS11725, WS11726, WS11771 were collected in 24.viii.2018, but all agree in locality and collector with holotype WS11702 and paratype WS11703 Type locality: White Sea, Kandalaksha Bay, Velikaya Salma Strait, vicinity of the N.A. Pertsov White Sea Biological Station (66°34'N, 33°08'E). Material studied: 54 specimens. Molecular data - 54 specimens, external morphology - 54 specimens, skeleton organisation - 3 specimens (WS11655, WS11728, WS11762), spicules (SEM) – 5 specimens (WS11579, WS11605, WS11704, WS11729, WS11775), cytology (TEM) - 3 specimens (WS11579, WS11600, WS11698) (Table S1). 

*Etymology:* From English "creep", referring to specific decumbent cormus and unusual growth
form of this species in contrast to sympatrically living *Leucosolenia corallorrhiza*.

*External morphology:* Length of cormus up to 5 cm. Cormus formed by basal reticulations of
creepy tubes with one or several oscular tubes (Fig. 19A). Tubes brittle. Oscular tubes creepy
slightly curved and erect at distal end, sometimes with few diverticula. Oscular rim gradually
narrows, possessing prominent spicular crown (Fig. 19A). Surface echinate. Colouration of living
and preserved specimens grayish white (Fig. 19A).

*Spicules:* Diactines (Fig. 20A, B). Spiny diactines, mean length 194.9 μm, mean width 5.1 μm
814 (Table 8). Extremely variable in length, without lanceolate tips, spiny. Largest diactines slightly
815 curved, intermediate and short diactines straight. Spines in distinct rows at one end of diactines,
816 more or less reduced in large ones (Fig. 20B).

 $^{52}$ 817Triactines (Fig. 20C, D). Sagittal, T-shaped and V-shaped (mean angle 131.1°) (Table 8), $^{54}$ usually recurved, unpaired actine variable in length: most frequently shorter then paired actines, $^{56}$ but equal and longer unpaired actines rarely occur (mean length: 80.7 µm – unpaired, 94.9 µm – $^{59}$ 820 $^{50}$ paired) (Table 8). Aberrant T- and V-shaped triactines present, sometimes with undulated rays

821 (Fig. 20D). Unpaired actine often more slender than paired actines (mean width: 5.4 μm –
822 unpaired, 5.9 μm – paired) (Table 8).

 823Tetractines (Fig. 20E). Quite rare. Sagittal, T-shaped and V-shaped (mean angle 139.5°)824(Table 8), variable in size and proportions. Unpaired actine variable in length: longer, shorter and825equal to unpaired actines (mean length: 85.1 µm – unpaired, 95.3 µm – paired, 25.6 µm – apical)826(Table 8). Paired and unpaired actines equal in width (mean width: 6.2 µm – unpaired, 6.3 µm –827paired) (Table 8). Apical actine curved, smooth and slender (mean width 5.2 µm) (Table 8).

*Skeleton:* Skeleton of oscular rim predominantly formed by both tri- and tetractines, while in other body parts tetractines absent (Fig. 19C, D). In oscular tubes spicules constitute organized array with their unpaired actines directed toward cormus and oriented more or less in parallel to proximo-distal axis of oscular tube (Fig. 19C). In cormus tubes spicule array completely disordered (Fig. 19D). Diactines forms extending oscular crown up to 500 μm (Fig. 19B) and cover tubes surface in large number, orienting in different directions and making it hispid.

*Cytology:* Body wall, 9-14 μm thick, three layers: exopinacoderm, loose mesohyl, choanoderm
(Fig. 21A, B; Table S2). Flat endopinacocytes located only in distal part of oscular tube (oscular
ring) replacing choanocytes. Inhalant pores scattered throughout exopinacoderm, except the
oscular ring area.

Exopinacocytes non-flagellated T-shaped, rarely flat (Fig. 21C). External surface covered
by glycocalyx. Cell body (height 5.8 μm, width 2.8 μm), containing spherical to oval nucleus
(diameter 2.7 μm), submersed in mesohyl. Cytoplasm with specific spherical electron-dense
inclusions (0.25-0.35 μm diameter) (Fig. 21C).

842 Endopinacocytes non-flagellated flat cells, size 16.2  $\mu$ m × 2.7  $\mu$ m (Fig. 21D). External 843 surface covered by glycocalyx. Nucleus (diameter 2.7  $\mu$ m) spherical without nucleolus. Cytoplasm 844 without specific inclusions (Fig. 21D).

57 845 Choanocytes flagellated trapeziform or prismatic (height 11.4 μm, width 3.6 μm) (Fig. 21E).
 58 59 60 846 Flagellum surrounded by collar of microvilli. Characteristic pyriform nucleus (diameter 2.3 μm)
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| 1              |     |   |
|----------------|-----|---|
| 2              | 847 | in apical position. Cytoplasm with phagosomes and small vacuoles (Fig. 21E).                                  |
| 4<br>5         | 848 | Porocytes tubular cylindrical (height 4.6-8.9 µm, width 2.8-2.9 µm), connecting external                      |
| 6<br>7         | 849 | milieu with choanocyte tube (Fig. 21F). Nucleus oval to spherical (diameter 2.5 $\mu$ m), sometime            |
| 8<br>9         | 850 | with nucleolus. Cytoplasm with spherical electron-dense inclusions identical with inclusions of               |
| 10<br>11<br>12 | 851 | exopinacocytes, phagosomes and small vacuoles (Fig. 21F).   |
| 13<br>14       | 852 | Sclerocytes amoeboid, size 7.6 $\mu$ m × 2.9 $\mu$ m (Fig. 22A). Nucleus usually oval or pear-shaped          |
| 15<br>16       | 853 | (diameter 2.3 $\mu$ m), sometimes with single nucleolus. Well-developed Golgi apparatus and rough             |
| 17<br>18       | 854 | endoplasmic reticulum. Cytoplasm usually with phagosomes and/or lysosomes (Fig. 22A).                         |
| 19<br>20<br>21 | 855 | Amoebocytes of different shape (from oval to amoeboid) without special inclusions, size 5.7                   |
| 22<br>23       | 856 | $\mu$ m × 2.6 $\mu$ m (Fig. 22B). Nucleus spherical (diameter 2.5 $\mu$ m), sometimes with nucleolus.         |
| 24<br>25       | 857 | Myocytes fusiform cells, size 16.5 $\mu$ m × 3.3 $\mu$ m, located in mesohyl. Nucleus oval (2.4 $\mu$ m       |
| 26<br>27       | 858 | $\times$ 1.9 µm), without nucleolus (Fig. 21D). Cytoplasm with mitochondria, ribosomes, small vesicles        |
| 28<br>29<br>30 | 859 | and cytoplasmic myofilaments. Myofilaments grouped in bundles (diameter 0.25–0.32 µm) that                    |
| 31<br>32       | 860 | are located along the long axis of the cell (Fig. 21D).   |
| 33<br>34       | 861 | Two morphotypes of bacterial symbionts in mesohyl. Morphotype 1 numerous (Fig. 22C,                           |
| 35<br>36<br>37 | 862 | D). Bacteria large rod-shaped slightly curved, diameter 0.4-0.5 μm, length 2.7 μm. Cell is double,            |
| 38<br>39       | 863 | smooth and covered with fibers, cytoplasm transparent with vacuolar inclusions, nucleoid region               |
| 40<br>41       | 864 | filamentous (Fig. 22C, D).  |
| 42<br>43       | 865 | Morphotype 2 abundant (Fig. 22E, F). Bacteria small rod-shaped bacteria, diameter 0.19 µm.                    |
| 44<br>45<br>46 | 866 | length 1 9-2 1 µm Cell wall smooth cytoplasm transparent nucleoid region filamentous (Fig                     |
| 40<br>47<br>48 | 867 | 22E F)  |
| 49<br>50       | 868 | <i>Distribution</i> : Arctic species. In the White Sea quite rare, found in low intertidal and upper subtidal |
| 51<br>52       | 869 | zones up to 5-10 m denth, on kelns and rocks  |
| 53<br>54       | 870 | <i>Reproduction</i> : In White See specimens, collected in mid. June to the end of the August contained       |
| 55<br>56<br>57 | 070 | neproduction. In write Sea specificity, confected in find-June to the end of the August contained             |
| 5,             | ŏ/⊥ | oucytes at uniferent stages (mostly, early) of development (Figs 21B, 22G).                                   |

Remarks: Leucosolenia creepae sp. nov. well differs from other Leucosolenia species in both external characters and the morphology of spicules. In L. somesii, diactines are of two types: (1) smooth diactines, which are variable in length and (2) short and highly spined ones. In Leucosolenia creepae sp. nov., we identified only one type of diactines, which have spines on the outer tip and are variable in length. However, in *Leucosolenia creepae* sp. nov., spines are more expressed in small and medium-size diactines but become hardly visible in longer diactines of ~250-300 µm in length. In L. somesii, all medium-size diactines have smooth tips (Fig. 24B). Leucosolenia creepae sp. nov. forms a sparse basal reticulation with few oscular tubes, while the cormus of *L. somesii* is formed by a dense reticulation of extremely branched winding tubes. From all other North Atlantic and Arctic Leucosolenia species, Leucosolenia creepae sp. nov. differs by the absence of lanceolate diactines. The mesohyl cell composition of *Leucosolenia creepae* sp. nov, includes only amoebocytes, myocytes and sclerocytes, differing this species from sympatrically living *L. corallorrhiza* and *L. variabilis* (Table S2). 

