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1	Lithothamnion species (Hapalidiales, Rhodophyta) in the Arctic and Subarctic:
2	providing a systematics foundation in a time of rapid climate change based on DNA
3	sequencing of type and recent specimens*
4	
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- 43 \*We dedicate this paper to Walter H. Adey and William J. Woelkerling for their cataloging
- 44 and scholarship on the Foslie collection and for their mentorship of the next generation of
- 45 corallinologists.

46	
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50	Running Head: Arctic and Subarctic Lithothamnion species
51	

- 52 Abstract

53 54	Coralline red algae in the non-geniculate genera Clathromorphum, Phymatolithon and
55	Lithothamnion are important benthic ecosystem engineers in the photic zone of the Arctic
56	and Subarctic. In these regions, the systematics and biogeography of Clathromorphum and
57	Phymatolithon species have mostly been resolved whereas Lithothamnion species have not,
58	until now. Seventy-three specific and infraspecific names have been given to Arctic and
59	Subarctic Lithothamnion specimens in the late 19th and early 20th century by Frans R.
60	Kjellman and particularly by Mikael H. Foslie. DNA sequences from 36 type specimens,
61	five historical specimens, and an extensive sampling of recent collections, resulted in the
62	recognition of four Arctic and Subarctic Lithothamnion species, L. glaciale, L. lemoineae,
63	L. soriferum and L. tophiforme. Three genes were sequenced, two plastid encoded, rbcL
64	and <i>psbA</i> , and the mitochondrial encoded COI-5P; <i>rbcL</i> and COI-5P segregated <i>L</i> . <i>glaciale</i>
65	from L. tophiforme but psbA did not. Partial rbcL sequences obtained from type collections
66	enabled us to correctly apply the earliest available names and to correctly place the
67	remainder in synonymy. We were unable to sequence another 22 type specimens, but all of
68	these are more recent names than those that are now applied. It is difficult to identify these
69	species solely on morpho-anatomy as they can all occur as encrusting corallines or as maerl
70	(rhodoliths). We demonstrate the importance of sequencing historical type specimens by
71	showing that the recently proposed northeast Atlantic L. erinaceum is a synonym of one of
72	the earliest published Arctic species of Lithothamnion, L. soriferum, itself incorrectly
73	placed in synonymy under L. tophiforme based on morpho-anatomy. Based on sequenced
74	specimens, we update the distributions and ecology of these species.

- **Key words**: coralline red algae; cox1; DNA barcoding; distribution; morpho-anatomy;
- *psbA*; *rbc*L; systematics; taxonomy; type collections.

## 78 Introduction

79 Coralline algae are important ecosystems engineers worldwide through the formation of 80 extensive and biodiverse cover on hard substrata and as unattached maerl (rhodoliths, 81 Freiwald & Henrich, 1994; Foster, 2001; Amado-Filho et al., 2010; Riosmena et al., 2017). 82 Approximately one-third of the total continental carbonate production takes place in 83 temperate and polar coastal waters with a significant amount coming from coralline algae 84 (Nelson, 2009). In cold-water habitats, coralline algae can live for hundreds of years 85 (Freiwald & Henrich, 1994; Halfar et al., 2013; Adey et al., 2015a), providing habitats for 86 other seaweeds (Peña et al., 2014a) and for many epibenthic and cryptic macrofauna 87 (Gagnon et al., 2012; Teichert, 2014). Over the past two decades, surveys have shown that 88 rhodolith beds are widespread in the NE Pacific (Robinson et al., 2017), NW Atlantic 89 (Gagnon et al., 2012; Copeland et al., 2013; Adey et al., 2015a), Labrador Sea and Western 90 Greeland (Jørgensbye & Halfar, 2017; Schoenrock et al., 2018a,b) and the Arctic (Teichert 91 et al., 2012; 2014); this habitat is clearly much more abundant in Arctic environments than 92 was previously assumed. 93 Adey & Steneck (2001) identified as Arctic those marine habitats ranging in temperature

94 from  $\leq$  5 °C in summer to ~ -1.5 °C in winter, and as Subarctic, those experiencing 5-15 °C

95 in summer and -1.5 to +1 °C in winter. This characterization also applies to the NW Pacific

96 Subarctic, but in the NE Pacific Subarctic, summer temperatures range from 10-15 °C and

97 winter temperatures -1.5 to  $\sim$ 5 °C (based on oceanographic conditions where the species

98 occur). The Arctic and Subarctic are warming faster than most of the world's oceans, but

99 the impact this will have on marine photosynthetic organisms is largely unknown.

100 Wassmann et al. (2011) cited 51 reports of documented changes in the Arctic marine biota

101 in response to ocean warming, but most focused on marine mammals and fish. Two of 102 these papers focused on benthic marine algae, but neither on the corallines, the group that 103 provides the dominant benthic cover of seabed habitats in the photic zone (Adey & Hayek, 104 2011). Based on research carried out elsewhere, Brodie et al. (2014), however, projected a 105 significant decrease of coralline algae in the Arctic because anthropogenic carbon dioxide 106 emissions are causing ocean acidification, which in turn is causing waters in the photic 107 zone of the Arctic to become undersaturated with aragonite. Many coralline algae are 108 susceptible to reductions in the concentration of aragonite as this can make seawater 109 corrosive to their high magnesium calcite skeletons, a response that is mediated by the rate 110 of environmental change (Kamenos et al., 2013; 2016; Martin & Hall-Spencer, 2017; Chan 111 et al., 2020). Climate-change induced permafrost thawing and snow melting at high 112 latitudes also increase freshwater runoff and coastal nutrient inputs (Walvoord & Striegl, 113 2007; Kendrick et al., 2018), which in turn can alter calcification rates and subsequent 114 coralline growth (McCoy & Kamenos, 2018; Bélanger & Gagnon, 2020) and 115 photophysiology (Schoenrock et al. 2018a). In this regard, Williams et al. (2020) observed 116 different responses among species of Clathromorphum Foslie related to their sensitivity to 117 environmental change; thus, the widely distributed C. compactum (Kjellman) Foslie might 118 expand its northern limit whereas the narrow-range C. nereostratum Lebednik is expected 119 to decline. 120 In Arctic and Subactic regions, Lithothamnion Heydrich species often dominate 121 coralline algal assemblages from the low intertidal to the lower limit of the photic zone,

122 contributing significantly to shelf carbonate budgets (Freiwald & Henrich, 1994; Nelson,

123 2009; Adey & Hayek, 2011; Teed et al., 2020). Several Lithothamnion species form maerl,

124 or branched crusts, and these structures significantly increase benthic habitat complexity

125 and biodiversity (Gagnon *et al.*, 2012; Teichert *et al.*, 2014; Jørgensbye & Halfar, 2017;

126 Schoenrock et al., 2018b). Because several Lithothamnion species (like most coralline

127 algae) also induce larval settlement and metamorphosis in invertebrates with important

128 functional roles, the genus is considered an ecosystem engineer (Steneck, 1982; Rowley,

129 1989; Pearce & Schiebling, 1990; Nelson, 2009).

130 Adey and co-workers have been studying Arctic and Subarctic subtidal benthic non-131 geniculate coralline communities for the past 50+ years, publishing numerous papers on the 132 ecology (Adey, 1964; 1965; 1966a, b; 1970a; 1971; Adey & McKibbin, 1970; Adey & 133 Adey, 1973; Adey et al., 2005), physiology (Adey, 1970b; 1973; Adey et al., 2013; 2015a) 134 and biogeography (Adey, 1966b; Adey et al., 1976; 2008; Adey & Steneck, 2001) of these 135 algae, and recently have added DNA-based taxonomic and phylogenetic studies. Thus, 136 Arctic and Subarctic species of Clathromorphum (Adey et al., 2015b), Neopolyporolithon 137 W.H.Adey & H.W.Johansen (Gabrielson et al., 2019) and Phymatolithon Foslie (Adey et 138 al., 2018) have largely been resolved, but Lithothamnion species still need clarification. 139 This is primarily due to the large number of species and infraspecific taxa that were named 140 in the late 19th and early 20th century primarily by the Norwegian corallinologist Mikael 141 Heggelund Foslie, but also by the Swedish phycologist Frans Reinhold Kjellman. Some of 142 these taxa have been placed in synonymy based on morpho-anatomy, but many are still 143 recognized (Guiry & Guiry, 2020) or are considered Incertae Sedis (Athanasiadis, 2016). 144 Studies of other coralline genera have shown increasingly that morpho-anatomy alone 145 cannot distinguish species (Sissini et al., 2014; Peña et al., 2014b; 2015a; Hernández-146 Kantún et al., 2016; Gabrielson et al., 2018). Here, we assess many of the unresolved

147 species and infraspecific taxa of Arctic and Subarctic *Lithothamnion* to provide

148 fundamental taxonomic, ecological and biogeographic knowledge of these species in the

face of the anticipated but unknown effects of climate change on the marine flora of theseregions.

151

## 152 Materials and Methods

153 Collections studied. Fifty-eight type specimens of Lithothamnion species and infraspecific 154 taxa housed in TRH and UPS as well as eleven historical specimens in TRH (herbarium 155 acronyms follow Thiers, 2020) were considered for DNA analysis (Supplementary table 156 S1, Supplementary note S1). One of us (SCL), as a guest of the Department of Botany, 157 Stockholm University, located Kjellman's type specimens in UPS that are housed in a room 158 separate from the main algal collection. This may explain why earlier investigators were 159 unable to locate them. These specimens, described by F. R. Kjellman between 1877 and 160 1889 were later received on loan by PWG; specimens described by M. H. Foslie between 161 1891 and 1908 were examined by VP or PWG. Most of the specimens had their type 162 localities along the Norwegian coast but some of them were described from Svalbard, 163 Scotland, Greenland, Canada and USA (Kjellman, 1883; 1889; Foslie, 1891; 1895; 1896; 164 1900; 1905a, b; 1908). In addition, 440 recent collections from Norway, Svalbard, 165 Greenland, and the Atlantic and Pacific coasts of Canada and USA have also been 166 sequenced (Supplementary table S2). Most of these specimens were collected subtidally in 167 coralline algal beds (known as maerl or rhodolith beds) or as crusts, and they are preserved 168 in NCU, TRH, SANT, UBC and UNB (see collection details in Supplementary table S2).

169	DNA sequencing and analyses. Herbarium material was extracted and amplified at five
170	different institutions: the Muséum National d'Histoire Naturelle, Paris (MNHN), the
171	University of North Carolina, Chapel Hill (UNC), Hartnell University (HC), the University
172	of British Columbia (UBC), and the University of New Brunswick (UNB). Extractions and
173	amplifications of types and historical collections were accompanied by negative controls at
174	every step, and they were performed separately from recent collections. At the MNHN,
175	DNA of type specimens and historical collections were extracted using QIAamp®DNA
176	Micro Kit (Qiagen S.A.S., Les Ulis, France) following the manufacturer's protocol for
177	tissues; recent collections were DNA-extracted using a NucleoSpin® 96 Tissue kit
178	(Macherey-Nagel, GmbH and Co. KG, Germany). At UNC type material and recent
179	collections were extracted following Gabrielson et al. (2011); at HC type material was
180	extracted according to Hernández-Kantún et al. (2016) following the precautionary
181	guidelines proposed by Hughey & Gabrielson (2012); at UNB recent collections were
182	extracted following Saunders & McDevit (2012); at UBC recent collections were extracted
183	following Lindstrom & Fredericq (2003). Three genes (rbcL, psbA and COI) were
184	amplified in this study. For type specimens and historical collections, <i>rbc</i> L sequences were
185	obtained with two primer combinations, F1150Cor-R1460 or F1150Cor - RbcS-Start,
186	yielding a fragment trimmed to 263 bp (1172-1434) or 293 bp (1172-1464), respectively;
187	for recent collections, <i>rbc</i> L sequences of 1383 bp were obtained with two overlapping
188	primer combinations F57-R1150 and F753-RrbcS or with primer combination F753/RrbcS-
189	Start trimmed to 691 bp (772-1464), or followed Saunders & Moore (2013) for
190	amplifications completed at UNB. For recent collections and for some type specimens and
191	historical collections, <i>psbA</i> sequences were obtained by the institutions mentioned above,

192	using the primer pairs psbA-F1/psbA-R2 and psbA-F1/psbA-600R (Yoon et al., 2002),
193	following Peña et al. (2015b) or Adey et al. (2015b). COI-5P sequences were obtained only
194	for recent collections using the primer pairs Gaz-F1/Gaz-R2 and Gaz-F1/GCorR3,
195	following Saunders & Moore (2013) or Peña et al. (2015b). PCR products were purified
196	and sequenced at MNHN by Eurofins (Eurofins Scientific, Nantes, France); at UNC
197	according to Hughey et al. (2001) and sequenced at the DNA Analysis Core Facility,
198	Center for Marine Sciences, University of North Carolina, Wilmington; and at HC by
199	Functional Biosciences, Inc. (Madison, WI, USA). Sequences were assembled and aligned
200	with the assistance of CodonCode Aligner® (CodonCode Corporation, USA) or with
201	Sequencher (Gene Codes Corp., Ann Arbor, MI, USA) and adjusted manually using
202	SeaView version 4 (Gouy et al., 2010) or using Sequence Alignment Editor
203	(http://tree.bio.ed.ac.uk/software/seal/); sequences were submitted to the Barcode of Life
204	Data Systems (BOLD projects 'NCCAB", "NGCOR" and dataset "LITHOTH1",
205	http://www.boldsystems.org; Ratnasingham & Hebert, 2007) and/or to GenBank (accession
206	numbers listed in Supplementary tables S1 and S2).
207	DNA sequencing and analyses. Three data sets were built, one for each gene (rbcL, psbA
208	and COI-5P), comprising ca. 526 sequences obtained in this study (Supplementary tables
209	S1 and S2) and supplemented with GenBank sequences publicly available for Arctic and
210	Subarctic collections of Lithothamnion as well as for other Hapalidiales genera
211	(Clathromorphum and Phymatolithon) for which relevant matches were found
212	(Supplementary table S3). As out-group we used <i>rbcL</i> and <i>psbA</i> sequences linked to the
213	generitype Lithophyllum incrustans Philippi, order Corallinales; for COI-5P we used a
214	sequence generated from the neotype of <i>Phymatolithon calcareum</i> (Pallas) Adey &

215 McKibbin, order Hapalidiales (Supplementary table S3). Phylogenetic relationships were 216 inferred with maximum likelihood (RAxML) and Bayesian inference (BI) using Mega 6.06 217 (Tamura et al., 2013), RAxML 8.1.11 (Stamatakis, 2014; available in CIPRES Science 218 Gateway, Miller et al., 2010) and MrBayes 3.2.1 (Ronquist & Huelsenbeck, 2003). Models 219 of sequence evolution were estimated using the Akaike Information Criterion (AIC) and the 220 Bayesian Information Criterion (BIC) obtained in jModeltest 2.1.3 (Darriba et al., 2012). 221 Maximum likelihood for the *rbc*L, COI-5P and *psbA* alignments were performed under a 222 generalized time-reversible with invariant sites heterogeneity model (GTR+I+G). The 223 Bayesian analyses for the *rbc*L and *psb*A alignments were performed under the same model 224 (GTR+I+G) with four Markov Chain Monte Carlo method for 10 million generations, and 225 tree sampling every 1,000 generations. 226 Distribution of Arctic/Subarctic Lithothamnion species studied. Geographic coordinates 227 were obtained by GPS for each sequenced collection and were estimated for types and 228 historical specimens using Google Earth Pro 7.3.3.7786 (© 2020 Google LLC). 229 Distribution maps were created by projecting latitude and longitude of all specimens 230 delimited for each species using QGIS3.10 (QGIS.org, 2020) with North Pole Lambert 231 Azimuthal Equa Area projection. The following shape file was used for the map 232 background: https://www.data.gouv.fr/fr/datasets/continents/ 233 234 Results

Of the type specimens (58) and historical collections (11) that we attempted to sequence,

we successfully amplified and sequenced 62% of them for *rbc*L (36 types and 5 historical

237 specimens) and *psbA* (5 types) (Supplementary table S1).