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- *Leucosolenia somesii* (Bowerbank, 1874)
- 887 (Figs 23-24; Table 9)

38
 39
 888 *Type material:* Lectotype BMNH 1925.11.2.24, paralectotype BMNH 1925.11.2.25, slides of the
 40
 41
 889 same: BMNH 1956.4.26.35.

*Type locality:* Brighton Aquarium.

*Material studied:* 1 specimen, ZMA Por. 17572 (external morphology, skeleton organisation,
892 spicules) (Table S1).

External morphology: Length up to 12 cm. Cormus formed by dense reticulation of extremely branched winding tubes (Fig. 23A). Surface hispid. Colouration of living and preserved specimens gravish white. Examined specimen lacks oscular tubes. According to the original description (Bowerbank, 1874), sponge have numerous small and large oscular tubes, bearing spicular crown.

<sup>59</sup> 897 Oscular tubes erect and slightly curved, gradually narrowing to oscular rim.

1

| 2                        | 898 | Spicules: Diactines (Fig. 24A-C). Two populations of diactines: (1) Curved smooth diactines (Fig.            |
|--------------------------|-----|--|
| 4<br>5                   | 899 | 24A, C), mean length 425.5 $\mu$ m, mean width 9.9 $\mu$ m (Table 9). Slightly curved, smooth, variable      |
| 6<br>7                   | 900 | in length, lacking lanceolate tips, with undulated tip. (2) Straight spiny diactines (Fig. 24B, C),          |
| 8<br>9<br>10<br>11<br>12 | 901 | mean length 90.0 $\mu$ m, mean width 3.3 $\mu$ m (Table 9). Short, strait, lacking lanceolate tips, with     |
|                          | 902 | numerous spines in distinct rows (Fig. 24B).   |
| 13<br>14                 | 903 | Triactines (Fig. 24D, E). Sagittal, mostly T-shaped, but V-shaped also occurs (mean angle                    |
| 15<br>16<br>17           | 904 | 131.7°) (Table 9), unpaired actine usually shorter then paired actines, but longer unpaired actines          |
| 17<br>18<br>19           | 905 | rarely occur (mean length: 127.4 $\mu$ m – unpaired, 155.2 $\mu$ m – paired) (Table 9). Paired and unpaired  |
| 20<br>21                 | 906 | actines equal in width (mean width: 8.2 $\mu$ m – unpaired, 8.0 $\mu$ m – paired) (Table 9). Abnormal        |
| 22<br>23                 | 907 | triactines in common, sometimes with undulated rays (Fig. 24E).  |
| 24<br>25<br>26           | 908 | Tetractines (Fig. 24F). Quite rare. Sagittal, mostly T-shaped, but V-shaped also occurs                      |
| 27<br>28                 | 909 | (mean angle 139.3°) (Table 9), unpaired actine usually shorter then paired actines, but longer               |
| 29<br>30                 | 910 | unpaired actines rarely occur (mean length: 156.0 $\mu$ m – unpaired, 178.7 $\mu$ m – paired, 21.5 $\mu$ m – |
| 31<br>32<br>33           | 911 | apical) (Table 9). Apical actine curved and smooth. All actines more or less equal in width (mean            |
| 34<br>35                 | 912 | width: 9.1 $\mu$ m – unpaired, 9.5 $\mu$ m – paired, 10.0 $\mu$ m – apical) (Table 9).                       |
| 36<br>37                 | 913 | Skeleton: Very dense net predominantly formed by triactines, oriented in different directions,               |
| 38<br>39<br>40           | 914 | tetractines rare (Fig. 23B). Both trichoxea populations cover tubes surface in large number,                 |
| 40<br>41<br>42           | 915 | orienting in different directions and making it hispid. Skeleton of osculum was not studied.                 |
| 43<br>44                 | 916 | Cytology: No material was available for cytological studies.   |
| 45<br>46                 | 917 | Distribution: Boreal species. Described from Brighton Aquarium with confirmed reports from                   |
| 47<br>48<br>49           | 918 | Netherlands (van Soest et al., 2007). Probably it has wider distribution in North-East Atlantic.             |
| 50<br>51                 | 919 | Reproduction: No data on reproduction time are available.  |
| 52<br>53                 | 920 | Remarks: Leucosolenia somesii was considered a minor synonym of L. variabilis until a recent                 |
| 54<br>55<br>56           | 921 | study by van Soest et al. (2007) was published. They showed valuable differences between these               |
| 57<br>58                 | 922 | two species, based on a large number of specimens, including the type material. Here we provide              |
| 59<br>60                 | 923 | its first molecular data and an updated morphological description. Our novel data confirms that L.           |
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somesii represents a distinct species, based on both morphological and molecular analyses. The reexamination of spicules of specimen ZMA Por. 17572 studied by van Soest et al. (2007) confirms the strong correspondence of its specular characteristics to the paralectotype BMNH 1956.4.26.35 (Fig. 23C, D). According to our phylogenetic reconstruction, the most closely related species is Arctic *Leucosolenia creepae* sp. nov., with which *Leucosolenia somesii* shares some specific morphological features: echinate external appearance due to the high number of non-lanceolate diactines protruding to the external surface, dimensions of tri- and tetractines. The discussion of their differences is given above under Leucosolenia creepae sp. nov. description. From all other North Atlantic and Arctic Leucosolenia species, L. somesii differs by the absence of lanceolate diactines.

#### 

**Discussion** 

#### 935 Leucosolenia taxonomy

The calcarean biodiversity in the Arctic region remains poorly studied. In the beginning of the XX century, only three asconoid species were detected in the White and Barents Seas (Breitfuss, 1898a): Ascandra variabilis, A. contorta and A. fabricii. This view was even more simplified in subsequent works, designating that only Leucosolenia complicata occurs in these seas (Koltun, 1952). The present study demonstrates that L. complicata is restricted to the North-East Atlantic, while in the Arctic, the Leucosolenia diversity is represented by at least four species: Leucosolenia corallorrhiza, L. variabilis, Leucosolenia creepae sp. nov. and yet undescribed *Leucosolenia* sp. A. These species represent different clades in the present phylogenetic analysis (Fig. 1): (1) Clade I, which includes the nominative species, L. corallorrhiza, Leucosolenia sp. A and several phylogenetically distinct lineages formed by Norway and Greenland specimens (Leucosolenia sp. B-D), (2) Clade II with the closely related North-Atlantic L. somesii and Leucosolenia creepae sp. nov. restricted to the White Sea. Leucosolenia botryoides, the type species of the genus, shows sister relationships with the latter group, and L. complicata is sister to this clade. The phylogenetic signal from different molecular markers gave a similar result (Data S1), supporting the chosen species hypothesis from the concatenated phylogenetic dataset (Figs 1, S1, S2), and the lack of heterozygous sites in the studied markers indicates the absence of hybridization between different species and supports well-established species boundaries.

Overall, species within each group show several common characters, which correlate with the level of molecular divergence. Most of the traits traditionally used in calcarean systematics (the general skeleton composition, the spicular set and their fine morphology) supported the species hypothesis obtained from the molecular phylogenetic analysis. Spicular set is the most useful character to delimit species, and general spicular composition and proportions of spicules may also have a certain phylogenetic signal: (i) Lanceolate diactines with or without spines are found within the Clade I (Fig. 1) and in *L. complicata;* (ii) T-shaped tri- and tetractines with shorter

960 unpaired actines are common only in species from the Clade I; (iii) sister species *L. somesii* and
961 *L. creepae* sp. nov. bear generally thinner spicules than other species of the genus; these species
962 also have an echinate appearance due to the high number of long diactines protruding through the
963 surface.

Here we also show that modern techniques like scanning electron microscopy may give a new insight into understanding actual biodiversity, as in closely related species, similar spicular types differ by fine features like the presence or absence of spines. This micromorphological approach has been successfully applied in the taxonomy of calcareous sponges from subclass the Calcinea (Azevedo et al., 2009, 2015; Klautau et al., 2016). For example, although both *Leucosolenia creepae* sp. nov. and *L. somesii* have similar specular sets, the latter species has two populations of non-lanceolate diactines (spined and smooth), while only a single population of spined diactines with a continuous reduction of spines in larger diactines is found in *Leucosolenia* creepae sp. nov. The same was shown for L. variabilis and L. corallorrhiza: both species have lanceolate and sometimes spined diactines, but in L. variabilis we also detected thin long and highly spined trichoxeas. These differences in diactine types are obvious with the help of SEM techniques but may be overlooked during investigation using light microscopy alone.

The proportion of spicular types in different parts of cormus and the form of actines in tri-and tetractines (straight vs. bent) may also be important taxonomical characters. Haeckel (1872) recognised several species as distinct based on these differences. For example, Ascandra fabricii differs from *L. complicata* by the absence of tetractines, and *Asculmis armata* Haeckel, 1870 – by a low amount of triactines. Furthermore, Haeckel described two varieties of L. complicata: L. complicata var. hispida with straight paired actines and L. complicata var. ameboides in which paired actines are bent. Also, some varieties – cervicornis, confervicola, arachnoides, hispidissima - were distinguished for *L. variabilis* (Haeckel, 1872). Although these names are currently accepted as valid species within Leucosolenia (de Voogd et al., 2023), these forms were for a long time designated as morphotypes of either L. complicata or L. variabilis (Minchin, 1904; Burton,

986 1963) and no integrative study of Norway species has been conducted yet to support or reject their 987 validity. At the same time, our results show that *Leucosolenia* sp. A differs from *L. corallorrhiza* 988 by the low amount of tetractines and by bent paired actines in triactines. Since we suspect that 989 *Leucosolenia* sp. A also represents a distinct species, all the above-mentioned characters should 990 be taken into account for further taxonomical revisions of the North Atlantic and Arctic *Leucosolenia*.

In addition to the proportion of spicules in different parts of a skeleton, we identified several differences in the spiculation of the oscular crown among different species. The oscular rim bears numerous diactines protruding through the surface and forming a crown. The crown may be short (max length 50-100 µm, in most species) or extremely long (up to 500 µm, in *Leucosolenia creepae* sp. nov.). This character was not tracked in previous works on *Leucosolenia* systematics, as researchers commonly pointed at the smooth or echinate surface of the oscular tube without considering the spiculation of the oscular rim. The only exception is a recent description of *Leucosolenia salpinx* van Soest, 2017 as in this species the extended, long oscular crown is a notable diagnostic trait (van Soest, 2017). At the same time, forms with more or less echinate surface may be found in *L. corallorrhiza* in the White Sea, making its external appearance similar to that of sympatrically occurring *Leucosolenia creepae* sp. nov. In this case, the oscular rim spiculation is more useful for accurate identification of these species in the field. However, the phylogenetic value of this character is a subject for further studies, as the current results lack data on L. somesii, L. botryoides and Leucosolenia sp. A.

Another source of species-specific traits is the cytological structure of the studied
Leucosolenia species. It is well known that cytological characters, such as the cell types with
inclusions, are very important for Demospongiae and Homoscleromorpha species without skeleton
identification (*e.g.*, Muricy *et al.*, 1996; Ereskovsky *et al.*, 2011, 2017b; Gazave *et al.*, 2013;
Willenz *et al.*, 2016). Despite the fact, that the set of cell types is generally similar among studied *Leucosolenia* species, some species have characteristic cytological features. Both *L. corallorrhiza*

and L. variabilis have unique types of mesohyl cells - cells with inclusions (granular and/or spherulous) (Table S2), while L. complicata and Leucosolenia creepae sp. nov. lack such cells. There are no cytological data for *L. somesii* and *Leucosolenia* sp. A., but it is reasonable to assume that considering the phylogenetic position of these species, *Leucosolenia* sp. A. should have cells with inclusions and L. somesii should not. We have for the first time clearly shown the presence of myocytes in all species studied, as well as the presence of endopinacocytes in the oscular ring. However, further broad studies of cytology in the genus Leucosolenia are required to evaluate the phylogenetic value of these characters. 

The composition of symbiotic bacteria also shows variation among studied species, both in the number of bacterial morphotypes and their morphology (Table S2). The differences in composition of symbiotic bacteria are obvious even in the case of closely related sympatric species, e.g., L. corallorrhiza and L. variabilis. Considering the stability of the core microbiome for a particular sponge species across various localities and different environmental conditions (Webster & Thomas, 2016), it could become a useful character for delimiting calcareous sponge species. Species-specific traits in microbiome composition could already be revealed by cytological studies, but the metabarcoding approach enables much more detailed analysis and precise comparison of microbiome composition (Ribeiro et al., 2023), and should be preferred when possible.

#### Integrative taxonomy of Calcaronea

1031 The systematics of subclass Calcaronea is currently facing many challenges due to the 1032 broad implementation of molecular methods in taxonomical studies. Traditional taxonomical 1033 schemes were primary typologic (Borojevic *et al.*, 1990, 2000) and the high level of morphological 1034 homoplasy obscured the actual evolutionary relationships of high-level taxa within Calcarea 1035 (Manuel *et al.*, 2003). As a result, contemporary studies have partially shown the broad 1036 incongruence of morphology-based classification with newly reconstructed molecular phylogenies 1037 (Dohrmann *et al.*, 2006; Voigt *et al.*, 2012; Voigt & Wörheide, 2016; Alvizu *et al.*, 2018). Page 43 of 92

However, these analyses strongly supported the monophyly of both subclasses. Calcinea and Calcaronea. In the most recent molecular phylogeny of Calcaronea based on broad taxon sampling and two ribosomal markers, the para- and polyphyly of most families and genera were shown, along with the high hidden diversity (Alvizu et al., 2018). The phylogenetic studies within Calcaronea are also complicated by the unusual architecture of the mitochondrial genome in this group, which is organised in individual linear chromosomes and has a unique genetic code and accelerated rates of sequence evolution (Lavrov et al., 2013, 2016b). This hampers using of the mitochondrial markers, in particular the standard barcode cytochrome c oxidase subunit I (COI). Accordingly, the phylogenetic and taxonomical studies were primarily based on the nuclear ribosomal genes and internal transcribed spacers (ITS1, ITS2) (Voigt et al., 2012; Klautau et al., 2013, 2020, 2021; Azevedo et al., 2015, 2017; Sanamyan et al., 2022). Among ribosomal markers, the C-region of 28S was suggested as a barcode marker for delimitation of taxa on species- and generic levels (Voigt & Wörheide, 2016) and was successfully applied in subsequent integrative studies (Alvizu et al., 2019; Cóndor-Luján et al., 2019; Klautau et al., 2020, 2021). In the present study, this marker was used for species delimitation as well, but it gave doubtful results at the lowest taxonomical level. There was an obvious genetic break between some closely related species, e.g., L. somesii and Leucosolenia creepae sp. nov., but in the case of Leucosolenia species from Clade I, the interspecific differences may be equal to intraspecific values, like in L. corallorrhiza, L. variabilis and Leucosolenia sp. A (Table 2, Fig. 2). As a result, automatic species delimitation approaches like ASAP failed to find a barcode gap and delimit species within this group, suggesting either oversplitting or overlamping scenarios. Therefore, while the C-region gives a good resolution at the species-level and is useful for detection of the genetic breaks among species, it cannot be considered equivalent to standard mitochondrial COI. The same challenges in detecting of a barcode gap in the C-region dataset were previously shown in the larger taxon sampling of Calcaronea (Alvizu et al., 2018). At the same time, SSU data do not provide sufficient divergence rates to test the automatic species delimitation methods (Data S1, S2, see also Alvizu

et al., 2018). Another part of the nuclear ribosomal operon, ITS1 and ITS2, often contain numerous allele indels in Calcaronea, which complicate the PCR and sequencing processes (Wörheide et al., 2004; Voigt & Wörheide, 2016) and therefore are not useful for testing the large taxon sampling. In this study, the first molecular analysis of calcareous sponges using the nuclear histone H3 marker was performed. Within Leucosolenia, the H3 showed higher substitution rates at both intra-and interspecific levels than ribosomal markers (Tables 2, 3; Fig. 2), and topologies of the H3-based and ribosomal trees were congruent. High substitution rates enable the usage of ASAP, and its results fully confirmed the phylogenetic species hypothesis even within Clade I (Data S2). This suggests the H3 marker may be a powerful tool for taxonomical studies within calcaronean sponges in addition to the widely used LSU, and is more useful for delimitation of closely related species. Also, histone H3 is a protein-coding gene with a conservative protein sequence, which eases a verification of possible incorrect base-calling during sequencing. The congruent topologies of single-gene trees based on LSU and H3 markers also indicate that the latter may further improve the resolution and support of multi-locus phylogenetic trees within Calcaronea. 

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#### Patterns of biodiversity and biogeography

The obtained results indicate that there is an obvious connectivity between the calcaronean fauna of the White Sea and Greenland, since specimens of L. corallorrhiza and Leucosolenia sp. from these localities are conspecific in the molecular phylogenetic analysis. Although both localities are parts of the Arctic regions, this is an unexpected result in the case of the genus Leucosolenia. Members of this genus have short-living larvae, which are not capable of dispersal to a great distance (Anakina, 1981) so such wide distribution ranges accommodated by long-distance deposition of larvae via existing marine currents seem very unlikely. The connectivity of adult forms is also prevented by the deep and wide Fram Straight (max depth 2545 m) between Greenland and Svalbard, hampering the migration of shallow water forms with short-living larvae or direct development via the North Atlantic route (Meyer-Kaiser et al., 2022). We may suggest two possible explanations for such connectivity. Likely both Leucosolenia corallorrhiza and

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1090 Leucosolenia sp. A have a circumpolar distribution and may be found in other regions across the 1091 Arctic. Another explanation is that the connectivity of distant populations is achieved by 1092 transportation via ballast waters or on ship bottoms as a common part of the Arctic fouling 1093 communities. Unfortunately, the exact mechanism cannot be evaluated based on the molecular 1094 data presented in this study, as the population structure is not evident from the conservative nuclear 1095 markers used.

1096 For now, possible calcarean faunal connections between the Arctic and the Boreal North-1097 East Atlantic waters cannot be evaluated due to the low number of sequenced specimens from these regions. Leucosolenia diversity in Norwegian waters estimates 10 species (de Voogd et al., 1098 2023), among which at least Leucosolenia variabilis and L. corallorrhiza inhabit the Arctic. At 1099 1100 the same time, neither L. complicata nor L. botryoides were found in the White Sea, and *Leucosolenia creepae* sp. nov. and *L. somesii* represent two closely related and morphologically 1101 similar but distinct species, suggesting the temperate Atlantic fauna is distinct. The integrative 1102 studies of the Barents Sea biodiversity are therefore of crucial importance for a precise account of 1103 1104 possible faunal links.

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   1130 Competing Interests
- 5556 1131 Authors declare no competing interests.