238	The <i>rbc</i> L alignment comprised 121 sequences of variable length resulting in 81
239	unique DNA sequences ranging from 205 to 313 bp, with 114 variable sites. Both ML and
240	Bayesian analyses resolved the type specimens and historical collections with moderate to
241	full support in different Hapalidiales lineages encompassing species of Lithothamnion,
242	Phymatolithon and Clathromorphum, Fig. 1). Most of the type specimens and historical
243	collections sequenced were situated within lineages represented by the lectotype of
244	Lithothamnion glaciale Kjellman (0-3 bp differences, up to 1.14 % divergence -uncorrected
245	p-distance-), followed by the lectotype of L. soriferum Kjellman (0-1 bp differences, up to
246	0.41% divergence) and then the neotype of <i>L. tophiforme</i> (0-2 bp, up to 0.68% divergence)
247	(Fig. 1). Three type specimens were placed in the genus Phymatolithon (Fig. 1,
248	Supplementary table S1): the holotype of Lithothamnion scabriusculum Foslie was
249	positioned within a clade encompassing collections of <i>P. rugulosum</i> W.H.Adey (1-3 bp
250	differences; 0.3-1% divergence); the lectotype of L. squarrulosum f. palmatifidum Foslie
251	was resolved within a clade represented by the neotype of <i>P. calcareum</i> (2 bp differences,
252	0.9% divergence); the holotype of Lithothamnion lenormandii f. squamulosum (Foslie)
253	Cotton was identical in sequence to the isotype of P. squamulosum (Foslie) W.H.Adey,
254	Hernández-Kantún & P.W.Gabrielson. Another two lectotypes (L. coalescens Foslie and L.
255	evanescens Foslie) and one historical collection of Clathromorphum circumscriptum
256	(Strömfelt) Foslie from Norway were identical in sequence to the epitype of C.
257	circumscriptum (0 bp differences, Fig. 1); the infraspecific variation within C.
258	circumscriptum ranged up to 13 bp differences (1.3% uncorrected p-distance) and it
259	increased up to 14 bp differences (1.4%) after comparison with two further $rbcL$ sequences
260	obtained from recent collections from Alaska (UBC A92115 and UBC A94120, not

included in Fig. 1). None of the types and historical collections sequenced were resolved
within the lineage of *Lithothamnion lemoineae* W.H.Adey; only three recent collections
(UBC A94112, Fig. 1, together with UBC A94113 and UBC A94121, as *Lithothamnion*sp., not included in Fig 1, Supplementary table S2) appeared to be closely related to *L. lemoineae*, showing at minimum 11 bp differences (1.15 % of divergence) between the
taxa.

267 The COI-5P alignment comprised 151 sequences that consisted of 81 unique 268 sequences ranging from 518 to 579 base pairs (bp), with 153 variable sites. The 269 phylogenetic tree obtained from the ML analysis resolved seven fully and two moderately 270 supported lineages (Supplementary fig. S1). Four of these lineages are represented by 271 recent collections of L. glaciale, L. tophiforme (Esper) Unger, L. lemoineae and L. 272 erinaceum Melbourne & J.Brodie (herein proposed as a synonym of L. soriferum Kjellman, 273 see next section below). The remaining three fully supported lineages (as L. sp. 2 to L. sp. 274 4) and another two moderately supported lineages (L. sp. 1, L. sp. 5) corresponded to recent 275 collections pertaining to five *Lithothamnion* species that did not return any relevant match 276 with publicly available GenBank sequences (Supplementary fig. S1, Supplementary tables 277 S2 and S3). The highest infralineage variation (uncorrected p-distance) was recorded in L. 278 glaciale (up to 2.07 %). 279 The *psbA* alignment comprised 421 sequences that resulted in 124 unique 280 sequences, ranging from 382 to 851 bp with 249 variable sites. The alignment encompassed 281 recent collections, five type collections generated in the present study and publicly 282 available sequences from GenBank such as the holotypes of L. erinaceum and L.

283 *lemoineae*, and the isotype of *P. rugulosum* (herein as *P. scabriusculum*, see next section of

284	taxonomic proposals) (Supplementary tables S1-S3). Both RAxML and Bayesian analyses
285	(Supplementary fig. S2) resolved our recent collections in different lineages pertaining to
286	the genera Clathromorphum (C. circumscriptum), Phymatolithon (P. squamulosum), and
287	Lithothamnion (L. lemoineae, L. tophiforme, L. glaciale and L. erinaceum [herein as L.
288	soriferum, see next paragraph of taxonomic proposals]). In agreement with results obtained
289	for <i>rbc</i> L, both RAxML and Bayesian analyses of <i>psb</i> A resolved these type collections in
290	three different genera (Supplementary fig. S2): Lithothamnion (neotype of L. glaciale f.
291	subsimplex Foslie), Clathromorphum (lectotypes of L. coalescens and L. evanescens) and
292	Phymatolithon (holotype of L. lenormandii f. squamulosa and lectotype of L. squarrulosum
293	f. palmatifida). However, one recent collection (as Lithothamnion sp., Supplementary fig.
294	S2, Supplementary table S2) was distinct in sequence from all other analyzed taxa, and it
295	did not return any relevant match with publicly available GenBank sequences. In contrast to
296	rbcL and COI analyses, the support values obtained were generally lower, particularly for
297	L. glaciale with weak support (Supplementary fig. S2).
298	Given the molecular evidence noted above, and in accordance with Article 11.4 of
299	the International Code of Nomenclature for algae, fungi, and plants (ICN, Turland et al.,
300	2018), we present the following taxa with the corresponding heterotypic synonyms from
301	the present study. Note that for each taxon below where it is stated "the lectotype is
302	narrowed" we are invoking Article 9.17 of the ICN (Turland et al. 2018).
303	
304	
305	Clathromorphum circumscriptum (Strömfelt) Foslie, 1898a: 5
306	BASIONYM: Lithothamnion circumscriptum Strömfelt, 1886: 20, pl. 1, figs. 4-8

- 307 HOMOTYPIC SYNONYMS:
- 308 *Phymatolithon compactum* f. *circumscriptum* (Strömfelt) Foslie, 1905a: 88.
- 309 *Clathromorphum compactum* f. *circumscriptum* (Strömfelt) Foslie, 1908: 11.
- 310 Lithothamnion compactum f. circumscriptum (Strömfelt) Lund, 1959: 200.
- 311 LECTOTYPE: S; seven microscope slides apparently from the original material designated
- 312 by Athanasiadis (2016: 251) as the holotype, but as Strömfelt (1886) designated syntype
- 313 localities, this material is better called a lectotype. This is a correctible error in accordance
- 314 with Art. 9.10 of the ICN (Turland *et al.*, 2018).
- 315 Comment: Adey et al. (2015b), while designating an epitype for C. circumscriptum
- 316 inadvertently omitted submitting this epitype sequence to GenBank. This has now been
- 317 rectified: GenBank XXXXXXX, an *rbc*L-263 (bp 1172-1434) sequence.
- 318
- 319 HETEROTYPIC SYNONYMS:
- 320 Lithothamnion coalescens Foslie, 1895: 162 (reprint 134), pl. 19, figs. 15-20.
- 321 *Clathromorphum coalescens* (Foslie) Foslie, 1898b: 8.
- 322 *Phymatolithon compactum* f. *coalescens* (Foslie) Foslie, 1905a: 8.
- 323 LECTOTYPE: TRH C21-3503; 12.viii.1893, leg. unknown.
- 324 TYPE LOCALITY: Inderøen, Strømmen, Trondheimsfjord, Norway.
- 325 Lectotype DNA sequences: psbA and rbcL-263 (bp 1172-1434), GenBank XXXXXXXX
- and XXXXXXXX.
- 327 Comment: Following his description, Foslie (1895) transferred the species without
- 328 comment to *Clathromorphum* (Foslie, 1898b). Later, Foslie (1905a) reduced
- 329 Clathromorphum to a subgenus of Phymatolithon and reduced C. coalescens to a form of