60 1133 Data Availability Statement

All new sequences obtained in the study were deposited in GenBank, GenBank accession numbers are listed in Table S1. Unedited Maximum Likelihood single-gene trees (LSU, H3, 18S and concatenated datasets) are available in the Supplementary materials. Generated raw data on the external morphology, skeleton organisation, spicule types and cytology of studied specimens are available in the Mendeley Data repository (10.17632/4pkf7x4jb9.1 and 10.17632/r7f8zmdh2n.1).

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| Marker     | Primers  | PCR conditions  | Reference                |  |
|------------|--|---|--------------------------|--|
| 28S rRNA   | 28S-C2FGAA AAG AAC TTT GRA RAG AGA GT28S-D2RTCC GTG TTT CAA GAC GGG  | 5 min - 94°C, 35x[1 min -<br>95°C, 45s - 50°C, 1 min -<br>72°C], 7 min - 72°C   | Chombard et al.,<br>1998 |  |
| 18S rRNA   | 18S-328FCCTGGTGATCCTGCCAG18S-HI + RCAACTAAGAACGGCCATGCAC18S-329RTAA TGA TCC TTC CGC AGG TT18S-A-FCAG CMG CCG CGG TAA TWC | 5 min - 94°C, 35x[1 min -<br>94°C, 1 min - 50°C, 2 min -<br>72°C], 7 min - 72°C | Alvizu et al., 201       |  |
| Histone H3 | Por_h3f<br>ATG GCC CGT ACC AAG CAG ACT GC<br>Por_h3r   | 5 min - 94°C, 35x[15 s - 95°C,<br>30s - 50°C, 45 s - 72°C], 7 min -<br>72°C     | This study               |  |

# Table 2. Uncorrected intra- and interspecific *p*-distances of 28S marker in the genus *Leucosolenia*. Intraspecific distances are highlighted in bold.

|                    | L. complicata | L. corallorrhiza | Leucosolenia sp. A | L. variabilis | L. creepae | L. somesii |
|--------------------|---------------|------------------|--------------------|---------------|------------|------------|
| L. complicata      | 0-0.4         |                  |                    |               |            |            |
| L. corallorrhiza   | 6.3-8         | 0-1.3            |                    |               |            |            |
| Leucosolenia sp. A | 6.8-7.6       | 0.4-1.7          | 0                  |               |            |            |
| L. variabilis      | 5.1-7.2       | 0.4-1.3          | 0.8-1.3            | 0-0.4         |            |            |
| L. creepae         | 5.1-5.9       | 8.9-10.5         | 8.9-9.3            | 8.9-9.3       | 0-0.4      |            |
| L. somesii         | 5.5-5.9       | 10.5-11.8        | 10.1               | 10.1          | 2.5-3      | -          |

| 1387                    | Table 3.                  | Unco         | rrected      | l intra-<br>s are h | and in       | ntersp<br>abted i | ecific <i>p</i><br>n bold | -dista     | nces of         | °H3 m        | arke                                    | r in the ge | enus <i>L</i> | eucosole   | enia. |   |
|-------------------------|---------------------------|--------------|--------------|---------------------|--------------|-------------------|---------------------------|------------|-----------------|--------------|---|-------------|---------------|------------|-------|---|
| 1300                    |                           |              | L            | . complic           | ata          | L. corallorrhiza  | Le                        | ucosolen   | <i>ia</i> sp. A | L. variabili |   |             | L. creepae    | 2          |       |   |
|                         | L. compli                 | cata         |              | 0-0.4               |              |                   |                           |            |                 |              |   |             |               |            |       |   |
|                         | L. corallo                | orrhiza      |              | 7.9-8.8             |              | 0                 | )-0.8                     |            |                 |              |   |             |               |            |       |   |
|                         | Lauranal                  |              | ٨            | 9406                |              | 2                 | 5 7 7                     |            | 040             | 0            |   |             |               |            |       |   |
|                         | Leucosoie                 | enia sp. 1   | A            | 8.4-9.0             |              | ۷.                | 3-3.3                     |            | 0.4-0           | .ð           |   |             |               |            |       |   |
|                         | L. variabi                | ilis         |              | 7.9-9.2             |              | 3                 | 3.3-5                     |            | 2.9-4           | .2           |   | 0-1.3       |               |            |       |   |
|                         | L. creepa                 | е            |              | 3.8-4.6             |              | 9.0               | 5-10.5                    |            | 9.6-10          | ).5          |   | 9.2-10.9    |               | 0-1.7      |       |   |
| 1389                    |                           |              |              |                     |              |                   |                           |            |                 |              |   |             |               |            |       |   |
| 1390                    | Table 4                   | Snici        | ıle dim      | ension              | s of I       | eucos             | olenia d                  | romnl      | icata (         | Monta        | on 1                                    | 814)        |               |            |       |   |
| Spic                    | ule                       | spice        | Ler          | ngth (µ             |              | encos             | orenia                    | Wi         | dth (µ          | m)           | 54, 1                                   | 011)        | A             | ngle (°)   |       |   |
| <b>i</b>                |                           | Min          | Mean         | Max                 | SD           | n                 | Min                       | Mean       | Max             | SD           | n                                       | Min         | Mean          | Max        | SD    |   |
| Curved lan<br>diactines | iceolate                  | 147.1        | 263.7        | 389.3               | 58.8         | 46                | 6.6                       | 9.5        | 14.4            | 1.4          | 47                                      | -           | -             | -          | -     |   |
| Trichoxeas              | 5                         | 66.0         | 127.3        | 248.7               | 40.3         | 18                | 1.5                       | 2.4        | 3.5             | 0.6          | 17                                      | -           | -             | -          | -     |   |
| Triactines              |                           |              |              |                     |              |                   |                           |            |                 |              |   |             |               |            |       | _ |
| Unpair                  | ed actine                 | 75.3         | 113.5        | 149.4               | 19.4         | 40                | 3.7                       | 6.3        | 10.5            | 1.3          | 41                                      | -           | -             | -          | -     |   |
| Paired                  | actines                   | 62.3         | 94.9         | 122.0               | 13.1         | 81                | 4.1                       | 6.8        | 11.8            | 1.3          | 81                                      | 120.4       | 125.7         | 137.9      | 4.4   | 2 |
| Tetractines             | 5                         |              |              |                     |              |                   |                           |            |                 |              |   |             |               |            |       |   |
| Unpair                  | ed actine                 | 30.6         | 109.3        | 156.7               | 24.1         | 56                | 3.7                       | 6.7        | 9.7             | 1.2          | 57                                      | -           | -             | -          | -     |   |
| Paired                  | actines                   | 14.2         | 93.7         | 170.2               | 20.4         | 109               | 3.5                       | 6.9        | 10.0            | 1.2          | 108                                     | 115.7       | 123.5         | 134.0      | 4.4   | 4 |
| <i>Apical (</i><br>1391 | actine                    | 8.9          | 23.8         | 49.7                | 11.1         | 56                | 2.0                       | 5.3        | 10.1            | 1.4          | 57                                      | -           | -             | -          | -     |   |
| 1392                    | Table 5                   | Snici        | ıle dim      | ension              | s of I       | eucos             | olenia d                  | corall     | orrhiza         | ı (Haec      | ekel                                    | 1872)       |               |            |       |   |
| Spic                    | ule                       | spiec        | Ler          | ngth (u             |              | eneos             | 0101114                   | Wi         | dth (u          | m)           | ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,, | 10/2)       | A             | ngle (°)   |       |   |
|                         |                           | Min          | Mean         | Max                 | SD           | n                 | Min                       | Mean       | Max             | SD           | n                                       | Min         | Mean          | Max        | SD    |   |
| Curved lan<br>diactines | nceolate                  | 73.5         | 179.0        | 455.0               | 74.2         | 27                | 2.8                       | 6.0        | 11.4            | 1.7          | 27                                      | -           | -             | -          | -     |   |
| Triactines              |                           |              |              |                     |              |                   |                           |            |                 |              |   |             |               |            |       |   |
| Unpair                  | ed actine                 | 37.1         | 70.5         | 100.0               | 14.9         | 91                | 3.5                       | 6.5        | 10.5            | 1.4          | 91                                      | -           | -             | -          | -     |   |
| Paired                  | actines                   | 36.1         | 82.7         | 133.3               | 20.4         | 165               | 3.1                       | 6.5        | 12.5            | 1.5          | 164                                     | 124.6       | 142.9         | 151.2      | 6.0   |   |
| 1 011 001               |                           |              |              |                     |              |                   |                           |            |                 |              |   |             |               |            |       |   |
| Tetractines             | 5                         |              |              |                     |              |                   |                           |            |                 |              |   |             |               |            |       |   |
| Tetractines<br>Unpair   | s<br>ed actine            | 46.8         | 68.8         | 98.8                | 16.0         | 14                | 3.5                       | 5.6        | 7.9             | 1.2          | 15                                      | -           | -             | -          | -     |   |
| Tetractines<br>Unpaired | s<br>ed actine<br>actines | 46.8<br>37.6 | 68.8<br>80.7 | 98.8<br>128.1       | 16.0<br>22.5 | 14<br>23          | 3.5<br>2.4                | 5.6<br>5.8 | 7.9<br>8.4      | 1.2<br>1.6   | 15<br>25                                | - 144.8     | -<br>151.4    | -<br>159.3 | - 4.0 |   |