331	(1977) nor by Adey et al. (2015b), but was listed by Athanasiadis (2016) as a synonym of
332	C. compactum (Kjellman) Foslie. According to Woelkerling et al. (2005), the lectotype of
333	L. coalescens is a blue box with five specimens that was earlier designated by Woelkerling
334	(1993: 52) as the lectotype, noting that Foslie (1895: 163) had cited two syntype localities.
335	Upon examination, the box contained four specimens illustrated in Foslie (1895, pl. 19,
336	among figs. 15-20), one of which was sequenced, and herein the lectotype is narrowed to
337	that sequenced specimen (GenBank XXXXXXX and XXXXXX). Both the <i>rbcL</i> and
338	psbA sequences obtained are identical to GenBank sequences of C. circumscriptum
339	(voucher US 169083), which were confirmed as identical to the epitype (voucher US
340	170939, Adey et al., 2015b: 195).
341	
342	Lithothamnion durum Kjellman, 1889: 22, pl 1. figs. 3-5.
343	Clathromorphum durum (Kjellman) Foslie, 1898b: 8.
344	HOLOTYPE: UPS A-000297, vii.1877, leg. F. R. Kjellman.
345	TYPE LOCALITY: Port Clarence, Alaska, USA.
346	Holotype DNA sequence: The rbcL-263 (bp 1172-1434) sequence was obtained from the

Phymatolithon, as P. compactum f. coalescens. This name was not treated by Lebednik

- 347 holotype specimen, and over this sequence length differed by 1 bp from the epitype of
- 348 *Clathromorphum circumscriptum.* This base pair position is variable in *C. circumscriptum*,
- 349 with specimens from Iceland, Labrador, Newfoundland and Maine sharing the same single
- 350 nucleotide polymorphisms (SNP) and likewise those from Greenland and Alaska (Port
- 351 Clarance and Juneau) sharing the same SNP.

- 352 *Comment*: This synonymy was first proposed by Foslie (1900: 10) and was accepted by
- Lebednik (1977: 64); the latter noted that a fragment, apparently from the holotype, was in
- TRH. This fragment is now considered an isotype (Art. 8.3, Turland *et al.* 2018). The
- holotype illustrated by Kjellman (1889, pl. 1, fig. 3), was found in UPS, and the DNA
- 356 sequence was obtained from the specimen labeled "b" (Kjellman, 1889, pl. 1, fig. 3). We
- did not sequence the fragment in TRH.
- 358
- 359 *Lithothamnion evanescens* Foslie, 1895: 137.
- 360 *Clathromorphum evanescens* (Foslie) Foslie, 1898b: 8.
- 361 *Phymatolithon evanescens* (Foslie) Foslie, 1905a: 92.
- 362 LECTOTYPE: TRH C21-3518, iv.1889, leg. F.S. Collins.
- 363 TYPE LOCALITY: Marblehead, Massachusetts, USA.
- 364 *Lectotype DNA sequences: psbA* and *rbcL-263* (bp 1172-1434), GenBank XXXXXXXX
- and XXXXXXXX.
- 366 *Comment*: Foslie (1895: 137, Pl. 22 figs. 6, 7) cited and illustrated specimens from two
- 367 syntype localities, Marblehead, Massachusetts, USA collected by F. S. Collins and from
- 368 Mastervik, Malangen, Norway collected by himself. Foslie transferred the species twice,
- 369 first, without comment, to *Clathromorphum* (Foslie 1898b), and later to *Phymatolithon*
- after admitting that *Clathromorphum* should be considered a subgenus of the latter (Foslie,
- 371 1905a: 87). Lebednik (1977) first proposed that this species was a synonym of C.
- 372 *circumscriptum*, and this was accepted by Athanasiadis (2016). Woelkerling (1993: 87)
- 373 designated as lectotype a single specimen collected by Collins in Marblehead,
- 374 Massachusetts and illustrated by both Foslie (1895, Pl. 22, fig 6) and Printz (1929, pl. 41,

375	fig. 13). Woelkerling (1993) justified the selection of this specimen as lectotype because
376	" it was in better condition and had numerous conceptacles". Both <i>rbcL</i> and <i>psbA</i>
377	sequences obtained for this lectotype specimen are identical to GenBank sequences of C
378	circumscriptum (voucher US 169083), which was confirmed as identical to the epitype
379	(voucher US 170939, Adey et al., 2015b: 195).
380	
381	Historical collection:

- 382 TRH C20-3495, as *Lithothamnion circumscriptum*. Tamsøya, Finnmark, Norway,
- 383 31.vii.1897, no habitat data, leg. M. H. Foslie. DNA sequence: rbcL-263 (bp 1172-1434),
- 384 GenBank XXXXXXXX (Supplementary table S1). The largest fragment of the four in the
- box was sequenced.
- 386

- 388 Norway: Porsangerfjorden (Finnmark) and Krøttøya (Troms). Intertidal to subtidal (6 m
- depth), encrusting pebbles and pottery, on hard substrata and associated with maerl beds.
- 390 One specimen collected in Krøttøya had uniporate conceptacles (gametangial or

391 carposporangial). *DNA sequences: psbA* (Supplementary table S2).

- 393 Lithothamnion glaciale Kjellman, 1883: 123-127, pls. 2, 3.
- 394 LECTOTYPE, herein designated: UPS A-000202, xi-xii.1872, leg. F. R. Kjellman.
- 395 TYPE LOCALITY: Mosselbay, Spitsbergen.
- 396 *Lectotype DNA sequence: rbc*L-263 (bp 1172-1434), GenBank XXXXXXXX.

<sup>387</sup> *Recent collections*:

397	Comment: Adey (1970a) made a provisional lectotypification based on " a Spitzbergen
398	specimen (No. 241, Institute of Taxonomy, Uppsala) collected by Kjellman in 1872-1873."
399	Adey (1970a) further stated that this specimen was not one illustrated by Kjellman (1883)
400	that accompanied the original description. The ICN does not accept provisional lectotypes
401	(Art. 7.11, Turland et al. 2018). Chamberlain & Irvine (1994), repeated Adey's (1970a)
402	lectotypification, thus making it acceptable, but stated that they did not see the specimen.
403	Among Kjellman's type collections was the single individual rhodolith of L.
404	glaciale illustrated by Kjellman (1883, pls. 2, 3), with some artistic license (Supplementary
405	fig. S3A), along with a collection label stating the type locality of Mosselbay on the island
406	of Spetsbergen (Spitzbergen) and dated, November and December 1872, collected while
407	the expedition aboard the Polhem was iced in until August 1873 (Wynne, 1995). We here
408	designate this specimen from which we obtained a partial <i>rbc</i> L sequence as the lectotype of
409	L. glaciale. All other sequences of L. glaciale differ by 1 bp from the lectotype sequence,
410	including all of the type sequences of synonyms listed below.
411	Lectotype SEM observations: A cross-section through a protuberance showed radial
412	construction and a buried conceptacle (Supplementary fig. S3B). Thallus construction was
413	monomerous with elongate hypothallial cells (Supplementary fig. S3C-D). Abundant
414	fusions linked cells from adjacent perithallial filaments (Supplementary fig. S3E) and
415	secondary pit connections were absent. The epithallus was single layered and epithallial
416	cells were flared; intercalary meristematic cells (subepithallial initials) were shorter or
417	about the same length as subtending perithallial cells (Supplementary fig. S3F).
418	