| 3         | <sup>3</sup> Spicule |       | Length (µm) |            |        |       |        | Width (µm) |        |     |     |       | Angle (°) |         |     |    |  |  |
|-----------|----------------------|-------|-------------|------------|--------|-------|--------|------------|--------|-----|-----|-------|-----------|---------|-----|----|--|--|
| 4         | •                    | Min   | Mean        | Max        | SD     | n     | Min    | Mean       | Max    | SD  | n   | Min   | Mean      | Max     | SD  | n  |  |  |
| 5 '<br>6  | Curved smooth        |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 7         | lanceolate           | 123.7 | 306.7       | 482.4      | 100.1  | 27    | 5.7    | 9.8        | 13.9   | 1.8 | 27  | -     | -         | -       | -   | -  |  |  |
| 8         | diactines            |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 9         | Strait spiny         |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 10        | lanceolate           | 111.4 | 151.4       | 191.2      | 28.7   | 5     | 2.4    | 3.1        | 3.8    | 0.5 | 10  | -     | -         | -       | -   | -  |  |  |
| 12.       | diactines            |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 13        | Trichoxeas           |       | fragme      | nts, up to | 362.4  |       | 0.6    | 0.9        | 1.3    | 0.2 | 17  | -     | -         | -       | -   | -  |  |  |
| 14        | Triactines           |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 15        | Unpaired actine      | 46.5  | 122.3       | 208.6      | 31.7   | 63    | 5.0    | 8.1        | 11.6   | 1.5 | 63  | -     | -         | -       | -   | -  |  |  |
| 16<br>17. | Paired actines       | 39.0  | 127.9       | 180.2      | 29.3   | 130   | 4.0    | 8.5        | 12.7   | 1.7 | 128 | 122.1 | 138.5     | 150.6   | 4.8 | 42 |  |  |
| 18        | Tetractines          |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 19        | Unpaired actine      | 101.3 | 147.6       | 196.2      | 28.4   | 12    | 6.4    | 8.5        | 11.1   | 1.5 | 12  | -     | -         | -       | -   | -  |  |  |
| 20        | Paired actines       | 92.5  | 142.0       | 172.5      | 22.3   | 25    | 6.0    | 9.1        | 11.5   | 1.5 | 26  | 134.0 | 142.2     | 155.3   | 4.6 | 34 |  |  |
| 21        | Apical actine        | 16.5  | 22.8        | 31.1       | 5.2    | 14    | 4.7    | 5.9        | 8.5    | 1.4 | 15  | -     | -         | -       | -   | -  |  |  |
| 22        | 1396                 |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 23        | 1397 Table 7.        | Spice | ıle dim     | ension     | s of L | eucos | olenia | sp. A.     |        |     |     |       |           |         |     |    |  |  |
| 25        | Spicule              |       | Ler         | ngth (µ    | ιm)    |       |        | Wie        | lth (µ | m)  |     |       | Ang       | gle (°) |     |    |  |  |
| 26        |                      | Min   | Mean        | Max        | SD     | n     | Min    | Mean       | Max    | SD  | n   | Min   | Mean      | Max     | SD  | n  |  |  |
| 27        | Curved spiny         |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 28        | lanceolate           | 131.9 | 189.1       | 311.0      | 48.2   | 15    | 5.8    | 7.2        | 8.8    | 0.8 | 15  | -     | -         | -       | -   | -  |  |  |
| 30.       | diactines            |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 31        | Curved smooth        | _     | 515.0       | _          | 83.4   | 2     |        | 11.6       | _      | 17  | 2   |       |           |         |     |    |  |  |
| 32        | diactines            | _     | 515.0       | _          | 00.4   | 2     |        | 11.0       | -      | 1.7 | 2   |       |           |         |     |    |  |  |
| 33        | Triactines           |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 34<br>35. | Unpaired actine      | 81.6  | 118.5       | 179.7      | 22.2   | 42    | 6.3    | 8.9        | 11.1   | 1.1 | 43  | -     | -         | -       | -   | -  |  |  |
| 36.       | Paired actines       | 80.5  | 125.1       | 174.8      | 22.4   | 74    | 6.6    | 9.1        | 11.6   | 1.0 | 84  | 132.9 | 146.5     | 160.7   | 5.6 | 44 |  |  |
| 37        | Tetractines          |       |             |            |        |       |        | (          |        |     |     |       |           |         |     |    |  |  |
| 38        | Unpaired actine      | 82.4  | 114.3       | 157.1      | 27.2   | 5     | 7.9    | 8.6        | 9.9    | 0.7 | 7   | -     | -         | -       | -   | -  |  |  |
| 39        | Paired actines       | 79.3  | 113.2       | 132.0      | 16.0   | 14    | 7.0    | 8.5        | 10.2   | 0.9 | 14  | 137.4 | 140.8     | 146.2   | 3.3 | 5  |  |  |
| 40<br>41  | Apical actine        | 25.7  | 30.0        | 34.2       | 4.2    | 3     | 5.8    | 7.1        | 8.5    | 1.2 | 5   | -     | -         | -       | -   | -  |  |  |
| 41        | 1398                 |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 43        | 1399                 |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 44        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 45        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 46<br>47  |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 47        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 49        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 50        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 51        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 52        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 53<br>51  |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 55        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 56        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 57        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 58        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 59        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |
| 60        |                      |       |             |            |        |       |        |            |        |     |     |       |           |         |     |    |  |  |

| Spicule               | Length (µm) |         |         |        |       | Width (µm) |        |         |        |         |       | Angle (°) |         |     |   |  |
|-----------------------|-------------|---------|---------|--------|-------|------------|--------|---------|--------|---------|-------|-----------|---------|-----|---|--|
|                       | Min         | Mean    | Max     | SD     | n     | Min        | Mean   | Max     | SD     | n       | Min   | Mean      | Max     | SD  | 1 |  |
| Spiny diactines       | 83.3        | 194.9   | 478.8   | 98.6   | 81    | 2.9        | 5.1    | 9.6     | 1.4    | 81      | -     | -         | -       | -   |   |  |
| <b>Triactines</b>     |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| Unpaired actine       | 29.9        | 80.7    | 125.8   | 18.8   | 87    | 3.0        | 5.4    | 8.8     | 1.1    | 89      | -     | -         | -       | -   |   |  |
| Paired actines        | 37.9        | 94.2    | 148.6   | 20.9   | 169   | 2.7        | 5.9    | 9.0     | 1.2    | 173     | 118.5 | 131.1     | 137.7   | 4.7 | 2 |  |
| <b>Tetractines</b>    |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| Unpaired actine       | 31.8        | 85.1    | 137.5   | 25.9   | 15    | 4.1        | 6.2    | 7.7     | 1.3    | 15      | -     | -         | -       | -   |   |  |
| Paired actines        | 52.9        | 95.3    | 136.2   | 20.6   | 30    | 3.7        | 6.3    | 8.1     | 1.2    | 31      | 131.0 | 139.5     | 146.1   | 4.5 |   |  |
| Apical actine         | 12.7        | 25.6    | 43.7    | 9.8    | 18    | 2.9        | 5.2    | 8.4     | 1.3    | 18      | -     | -         | -       | -   |   |  |
| 1401                  |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| 1402 <b>Table 9</b> . | . Spici     | ıle dim | ension  | s of L | eucos | olenia     | somesi | i (Bow  | verbar | ık, 18' | 74)   |           |         |     |   |  |
| Spicule               | 1           | Ler     | ngth (µ | ιm)    |       |            | Wie    | lth (µı | m)     | ,       | ,     | An        | gle (°) |     |   |  |
|                       | Min         | Mean    | Max     | SD     | n     | Min        | Mean   | Max     | SD     | n       | Min   | Mean      | Max     | SD  |   |  |
| Curved smooth         |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| diactines             | 148.0       | 424.5   | 649.8   | 116.0  | 68    | 6.5        | 9.9    | 12.3    | 1.3    | 88      | -     | -         | -       | -   |   |  |
| Straight spinv        |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| diactines             | 68.0        | 90.0    | 105.8   | 8.9    | 58    | 2.1        | 3.3    | 4.1     | 0.4    | 39      | -     | -         | -       | -   |   |  |
| Triactines            |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| Unnaired actine       | 58.3        | 127.4   | 186.1   | 27.0   | 60    | 6.5        | 8.2    | 10.3    | 1.0    | 64      | _     | -         | -       | _   |   |  |
| Paired actines        | 86.2        | 155.2   | 226.0   | 28.4   | 111   | 5.9        | 8.0    | 10.7    | 1.0    | 131     | 117.1 | 131.7     | 139.9   | 4.2 | 6 |  |
| Totractinos           |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| Unnaired actine       | 103.5       | 156.0   | 206.9   | 32.9   | 10    | 84         | 91     | 10.5    | 0.6    | 13      | _     | _         | _       | _   |   |  |
| Paired actines        | 1560        | 178.7   | 200.9   | 14.5   | 23    | 7.8        | 9.5    | 11.1    | 0.0    | 24      | 133.5 | 1393      | 149 1   | 3.9 | 1 |  |
| Anical actine         | 16.6        | 21.5    | 209.2   | 4.2    | 8     | 8.1        | 10.0   | 11.1    | 1.0    | 7       | -     | -         | -       | -   |   |  |
| 1/03                  | 10.0        | 21.5    | 27.5    | 7.4    | 0     | 0.1        | 10.0   | 11.2    | 1.0    | /       |       |           |         |     |   |  |
| 1403                  |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| 1405                  |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
| 1405                  |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |
|                       |             |         |         |        |       |            |        |         |        |         |       |           |         |     |   |  |

#### **Figure captions**

| 1407 | Figure 1. The molecular phylogenetic hypothesis of the genus <i>Leucosolenia</i> based on the Bayesian |
|------|--|
| 1408 | analysis of the concatenated dataset (28S, 18S and H3 markers). Initial species names are used on      |
| 1409 | the tips, bold font indicates original specimens used in this study. Each putative species-level clade |
| 1410 | is coloured according to the revised species hypothesis, the suggested revised species names are       |
| 1411 | provided on the right. All calcaronean species except representatives of the genus Leucosolenia        |
| 1412 | are collapsed into a single group, 'Calcaronea rest'. Numbers above branches indicate posterior        |
| 1413 | probabilities from the Bayesian Inference (> 0.9), numbers below branches – bootstrap support          |
| 1414 | from the Maximum Likelihood (>70).   |
|      |  |

Figure 2. Medium parsimony network analysis (TCS algorithm) of *Leucosolenia* species. The
relative size of circles is proportional to the number of sequences of that same genotype. A – 28S
alignment. Colours of the circles refer to the geographic origin of each genotype. Coloured
backgrounds and species names indicate the revised species hypothesis. B – H3 alignment. Colours
of the circles refer to the revised species hypothesis.