419 HETEROTYPIC SYNONYMS:

- 420 Lithothamnion apiculatum f. connatum Foslie, 1895: 54, pl. 15, figs. 9-13 (as 'connata').
- 421 LECTOTYPE: TRH B20-2669, 12.vii.1893, no habitat data, leg. H.H. Gran.
- 422 TYPE LOCALITY: Drøbak, Norway.
- 423 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 424 *Comment*: Woelkerling (1993) located four of five specimens as part of the protologue of *L*.
- 425 *apiculatum* f. *connatum* and designated these lectotype. They have been illustrated by
- 426 Foslie (1895, pl. 15, figs. 9-12) and by Printz (1929, pl. 21, figs. 11-15) under the name *L*.
- 427 *colliculosum* f. *pusilla*. One of these four specimens, branched and epilithic on a pebble
- 428 (among the specimens illustrated as figs. 9-11, Foslie 1895, pl. 15) was sequenced, and
- 429 herein this lectotype is narrowed to that specimen.
- 430
- 431 Lithothamnion battersii Foslie, 1896: 1, pl. 1, figs. 1-5.
- 432 HOLOTYPE: TRH C10-3098, viii.1891, leg. E. Batters.
- 433 TYPE LOCALITY: Cumbrae, Scotland.
- 434 *Holotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 435 *Comment*: The holotype collection comprises five individuals illustrated by Foslie (1896,
- 436 pl. 1, figs. 1-5); the specimen sequenced is depicted in Foslie's (1896) figure 2. Based on
- 437 morpho-anatomy, Chamberlain & Irvine (1994: 182) and Athanasiadis (2016: 217)
- 438 correctly listed *L. battersii* as a synonym of *L. glaciale*.
- 439
- 440 *Lithothamnion colliculosum* Foslie, 1891: 43, pl. 3, fig. 1.
- 441 LECTOTYPE: TRH B11-2311, 8.ix.1890, leg. M. H. Foslie (designated by Woelkerling,
- 442 1993: 53).

- 443 TYPE LOCALITY: Skorpen, Kvænangen, Norway.
- 444 *Lectotype DNA sequence: rbcL-293* (bp 1172-1464), GenBank XXXXXXXX.
- 445 *Comment*: Foslie (1891: 43-45, pl. 3, fig. 1) described and illustrated eight individual
- 446 specimens that he ascribed to this species. Adey & Lebednik (1967: 71) could not locate
- 447 any of these specimens, and therefore Adey (1970c) designated as neotype one specimen
- 448 from Kragerø collected in 1890. Later, Woelkerling (1993: 54) located in TRH numerous
- 449 specimens with the original collection data of *L. colliculosum*, including four of the eight
- 450 individuals comprising the holotype and depicted by Foslie (1891, pl. 3, fig. 1). Because the
- 451 other four individuals comprising the holotype remain missing, Woelkerling designated the
- 452 found specimens as the lectotype of *L. colliculosum*, superseding Adey's neotype. The
- 453 lectotype is narrowed herein to the individual sequenced crust among the original
- 454 specimens depicted by Foslie (1891, pl. 3, fig. 1, bottom row, second from right). Based on
- 455 morpho-anatomy Athanasiadis (2016: 224) listed L. colliculosum as Incertae Sedis; DNA
- 456 sequence data has confirmed the placement of the species in *L. glaciale*.
- 457
- 458 Lithothamnion colliculosum f. pusillum Foslie, 1905a: 35 (as 'pusilla').
- 459 LECTOTYPE: TRH B20-2706, 12.vii.1898, leg. H. H. Gran (designated by Woelkerling
- 460 1993: 185).
- 461 TYPE LOCALITY: Drøbak, Norway.
- 462 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 463 *Comment*: We sequenced one of the 14 specimens comprising the lectotype, which is
- 464 located separately within a blue box with label "Prep. 76-77". The lectotype is narrowed
- 465 herein to the sequenced specimen. Based on morpho-anatomy Athanasiadis (2016: 224)

- 466 listed *L. colliculosum* f. *pusillum* as *Incertae Sedis*; DNA sequence data has confirmed the
- 467 placement of the species in *L. glaciale*.
- 468
- 469 Lithothamnion congregatum Foslie, 1895: 142, pl. 20, figs. 1-6.
- 470 HOMOTYPIC SYNONYM: Lithothamnion nodulosum f. congregatum (Foslie) Foslie,
- 471 1900: 13.
- 472 LECTOTYPE: TRH C7-3062, 20.vii.1894, leg. M. H. Foslie (designated by Woelkerling,
- 473 1993: 60-61).
- 474 TYPE LOCALITY: Skjørn (now Stjørna), Trondheimsfjord, Norway (Woelkerling *et al.*,
  475 2005: 424).
- 476 *Lectotype DNA sequence: rbcL-293* (bp 1172-1464), GenBank XXXXXXXX.
- 477 *Comment*: Foslie (1895: 142-144, pl. 20, figs. 1-6) described and illustrated this species
- 478 based on six individuals from a single locality, and he compared the species to two other
- 479 species that he named in the same publication, *L. dehiscens* Foslie and *L. nodulosum* Foslie.
- 480 Later, Foslie (1900) reduced *L. congregatum* to a form of *L. nodulosum*. The sequenced
- 481 specimen is illustrated in Foslie (1895, pl. 20, fig. 2), and the lectotype is narrowed herein
- 482 to that sequenced specimen. Based on morpho-anatomy Athanasiadis (2016: 224) listed *L*.
- 483 *congregatum* as *Incertae Sedis*; DNA sequence data has confirmed the placement of the
- 484 species in *L. glaciale*.
- 485

486 Lithothamnion corallioides f. saxatile Foslie, 1895: 90, pl. 16, figs. 12-23 (as 'saxatilis').

487 LECTOTYPE: TRH C9-3097, 1.viii.1894, leg. M. H. Foslie (designated by Woelkerling,

488 1993: 195).

489	<b>TYPE LOCALITY:</b>	Røberg	(now Ra	udberget).	Norway	(Woelkerling	et al.	2005:413	۱
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- 490 *Lectotype DNA sequence: rbcL-293* (bp 1172-1464), GenBank XXXXXXXX.
- 491 *Comment*: Woelkerling (1993: 195-196) located and designated as the lectotype four of the
- 492 original 12 specimens included by Foslie (1895, pl. 16, figs. 14-17) in the original
- 493 protologue of this form. The specimen sequenced had a green label "Prep. 100-101"; it
- 494 resembled the specimen illustrated in Foslie (1895, pl. 16, fig. 16). The lectotype is here
- 495 narrowed herein to this single sequenced.
- 496
- 497 *Lithothamnion dimorphum* Foslie, 1895: 68, pl. 10, figs. 1-6.
- 498 HOMOTYPIC SYNONYM: Lithothamnion fornicatum f. dimorphum (Foslie) Foslie,
- 499 1905a: 38.
- 500 LECTOTYPE: TRH B25-2773, 10.vii.1894, 0-5.5 m depth on sandy and stony bottom, leg.
- 501 M. H. Foslie (designated by Woelkerling, 1993: 75).
- 502 TYPE LOCALITY: Frøjen (now Frøya), Rottingsundet, Trondeland, Norway (Woelkerling
  503 *et al.*, 2005: 375).
- 504 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 505 *Comment*: Foslie (1895, pl. 10, figs. 1-6) illustrated six specimens belonging to this species,
- 506 but did not designate a holotype. Woelkerling (1993: 74-75) designated as the lectotype
- four of the six specimens depicted in figs. 1, 3, 5 and 6 (Foslie, 1895, pl. 10). The lectotype
- is narrowed herein to the Foslie 1895, pl. 10, fig. 3 specimen that was sequenced. Based on
- 509 morpho-anatomy, Athanasiadis (2016: 225) listed L. dimorphum as Incertae Sedis; DNA
- 510 sequence data have confirmed the placement of the species in *L. glaciale*.

- 512 Lithothamnion divergens Foslie, 1895: 96, pl. 16, figs. 43-50.
- 513 HOMOTYPIC SYNONYMS: *Lithothamnion ungeri* f. *divergens* (Foslie) Foslie, 1900: 11;
- 514 *Lithothamnion tophiforme* f. *divergens* (Foslie) Foslie, 1905a: 51.
- 515 HOLOTYPE: C11-3167, 8.ix.1890, leg. M. H. Foslie.
- 516 TYPE LOCALITY: Kvaenangen, Skørpen (now Skorpa), Norway (Woelkerling et al.,
- 517 2005: 438).
- 518 *Holotype DNA sequence*: rbcL-293 (bp 1172-1464), GenBank XXXXXXXX.
- 519 *Comment*: The holotype material comprised two boxes (one round and one quadratic) with
- seven specimens and fragments (Woelkerling, 1993; 80; Woelkerling et al., 2005: 438).
- 521 The sequenced specimen is located in the round box with the green tag "Lith. Mon. pl. 20,
- 522 f. 8" and illustrated in Foslie (1895, pl. 16, fig. 48) and Printz (1929, pl. 20, fig. 8). As
- 523 allowed by Art. 9.17 of the ICN (Turland *et al.*, 2018), the holotype is here narrowed to the
- 524 one sequenced specimen. Based on morpho-anatomy, Athanasiadis (2016: 226) listed L.
- 525 *divergens* as *Incertae Sedis*; DNA sequence data confirm the placement of the species in L.
- 526 glaciale.
- 527
- 528 Lithothamnion fornicatum f. sphaericum Foslie, 1900: 12 (as 'sphaerica').
- 529 HOLOTYPE: TRH B26-2789, 20.vii.1894, no habitat data, leg. M. H. Foslie.
- 530 TYPE LOCALITY: Skjørn, Dalsøren (now Stjørna, Daleøra), Trondheimsfjorden, Norway
- 531 (Woelkerling *et al.*, 2005: 380).
- 532 *Holotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 533 *Comment*: Foslie (1900) only provided a reference to a previously published figure (Foslie
- 534 1895, pl. 12, fig. 1) for the protologue of this form, but Woelkerling (1993: 205) considered

- this a validly published name. We sequenced the same individual rhodolith cited by Foslie
- 536 (1900) and illustrated in Foslie (1895, pl. 12, fig. 1).
- 537
- 538 *Lithothamnion fruticulosum* f. *fastigiatum* Foslie, 1895: 46, pl. 5.
- 539 LECTOTYPE: TRH B25-2777, 6.vii.1894, leg. M. H. Foslie (designated by Woelkerling,
- 540 1993: 91).
- 541 TYPE LOCALITY: Bejan (now Beian), Beiskjaeret, Norway (Woelkerling *et al.*, 2005:
- 542 377).
- 543 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 544 Comment: Foslie (1895, pl. 5, figs. 1-7) illustrated seven individual rhodoliths that
- according to Woelkerling (1993) came from two collections from neighboring localities on
- 546 successive days. Woelkerling (1993: 91) designated as the lectotype collection nine
- 547 specimens comprising two of those illustrated in the protologue (Foslie, 1895, pl. 5, figs. 5
- and 7) and seven other specimens. The lectotype is narrowed herein to the sequenced
- specimen that is marked with label "nr. 2", which is cited as part of the lectotype in
- 550 Woelkerling *et al.* (2005: 377).
- 551
- 552 Lithothamnion fruticulosum f. flexuosa Foslie, 1895, pl. 7, figs. 1-3.
- 553 LECTOTYPE: TRH B27-2805, 15.viii.1890, leg. M. H. Foslie (designated by Woelkerling,

554 1993: 95).

- 555 TYPE LOCALITY: Tromsø, Norway.
- 556 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.

557	Comment: Foslie (1895) based this form on collections from several localities in Norway,
558	but did not designate a type. Woelkerling (1993) designated the specimen illustrated by
559	Foslie (1895, pl. 7, fig. 3) as the lectotype of this form. According to Woelkerling et al.
560	(2005), the lectotype has two green tags, and this was the specimen from which the DNA
561	sequence was obtained.
562	
563	Lithothamnion fruticulosum f. glomeratum Foslie, 1895: 46, pl. 4, fig. 3 (as 'glomerata').
564	LECTOTYPE: TRH B8-2153, 12.vi.1892, leg. unknown (designated by Printz, 1929, pl.
565	22, fig. 5 legend).
566	TYPE LOCALITY: Lyngø (now Lyngøya), near Tromsø, Norway (designated by
567	Woelkerling, 1993: 108, further information in Woelkerling et al. 2005: 288).
568	Lectotype DNA sequence: rbcL-293 (bp 1172-1464), GenBank XXXXXXXX.
569	Comment: Foslie (1895) cited specimens from two localities in Norway, Lyngø and Vardø,
570	but did not designate a type. Printz (1929) designated as lectotype the one rhodolith
571	illustrated by Foslie (1895: 46, pl. 4, fig. 3) and this is the specimen sequenced in the
572	current study.

- Lithothamnion gracilescens Foslie, 1895: 87, pl. 15, figs. 20-27, nom. illeg.
- HOMOTYPIC SYNONYM: Lithothamnion nodulosum f. gracilescens Foslie, 1900: 13,

nom. illeg.

- LECTOTYPE: TRH C6-3037, 6.vi.1894, leg. M. H. Foslie (designated by Woelkerling,
- 1993: 109; further information in Woelkerling et al., 2005: 419).

- 579 TYPE LOCALITY: Rotvold [now Rotvoll], Trondsheimsfjord, Norway (Woelkerling *et*580 *al.*, 2005: 419).
- 581 *Lectotype DNA sequence*: rbcL-293 (bp 1172-1464), GenBank XXXXXXXX.
- 582 *Comment*: Foslie (1895) proposed *Lithothamnion gracilescens* for the coralline that Unger
- 583 (1858: 19) had called *Lithothamnion byssoides*, but Kjellman (1883: 120) already had
- named this entity *Lithothamnion ungeri*. Foslie (1895: 90) listed three syntype localities,
- 585 Madal, Dröbak and Rotvold, and from the last location illustrated eight individuals (Foslie
- 586 1895, pl. 15, figs. 20-27). Later, Foslie (1900) without comment reduced L. gracilescens to
- 587 a form of L. nodulosum as L. nodulosum f. gracilescens. The specimen sequenced
- 588 corresponds to Foslie (1895, pl. 15, fig. 24), to which the lectotype is narrowed. Based on
- 589 morpho-anatomy Athanasiadis (2016: 228) listed *L gracilescens* as *Incertae Sedis*; DNA
- 590 sequence data have confirmed the placement of the species in *L. glaciale*.
- 591
- 592 Lithothamnion intermedium Kjellman, 1883: 127, pl. 4, figs. 1-10.
- 593 HOMOTYPIC SYNONYMS: Lithothamnion fruticulosum f. intermedium (Kjellman)
- 594 Foslie, 1895: 46; *Lithothamnion ungeri* f. *intermedium* (Kjellman) Foslie, 1898b: 5.
- 595 LECTOTYPE herein designated: UPS A648805, vi.1875, leg. F. R. Kjellman.
- 596 TYPE LOCALITY: Carlsö (now Karlsøy), Tromsø, Norway
- 597 *Lectotype DNA sequence*: rbcL-263 (bp 1172-1434), GenBank XXXXXXXX.
- 598 Comment: Kjellman (1883) did not designate a type specimen for his new species. In UPS
- there appears to be only one individual rhodolith with a label in Kjellman's hand and with
- 600 the specific locality of Karlsøy cited in the protologue of *L. intermedium*, although two