Figure 3. *Leucosolenia complicata* (Montagu, 1814). External morphology and skeleton. A –
general morphology (WS11661); B – skeleton of oscular rim (WS11662); C – skeleton of oscular
tube (WS11662); D – skeleton of cormus (WS11662). c – cormus, d – diverticulum, o – oscular,
or – oscular rim, ot – oscular tube.

1427 Figure 4. Leucosolenia complicata (Montagu, 1814). Spicule types, scanning electron
 1428 microscopy. A – curved lanceolate diactines; B, C – trichoxeas; D – triactines; E – tetractine.
 1429

Figure 5. Leucosolenia complicata (Montagu, 1814). Body wall structure and cell types of
 bordering tissues. A, B – semi-thin sections of sponge body wall; C – exopinacocyte; D –

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| 2  | 1432   | endopinacocyte; E – choanocytes; F – porocyte. Scale bars: A – 50 $\mu$ m, B – 20 $\mu$ m, C – F – 2 $\mu$ m.   |
|--|--|---|
| 4<br>5   | 1433   | ch – choanocytes, chd – choanoderm, en – endopinacocyte, ex – exopinacocyte, exp –  |
| 6<br>7   | 1434   | exopinacoderm, f – flagellum, m – mesohyl, mv – microvilli, n – nucleus, po – porocyte.   |
| 8<br>9   | 1435   |   |
| 10<br>11<br>12   | 1436   | Figure 6. Leucosolenia complicata (Montagu, 1814). Mesohyl cell types and symbiotic bacteria.   |
| 13<br>14   | 1437   | A – sclerocyte; B – amoebocyte; C – symbiotic bacteria, morphotype 1; D – symbiotic bacteria,   |
| 15<br>16<br>17   | 1438   | morphotype 2. Scale bars: A, B – 2 $\mu$ m, C, D – 0.5 $\mu$ m. am – amoebocytes, m – mesohyl, n –  |
| 17<br>18<br>19   | 1439   | nucleus, sp – spicule.  |
| 20<br>21   | 1440   |   |
| 22<br>23   | 1441   | Figure 7. Leucosolenia corallorrhiza (Haeckel, 1872). External morphology and skeleton. A –   |
| 24<br>25<br>26   | 1442   | general morphology (WS11642); B – skeleton of oscular rim (WS11653); C – skeleton of oscular  |
| 27<br>28   | 1443   | tube (WS11653); D – skeleton of cormus (WS11653). c – cormus, d – diverticulum, o – osculum,  |
| 29<br>30   | 1444   | oc – oscular crown, ot – oscular tube.  |
| 31<br>32<br>33   | 1445   |   |
| 34<br>35   | 1446   | Figure 8. Leucosolenia corallorrhiza (Haeckel, 1872). Spicule types, scanning electron  |
| 36<br>37   | 1447   | microscopy. A - curved lanceolate diactines; B - tips of diactines, I and II refer to the zones   |
| 38<br>39   | 1448   | marked on A, white arrowheads mark spines; C – tetractine; D – triactines.  |
| 40<br>41   |  |   |
| 4/   | 1449   |   |
| 42<br>43<br>44   | 1449<br>1450   | Figure 9. Leucosolenia corallorrhiza (Haeckel, 1872). Body wall structure and cell types of   |
| 42<br>43<br>44<br>45<br>46   | 1449<br>1450<br>1451                                 | <b>Figure 9.</b> <i>Leucosolenia corallorrhiza</i> (Haeckel, 1872). Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocyte; D –   |
| 42<br>43<br>44<br>45<br>46<br>47<br>48<br>49   | 1449<br>1450<br>1451<br>1452                         | <b>Figure 9.</b> <i>Leucosolenia corallorrhiza</i> (Haeckel, 1872). Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocyte; D – endopinacocyte; E – choanocytes, inset – junctions between choanocytes; F – porocyte. Scale bars:   |
| 42<br>43<br>44<br>45<br>46<br>47<br>48<br>49<br>50<br>51   | 1449<br>1450<br>1451<br>1452<br>1453                 | <b>Figure 9.</b> <i>Leucosolenia corallorrhiza</i> (Haeckel, 1872). Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocyte; D – endopinacocyte; E – choanocytes, inset – junctions between choanocytes; F – porocyte. Scale bars: $A - 50 \mu m$ , $B - 20 \mu m$ , $C - F - 2 \mu m$ , inset – 0.2 $\mu m$ . am – amoebocyte, ch – choanocytes, chd –  |
| 42<br>43<br>44<br>45<br>46<br>47<br>48<br>49<br>50<br>51<br>52<br>53                               | 1449<br>1450<br>1451<br>1452<br>1453<br>1454         | <b>Figure 9.</b> <i>Leucosolenia corallorrhiza</i> (Haeckel, 1872). Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocyte; D – endopinacocyte; E – choanocytes, inset – junctions between choanocytes; F – porocyte. Scale bars: $A - 50 \mu m$ , $B - 20 \mu m$ , $C - F - 2 \mu m$ , inset – 0.2 $\mu m$ . am – amoebocyte, ch – choanocytes, chd – choanoderm, en – endopinacocyte, exp – exopinacoderm, f – flagellum, m – mesohyl, mv –   |
| 42<br>43<br>44<br>45<br>46<br>47<br>48<br>49<br>50<br>51<br>52<br>53<br>54<br>55<br>55             | 1449<br>1450<br>1451<br>1452<br>1453<br>1454<br>1455 | <b>Figure 9.</b> <i>Leucosolenia corallorrhiza</i> (Haeckel, 1872). Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocyte; D – endopinacocyte; E – choanocytes, inset – junctions between choanocytes; F – porocyte. Scale bars: $A - 50 \mu m$ , $B - 20 \mu m$ , $C - F - 2 \mu m$ , inset – $0.2 \mu m$ . am – amoebocyte, ch – choanocytes, chd – choanoderm, en – endopinacocyte, exp – exopinacoderm, f – flagellum, m – mesohyl, mv – microvilli, n – nucleus, ph – phagosome, po – porocytes, sb – symbiotic bacteria. |
| 42<br>43<br>44<br>45<br>46<br>47<br>48<br>49<br>50<br>51<br>52<br>53<br>54<br>55<br>56<br>57<br>58 | 1449<br>1450<br>1451<br>1452<br>1453<br>1454<br>1455 | <b>Figure 9.</b> <i>Leucosolenia corallorrhiza</i> (Haeckel, 1872). Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocyte; D – endopinacocyte; E – choanocytes, inset – junctions between choanocytes; F – porocyte. Scale bars: A – 50 $\mu$ m, B – 20 $\mu$ m, C - F – 2 $\mu$ m, inset – 0.2 $\mu$ m. am – amoebocyte, ch – choanocytes, chd – choanoderm, en – endopinacocyte, exp – exopinacoderm, f – flagellum, m – mesohyl, mv – microvilli, n – nucleus, ph – phagosome, po – porocytes, sb – symbiotic bacteria.     |

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| 1457 | Figure 10. Leucosolenia corallorrhiza (Haeckel, 1872). Mesohyl cell types and symbiotic                                     |
|------|---|
| 1458 | bacteria. A - sclerocytes, inset - septate junctions between sclerocytes; B - amoebocytes; C, D -                           |
| 1459 | granular cells in the body wall; E – granular cell; F – degraded granular cell; G – myocytes, inset                         |
| 1460 | – bundles of myofilaments; H, I – symbiotic bacteria, morphotype 1. Scale bars: A, B, E, F – 2                              |
| 1461 | $\mu$ m, C – 50 $\mu$ m, C – 20 $\mu$ m, G – 5 $\mu$ m, H – 0.5 $\mu$ m, I – 1 $\mu$ m. ch – choanocytes, chd – choanoderm, |
| 1462 | ex - exopinacocyte, exp - exopinacoderm, f - flagellum, gc - granular cells gr - granule, m -                               |
| 1463 | mesohyl, mf - myofibrils, mv - microvilli, my - myocytes, n - nucleus, po - porocytes, sb -                                 |
| 1464 | symbiotic bacteria, sp – spicule.   |
| 1465 |   |
| 1466 | Figure 11. Leucosolenia variabilis Haeckel, 1870. External morphology and skeleton. A – general                             |
| 1467 | morphology (WS11731); B – skeleton of oscular rim (WS11731); C – skeleton of oscular tube; D                                |
| 1468 | – skeleton of cormus (WS11643). c – cormus, d – diverticulum, o – osculum, or – oscular rim, ot                             |
| 1469 | – oscular tube.   |
| 1470 |   |
| 1471 | Figure 12. Leucosolenia variabilis Haeckel, 1870. Spicule types, scanning electron microscopy.                              |
| 1472 | A – curved smooth lanceolate diactines; B – triactines; C – abnormal triactines; D – tetractine.                            |
| 1473 |   |
| 1474 | Figure 13. Leucosolenia variabilis Haeckel, 1870. Trichoxeas. A, B – general view of trichoxea                              |
| 1475 | fragments; C – enlarge view of trichoxea, white arrowhead marks spines. tx – trichoxeas.                                    |
| 1476 |   |
| 1477 | Figure 14. Leucosolenia variabilis Haeckel, 1870. Body wall structure and cell types of bordering                           |
| 1478 | tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocytes; D – choanocytes;                             |
| 1479 | E – porocyte; F – endopinacocyte. Scale bars: A – 50 $\mu$ m, B – 20 $\mu$ m, C, D – 5 $\mu$ m, E, F – 2 $\mu$ m.           |
| 1480 | am - amoeboid cell, ch - choanocytes, chd - choanoderm, ex - exopinacocyte, exp -   |
| 1481 | exopinacoderm, f - flagellum, m - mesohyl, mv - microvilli, n - nucleus, po - porocyte, sb -                                |
| 1482 | symbiotic bacteria, sc – spherulous cells.  |