- 601 individuals are illustrated by Kjellman (1883, pl. 4, figs. 1, 2). Thus, we designate UPS
- 602 A648805 as the lectotype.
- 603
- 604 Lithothamnion soriferum f. globosum Foslie, 1891: 41, pl. 3, fig. 3 (as 'globosa').
- 605 LECTOTYPE: C11-3142, 20.vi.1882, no habitat data, leg. M. H. Foslie (designated by
- 606 Woelkerling, 1993: 107).
- 607 TYPE LOCALITY: Honningsvaag (now Honningsvåg), Finnmark, Norway (Woelkerling
- 608 *et al.*, 2005: 434).
- 609 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 610 Comment: Woelkerling (1993: 107) located two of the four specimens illustrated by Foslie
- 611 (1891, pl. 3, fig. 3). The specimen sequenced is illustrated in Foslie (1891, pl. 3, fig. 3,
- 612 second specimen from the left), and it is this specimen to which the lectotype is narrowed
- 613 herein. Athanasiadis (2016: 234) lists as *Incertae Sedis*; DNA sequence data have
- 614 confirmed the placement of the species in *L. glaciale*.
- 615
- 616 *Lithothamnion tusterense* Foslie, 1905a: 65.
- 617 HOLOTYPE: TRH C9-3089, 10.viii.1898, leg. M. H. Foslie.
- 618 TYPE LOCALITY: Tusteren (now Tustna), Kristiansund, Norway (Woelkerling et al.,
- 619 2005: 428).
- 620 Holotype DNA sequence: rbcL-293 (bp 1172-1464), GenBank XXXXXXXX.
- 621 Comment: Woelkerling (1993: 229) considered the material in the Foslie herbarium
- 622 illustrated by Printz (1929, pl. 22, figs. 6-13) to be the holotype, and we agree. The
- 623 specimen sequenced is marked with a green tag "pl. 27, fig. 9", but the specimen

- 624 corresponds to the plate 22, figure 9 in Printz (1929), as was noted by Woelkerling *et al.*
- 625 (2005: 428). Based on morpho-anatomy, Athanasiadis (2016: 235) listed *L. tusterense* as
- 626 *Incertae Sedis*; DNA sequence data have confirmed the placement of the species in L.
- 627 glaciale.
- 628
- 629 Lithothamnion vardoeense Foslie, 1905b: 3 (as vardöense).
- 630 LECTOTYPE: TRH C8-3077, 6.ix.1897, leg. M. H. Foslie. Woelkerling (1993: 233)
- designated as lectotype the specimens illustrated by Printz (1929, pl. 33, figs. 12, 13 and
- 632 15).
- 633 TYPE LOCALITY: Svolvær, Lofoten, Norway.
- 634 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 635 Comment: Foslie (1905b) based this species on two collections, one dead, excavated from
- 636 Vardø, Norway, the other living from Lofoten, Svolvær, Norway. Adey & Lebednik (1967:
- 637 77) examined material in TRH under this name, but did not designate a lectotype; this was
- done by Woelkerling (1993: 233). The specimen sequenced corresponds to the one depicted
- 639 in Printz (1929, pl. 32, fig. 12), and the lectotype is narrowed herein to this specimen.
- 640 Based on morpho-anatomy, Athanasiadis (2016: 223) listed *L. vardoense* as *Incertae Sedis*;
- 641 DNA sequence data have confirmed the placement of the species in *L. glaciale*.
- 642
- 643 *Historical collections:*
- 644 TRH C12-3177, as *Lithothamnion tophiforme*. Sukkertoppen, Greenland, no date, leg.
- 645 Petersen. DNA sequence: rbcL-263 (bp 1172-1434), GenBank XXXXXXXX.

- 646 TRH B10-2305, as Lithothamnion glaciale f subfastigiatum (as 'subfastigiata').
- 647 Bekkarfjord, Alten [now Alta], Norway, 21.viii.1897, leg. M. H. Foslie. DNA sequence:
- 648 *rbc*L-293 (bp 1172-1464), GenBank XXXXXXX. *Comment*: The largest fragment in the
- 649 collection was sequenced. The sequence was identical to the lectotype sequence of
- 650 *Lithothamnion colliculosum* f. *pusillum*, shown herein as a heterotypic synonym of *L*.
- 651 glaciale. L. glaciale f. subfastigiatum is considered a superfluous substitute name for
- 652 *Lithothamnion varians* f. *varians* Foslie (Woelkerling *et al.*, 2005: 306); the lectotype of *L*.
- 653 varians f. varians (TRH C23-3649) was DNA-extracted but failed to amplify (see
- 654 Supplementary note S1).
- 655

## 656 Infraspecific taxa of *Lithothamnion glaciale* confirmed as *L. glaciale*.

- 657 *Lithothamnion glaciale f. subsimplex* Foslie, 1905a: 27.
- 658 NEOTYPE: TRH B9-2255, 22.vi.1900, leg. E. Bay. Woelkerling (1993: 215) designated as
- neotype the specimen illustrated by Printz (1929, pl. 23, fig. 5).
- 660 TYPE LOCALITY: the southern coast of Ellesmereland, Havnefjorden; Northwest
- 661 Territories, Canada (Woelkerling 1993: 215; Woelkerling *et al.*, 2005: 300).
- 662 *Neotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 663 Comment: Athanasiadis (2016) listed this taxon as a synonym of L. glaciale, but with a
- 664 query (?); DNA sequence data have confirmed this synonymy. The neotype sequence was
- 665 identical to the lectotype sequence of *Lithothamnion colliculosum* f. *pusillum*, shown herein
- as a heterotypic synonym of *L. glaciale*.
- 667

669 Lithothamnion soriferum Kjellman, 1883: 117, pl. 1, figs 1-19.

- 670 LECTOTYPE: herein designated, UPS A648809, viii.1876, leg. F. R. Kjellman.
- 671 TYPE LOCALITY: Maasö (now Måsøy), Finnmark, Norway.
- 672 Lectotype DNA sequences: rbcL-263 (bp 1172-1434), GenBank XXXXXXX,
- 673 XXXXXXXX and XXXXXXXX.

674 *Comment*: Kjellman (1883) did not designate a type specimen for his new species. As with

- 675 L. glaciale, one of us (SCL) located in UPS type material of L. soriferum that agrees, with
- 676 some artistic license, to the three entire specimens illustrated by Kjellman (1883, pl. 1, figs.
- 1-4). In the figure legends, Kjellman refers to the figure 1 specimen as young (ungt), the
- 678 figure 2 specimen as older (äldre) and seen from above (ofvanifrån), the figure 3 specimen,
- the same (samma) specimen (as in figure 2) seen from below (underifrån), and the figure 4
- 680 specimen as full grown (fullvuxet). An *rbc*L-263 sequence was obtained from each of these
- specimens, and the sequences are identical to each other. Kjellman (1883: 120) listed six
- 682 syntype localities Tromsö, Carlsö, Maasö, Magerö, Honningsvaag and Lebesby, the last
- two localities based on specimens sent to Kjellman by Foslie; the syntype corresponding to
- the latter locality –Lebesby– was also sequenced (TRH C13-3185, see below). The three
- specimens in UPS were found in a single envelope (Supplementary fig. S4A) with the
- 686 locality "Norway. Finnmark: Hammerfest, Måsö [Måsöya]". Following Article 9.3
- 687 (Turland et al. 2018) we herein designate these three specimens as the lectotype of L.
- 688 soriferum.