| 2<br>3         | 1483 |  |
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| 4<br>5         | 1484 | Figure 15. Leucosolenia variabilis Haeckel, 1870. Mesohyl cell types and symbiotic bacteria. A                           |
| 6<br>7         | 1485 | - amoebocytes; B - large amoeboid cell; C - granular cell; D - myocyte, inset - bundles of                               |
| 8<br>9<br>10   | 1486 | myofilaments; E - spherulous cell; F - spherulous cells in the body wall; G - sclerocyte; H -                            |
| 11<br>12       | 1487 | symbiotic bacteria, morphotype 1. I – symbiotic bacteria, morphotype 2. J – symbiotic bacteria,                          |
| 13<br>14       | 1488 | morphotype 3. Scale bars: A – 2 $\mu$ m, B – 5 $\mu$ m, C-E – 2 $\mu$ m, F – 50 $\mu$ m, G – 2 $\mu$ m, H-J – 1 $\mu$ m. |
| 15<br>16<br>17 | 1489 | am – amoebocyte, bam – large amoeboid cell, ch – choanocytes, gc – granular cell, gr – granule,                          |
| 18<br>19       | 1490 | m – mesohyl, mf – myofibrils, n – nucleus, sc – spherulous cells, sp - spicule.  |
| 20<br>21       | 1491 |  |
| 22<br>23<br>24 | 1492 | Figure 16. Leucosolenia variabilis Haeckel, 1870. Spicule types in the White Sea specimen (A)                            |
| 24<br>25<br>26 | 1493 | and in the specimens from the British Natural History Museum collection (B-D). A – spicules from                         |
| 27<br>28       | 1494 | WS11731; B – spicules from BMNH 1906.12.1.40; C – spicules from BMNH 1906.12.1.50; D –                                   |
| 29<br>30<br>21 | 1495 | spicules from the syntype BMNH 1910.1.1.421.a. d – diactines, tx – trichoxeas.   |
| 31<br>32<br>33 | 1496 |  |
| 34<br>35       | 1497 | Figure 17. Leucosolenia sp. A. External morphology and skeleton. A – general morphology                                  |
| 36<br>37       | 1498 | (WS11752); B – skeleton of oscular rim (WS11770); C – skeleton of oscular tube (WS11770); D                              |
| 38<br>39<br>40 | 1499 | - skeleton of cormus (WS11752). c - cormus, d - diverticulum, o - osculum, or - oscular rim, ot                          |
| 41<br>42       | 1500 | – oscular tube.  |
| 43<br>44       | 1501 |  |
| 45<br>46<br>47 | 1502 | Figure 18. Leucosolenia sp. A. Spicule types, scanning electron microscopy. A – curved smooth                            |
| 47<br>48<br>49 | 1503 | diactines; B – curved spiny lanceolate diactines triactines; C – tips of diactines, I and II refer to                    |
| 50<br>51       | 1504 | the zones marked on A and B, white arrowheads mark spines; D - triactines; E - abnormal                                  |
| 52<br>53       | 1505 | triactines; F – tetractine.  |
| 54<br>55<br>56 | 1506 |  |
| 57<br>58       | 1507 | Figure 19. Leucosolenia creepae sp. nov. External morphology and skeleton. A - general                                   |
| 59<br>60       | 1508 | morphology (WS11702, holotype); B – skeleton of oscular rim (WS11762); C – skeleton of                                   |
|                |      |  |

1509 oscular tube (WS11728); D – skeleton of cormus (WS11655). c – cormus, d – diverticulum, o –
1510 osculum, oc – oscular crown, ot – oscular tube.

Figure 20. *Leucosolenia creepae* sp. nov. Spicule types, scanning electron microscopy. A – spiny
diactines; B – tips of diactines, I, II, III and IV refer zones marked on A, white arrowheads mark
spines; C – triactines; D – abnormal triactines; E – tetractines.

**Figure 21.** *Leucosolenia creepae* sp. nov. Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocyte; D – endopinacocyte and myocyte, inset - bundles of myofilaments in the myocyte; E – choanocytes. F – porocyte. Scale bars: A – 50  $\mu$ m, B – 20  $\mu$ m, C – 2  $\mu$ m, D – 5  $\mu$ m, E, F – 2  $\mu$ m. am – amoeboid cell, ch – choanocytes, chd – choanoderm, en – endopinacocyte, ex – exopinacocyte, exp – exopinacoderm, f – flagellum, m – mesohyl, mf – myofibrils, mv – microvilli, my – myocytes, n – nucleus, oo – oocyte, ph – phagosome, po – porocyte, sb – symbiotic bacteria.

**Figure 22.** *Leucosolenia creepae* sp. nov. Mesohyl cell types and symbiotic bacteria. A – sclerocyte; B – amoebocyte; C, D – symbiotic bacteria, morphotype 1; E, F – symbiotic bacteria, morphotype 2; G – young oocyte. Scale bars: A, B – 2  $\mu$ m, C - F – 0.5  $\mu$ m, G – 5  $\mu$ m. am – amoeboid cell, ch – choanocytes, m – mesohyl, n – nucleus, oo – oocyte, sb – symbiotic bacteria, sp – spicule.

Figure 23. Leucosolenia somesii (Bowerbank, 1874). External morphology and skeleton. A –
general morphology (ZMA Por. 17572); B – skeleton of cormus (ZMA Por. 17572); C, D –
spicules from BMNH 1956.4.26.35. c – cormus, d – diverticulum.

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| 1<br>2   | 1534 | Figure 24. Leucosolenia somesii (Bowerbank, 1874). ZMA Por. 17572. Spicule types, scanning  |
|--|------|---|
| 3<br>4<br>5  | 1535 | electron microscopy. A - curved smooth diactines; B - straight spiny diactines; C - tips of |
| 6<br>7   | 1536 | diactines, I and II refer to the zones marked on A and B, white arrowheads mark spines; D - |
| 8<br>9<br>10<br>11<br>12<br>13<br>14<br>15<br>16<br>17<br>18<br>19<br>20<br>21<br>22<br>33<br>24<br>25<br>26<br>27<br>28<br>29<br>30<br>31<br>22<br>23<br>24<br>25<br>26<br>27<br>28<br>29<br>30<br>31<br>32<br>33<br>45<br>36<br>37<br>38<br>39<br>40<br>41<br>42<br>43<br>44<br>55<br>56<br>57<br>58<br>59<br>60 | 1537 | triactines; E – abnormal triactines; F – tetractines.                                       |

## 1538 SUPPLEMENTARY MATERIAL

- **1539 Data S1.** Unedited Maximum Likelihood trees (LSU, H3, 18S and concatenated datasets).
- **Data S2.** Results of ASAP analysis based on LSU and H3 datasets.
- **Table S1.** List of *Leucosolenia* specimens used in the study. Voucher numbers, collection dates
- and localities, and GenBank accession numbers are given.
- **Table S2.** The cell types and symbiotic bacteria of the studied *Leucosolenia* species.
- 1544 Figure S1. The molecular phylogenetic hypothesis of the genus *Leucosolenia* based on the
- 1545 Bayesian analysis of the concatenated dataset (28S, 18S and H3 markers). Numbers above
- 1546 branches indicate posterior probabilities from the Bayesian Inference (> 0.9), numbers below
- 1547 branches bootstrap support from the Maximum Likelihood (>70).
- 1548 Figure S2. The molecular phylogenetic hypothesis of the genus Leucosolenia based on the
- 1549 Bayesian analysis of the concatenated dataset (28S and 18S markers). Numbers above branches
- 1550 indicate posterior probabilities from the Bayesian Inference (> 0.9), numbers below branches –
- 1551 bootstrap support from the Maximum Likelihood (>70).

- <sup>0</sup> 1555



Figure 1. The molecular phylogenetic hypothesis of the genus Leucosolenia based on the Bayesian analysis of the concatenated dataset (28S, 18S and H3 markers). Initial species names are used on the tips, bold font indicates original specimens used in this study. Each putative species-level clade is coloured according to the revised species hypothesis, the suggested revised species names are provided on the right. All calcaronean species except representatives of the genus Leucosolenia are collapsed into a single group, 'Calcaronea rest'. Numbers above branches indicate posterior probabilities from the Bayesian Inference (> 0.9), numbers below branches – bootstrap support from the Maximum Likelihood (>70).

267x296mm (300 x 300 DPI)





Figure 2. Medium parsimony network analysis (TCS algorithm) of Leucosolenia species. The relative size of circles is proportional to the number of sequences of that same genotype. A – 28S alignment. Colours of the circles refer to the geographic origin of each genotype. Coloured backgrounds and species names indicate the revised species hypothesis. B – H3 alignment. Colours of the circles refer to the revised species hypothesis.

214x256mm (300 x 300 DPI)


Figure 3. Leucosolenia complicata (Montagu, 1814). External morphology and skeleton. A – general morphology (WS11661); B – skeleton of oscular rim (WS11662); C – skeleton of oscular tube (WS11662); D – skeleton of cormus (WS11662). c – cormus, d – diverticulum, o – osculum, or – oscular rim, ot – oscular tube.

174x151mm (300 x 300 DPI)



Figure 4. Leucosolenia complicata (Montagu, 1814). Spicule types, scanning electron microscopy. A – curved lanceolate diactines; B, C – trichoxeas; D – triactines; E – tetractine.

173x137mm (300 x 300 DPI)



Figure 5. Leucosolenia complicata (Montagu, 1814). Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of sponge body wall; C – exopinacocyte; D – endopinacocyte; E – choanocytes; F – porocyte. Scale bars: A – 50 µm, B – 20 µm, C - F – 2 µm. ch – choanocytes, chd – choanoderm, en – endopinacocyte, ex – exopinacocyte, exp – exopinacoderm, f – flagellum, m – mesohyl, mv – microvilli, n – nucleus, po – porocyte.

173x193mm (300 x 300 DPI)



Figure 6. Leucosolenia complicata (Montagu, 1814). Mesohyl cell types and symbiotic bacteria. A – sclerocyte; B – amoebocyte; C – symbiotic bacteria, morphotype 1; D - symbiotic bacteria, morphotype 2. Scale bars: A, B – 2 μm, C, D – 0.5 μm. am – amoebocytes, m – mesohyl, n – nucleus, sp - spicule.

173x123mm (300 x 300 DPI)



Figure 7. Leucosolenia corallorrhiza (Haeckel, 1872). External morphology and skeleton. A – general morphology (WS11642); B – skeleton of oscular rim (WS11653); C – skeleton of oscular tube (WS11653); D – skeleton of cormus (WS11653). c – cormus, d – diverticulum, o – osculum, oc – oscular crown, ot – oscular tube.

174x144mm (300 x 300 DPI)



Figure 8. Leucosolenia corallorrhiza (Haeckel, 1872). Spicule types, scanning electron microscopy. A – curved lanceolate diactines; B – tips of diactines, I and II refer to the zones marked on A, white arrowheads mark spines; C – tetractine; D – triactines.

173x160mm (300 x 300 DPI)



Figure 9. Leucosolenia corallorrhiza (Haeckel, 1872). Body wall structure and cell types of bordering tissues.
A, B – semi-thin sections of body wall of sponge; C – exopinacocyte; D – endopinacocyte; E – choanocytes, inset – junctions between choanocytes; F – porocyte. Scale bars: A – 50 μm, B – 20 μm, C - F- 2 μm, inset – 0.2 μm. am – amoebocyte, ch – choanocytes, chd – choanoderm, en – endopinacocyte, exp – exopinacoderm, f – flagellum, m – mesohyl, mv – microvilli, n – nucleus, ph – phagosome, po – porocytes, sb – symbiotic bacteria.

173x194mm (300 x 300 DPI)



Figure 10. Leucosolenia corallorrhiza (Haeckel, 1872). Mesohyl cell types and symbiotic bacteria. A – sclerocytes, inset – septate junctions between sclerocytes; B – amoebocytes; C, D – granular cells in the body wall; E – granular cell; F – degraded granular cell; G – myocytes, inset - bundles of myofilaments; H, I
symbiotic bacteria, morphotype 1. Scale bars: A, B, E, F – 2 μm, C – 50 μm, C – 20 μm, G – 5 μm, H – 0.5 μm, I – 1 μm. ch – choanocytes, chd – choanoderm, ex – exopinacocyte, exp – exopinacoderm, f – flagellum, gc – granular cells gr – granule, m – mesohyl, mf – myofibrils, mv – microvilli, my – myocytes, n – nucleus, po – porocytes, sb – symbiotic bacteria, sp - spicule.

173x260mm (300 x 300 DPI)



Figure 11. Leucosolenia variabilis Haeckel, 1870. External morphology and skeleton. A – general morphology (WS11731); B – skeleton of oscular rim (WS11731); C – skeleton of oscular tube; D – skeleton of cormus (WS11643). c – cormus, d – diverticulum, o – osculum, or – oscular rim, ot – oscular tube.

174x136mm (300 x 300 DPI)



Figure 12. Leucosolenia variabilis Haeckel, 1870. Spicule types, scanning electron microscopy. A – curved smooth lanceolate diactines; B – triactines; C – abnormal triactines; D – tetractine.

173x184mm (300 x 300 DPI)





Figure 14. Leucosolenia variabilis Haeckel, 1870. Body wall structure and cell types of bordering tissues. A, B – semi-thin sections of body wall of sponge; C – exopinacocytes; D – choanocytes; E – porocyte; F – endopinacocyte. Scale bars: A - 50 μm, B – 20 μm, C, D – 5 μm, E, F – 2 μm. am - amoeboid cell, ch – choanocytes, chd – choanoderm, ex – exopinacocyte, exp – exopinacoderm, f – flagellum, m – mesohyl, mv – microvilli, n – nucleus, po – porocyte, sb – symbiotic bacteria, sc – spherulous cells.

173x194mm (300 x 300 DPI)



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Figure 15. Leucosolenia variabilis Haeckel, 1870. Mesohyl cell types and symbiotic bacteria. A amoebocytes; B - large amoeboid cell; C - granular cell; D - myocyte, inset - bundles of myofilaments; E spherulous cell; F - spherulous cells in the body wall; G - sclerocyte; H - symbiotic bacteria, morphotype 1. I – symbiotic bacteria, morphotype 2. J – symbiotic bacteria, morphotype 3. Scale bars: A - 2  $\mu$ m, B – 5  $\mu$ m, C-E – 2 µm, F – 50 µm, G – 2 µm, H-J – 1 µm. am - amoebocyte, bam - large amoeboid cell, ch – choanocytes, gc - granular cell, gr - granule, m - mesohyl, mf - myofibrils, n - nucleus, sc - spherulous cells, sp - spicule.

173x253mm (300 x 300 DPI)



Figure 16. Leucosolenia variabilis Haeckel, 1870. Spicule types in the White Sea specimen (A) and in the specimens from the British Natural History Museum collection (B-D). A – spicules from WS11731; B – spicules from BMNH 1906.12.1.40; C – spicules from BMNH 1906.12.1.50; D – spicules from the syntype BMNH 1910.1.1.421.a. d – diactines, tx – trichoxeas.

174x173mm (300 x 300 DPI)



Figure 17. Leucosolenia sp. A. External morphology and skeleton. A – general morphology (WS11752); B – skeleton of oscular rim (WS11770); C – skeleton of oscular tube (WS11770); D – skeleton of cormus (WS11752). c – cormus, d – diverticulum, o – osculum, or – oscular rim, ot – oscular tube.

174x144mm (300 x 300 DPI)



Figure 18. Leucosolenia sp. A. Spicule types, scanning electron microscopy. A – curved smooth diactines; B
 – curved spiny lanceolate diactines triactines; C – tips of diactines, I and II refer to the zones marked on A and B, white arrowheads mark spines; D – triactines; E – abnormal triactines; F – tetractine.

173x97mm (300 x 300 DPI)



Figure 19. Leucosolenia creepae sp. nov. External morphology and skeleton. A – general morphology (WS11702, holotype); B – skeleton of oscular rim (WS11762); C – skeleton of oscular tube (WS11728); D – skeleton of cormus (WS11655). c – cormus, d – diverticulum, o – osculum, oc – oscular crown, ot – oscular tube.

174x144mm (300 x 300 DPI)



Figure 20. Leucosolenia creepae sp. nov. Spicule types, scanning electron microscopy. A – spiny diactines; B – tips of diactines, I, II, III and IV refer zones marked on A, white arrowheads mark spines; C – triactines; D – abnormal triactines; E – tetractines.

173x157mm (300 x 300 DPI)

no



bacteria.

173x193mm (300 x 300 DPI)



Figure 22. Leucosolenia creepae sp. nov. Mesohyl cell types and symbiotic bacteria. A – sclerocyte; B – amoebocyte; C, D – symbiotic bacteria, morphotype 1; E, F – symbiotic bacteria, morphotype 2; G – young oocyte. Scale bars: A, B – 2 μm, C - F – 0.5 μm, G – 5 μm. am - amoeboid cell, ch – choanocytes, m – mesohyl, n – nucleus, oo – oocyte, sb – symbiotic bacteria, sp – spicule.

173x121mm (300 x 300 DPI)



Figure 23. Leucosolenia somesii (Bowerbank, 1874). External morphology and skeleton. A – general morphology (ZMA Por. 17572); B – skeleton of cormus (ZMA Por. 17572); C, D – spicules from BMNH 1956.4.26.35. c – cormus, d – diverticulum.

174x152mm (300 x 300 DPI)



Figure 24. Leucosolenia somesii (Bowerbank, 1874). ZMA Por. 17572. Spicule types, scanning electron microscopy. A – curved smooth diactines; B – straight spiny diactines; C – tips of diactines, I and II refer to the zones marked on A and B, white arrowheads mark spines; D – triactines; E – abnormal triactines; F – tetractines.

173x213mm (300 x 300 DPI)