689 The identical *rbc*L-263 sequences of all three UPS specimens are an exact match to
690 GenBank sequences of the recently described *Lithothamnion erinaceum* (Melbourne *et al.*,

691 2017). Following Article 11.4 of the ICN (Turland *et al.*, 2018), the correct name is the

- 692 combination of the final epithet of the earliest legitimate name of the taxon at the same
- 693 rank. Consequently, L. soriferum has nomenclatural priority over L. erinaceum, and it is the
- 694 correct name for this taxon. Further information about the heterotypic synonym *L*.
- 695 *erinaceum* is in the entry below.
- 696 SYNTYPE: TRH C13-3185, 2.viii.1882, leg. M. H. Foslie (identified by Kjellman,
- 697 Woelkerling *et al.* 2005: 440).
- 698 TYPE LOCALITY: Lebesby, Finmarken (now Finnmark), Norway (Woelkerling et al.,
- *699* 2005: 441).
- 700 Syntype DNA sequence: rbcL-263 (bp 1172-1434), GenBank XXXXXXXX.
- 701 *Comment:* The specimen sequenced is the one that according to Woelkerling *et al.* (2005:
- 441) is illustrated in Foslie (1891, pl. 3, fig. 3, as *Lithothamnion soriferum* f. *globosa*).
- 703 *Lectotype SEM observations*: A vertical section through a protuberance showed the radial
- construction (Supplementary fig. S4B-D) and an extensive perithallus. Thallus construction
- 705 was monomerous with elongate hypothallial cells (Supplementary Fig. S4E). Fusions
- occurred between cells of adjacent perithallial filaments (Supplementary fig. S4F), and
- secondary pit connections were absent. There was a single layer of epithallial cells, and
- ros each epithallial cell had flared walls (Supplementary Fig. S4F-H). A single layer of
- intercalary meristematic cells (subepithallial initials) was composed of cells shorter than or
- as long as subtending perithallial cells (Supplementary fig. S4F-H).
- 711
- 712 HETEROTYPIC SYNONYMS:
- 713 Lithothamnion breviaxe Foslie, 1895: 44, pl. 2, figs. 1-2.

- 714 LECTOTYPE: TRH B12-2327, 3.viii.1887, leg. M. H. Foslie (designated by Adey &
- 715 Lebednik 1967: 63).
- 716 TYPE LOCALITY: Kjelmø (now Sør-Varanger, Kjelmøya), Sydvaranger, Finnmark,
- 717 Norway (Woelkerling *et al.* 2005: 310).
- 718 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 719 Comment: Foslie (1895: 44, pl. 2, figs. 1-2) described this species based on several
- specimens collected at one locality (Kjelmø, Norway), but illustrated only two. We
- obtained an *rbc*L sequence from the designated lectotype (Foslie 1895, pl. 2, fig. 1) and it
- was identical to the lectotype sequence of *L. soriferum*. Based on morpho-anatomy,
- 723 Athanasiadis (2016: 223) listed *L. breviaxe* as *Incertae Sedis*; DNA sequence data have
- confirmed the placement of the species in synonymy with *L. soriferum*.
- 725
- *Lithothamnion erinaceum* Melbourne & J.Brodie, 2017: 7, figs. 3, 7-9, 11.
- 727 HOLOTYPE: BM 001150576, 13.x.2014, leg. A. Mogg.
- 728 TYPE LOCALITY: Loch Creran, Oban, Scotland (Melbourne *et al.*, 2017).
- 729 Holotype DNA sequences: GenBank KX828452 (psbA) and KX828509 (COI-5P)
- (Melbourne *et al.*, 2017); GenBank MH697546 and MH697547 (*rbc*L, Hofman & Heesch
- 731 2018).
- 732 *Comment*: According to Melbourne *et al.* (2017), collections from Northern Ireland,
- 733 Iceland, Norway and British Columbia provided in Pardo et al. (2014) as Lithothamnion sp.
- 2 corresponded to *L. erinaceum*. Based on DNA sequences, these collections are also
- assigned to *L. soriferum*, as well as the remaining specimens from the UK identified as
- 736 *Lithothamnion* sp. in Melbourne *et al.* (2017, Table S1). Additionally, GenBank records

from Norway identified as *L. erinaceum* (specimens "NCCA" in Supplementary Table S3)

738 correspond to *L. soriferum* (Anglés d'Auriac *et al.*, 2019).

739

740 *Lithothamnion fornicatum* Foslie, 1891: 38, pl. 2 (bottom specimen).

LECTOTYPE: TRH B21-2712, 20.ix.1890, leg. unknown (designated by Adey & Lebednik
1967: 71).

743 TYPE LOCALITY: Melangen [now Malangen], Mestervik, Tromsø county, Norway

744 (Woelkerling *et al.*, 2005: 366).

745 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.

746 *Comment*: Foslie (1891) described and illustrated three specimens from the same locality.

Adey (1970c) noted that the lectotype, designated by Adey & Lebednik (1967: 71),

comprised two sets of specimens and that the selected set had a specimen pictured in the

original description, but Adey (1970c) did not indicate which of the three originally

pictured specimens is the designated lectotype. Woelkerling (1993: 97) and Woelkerling et

751 al. (2005: 366) provided information about the lectotype specimen illustrated in Foslie

752 (1891, pl. 2, bottom specimen) and marked with green tag "Præp. 151". Based on morpho-

anatomy, Athanasiadis (2016: 228) listed *L. fornicatum* as *Incertae Sedis*; DNA sequence

data have confirmed the placement of the species in synonymy with *L. soriferum*.

755

756 Lithothamnion granii (Foslie) Foslie, 1900: 11.

BASIONYM: *Lithothamnion flabellatum* f. *granii* Foslie, 1895: 98, pl. 17, figs. 1-7, pl. 22,
fig. 1.

- 759 HOMOTYPIC SYNONYMS: Lithothamnion glaciale var. granii (Foslie) Rosenvinge,
- 760 1917: 222, figs. 138-142, pl. 3, fig. 4; pl. 4: figs. 1-4; Lithothamnion glaciale f. granii
- 761 (Foslie) Foslie 1905a: 10.
- 762 LECTOTYPE: TRH C10-3114, 12.vii.1893, no habitat data, leg. H. H. Gran (designated by
- 763 Adey & Lebednik, 1967: 78).
- 764 TYPE LOCALITY: Drøbak, Norway.
- *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 766 *Comment*: The specimen sequenced is located in a large, round blue box; the label indicates
- 767 illustrations in Foslie (1895, pl. 17, fig. 3 and pl. 22, fig. 1) and Printz (1929, pl. 18, fig.
- 15). Foslie (1895: 98, pl. 17, figs. 1-7) described this new form, listed numerous syntype
- 769 localities and illustrated seven specimens. Later, Foslie (1900) elevated this form to specific
- rank. Adey & Lebednik (1967) lectotypified the specimens collected by Gran, who is
- honored by the form name. Subsequent to the original description, Foslie named five
- additional forms, f. robustum (Foslie, 1895), f. grandifrons, f. sphaericum (Foslie, 1900),
- and f. *obcrateriforme* and f. *tuberculatum* (Foslie, 1905a). We have not sequenced type
- material of any of these forms, hence they are not listed as synonyms. Based on morpho-
- anatomy, Athanasiadis (2016: 228) listed L. granii as Incertae Sedis; DNA sequence data
- have confirmed the placement of the species in synonymy with *L. soriferum*.
- 777
- 778 Lithothamnion nodulosum Foslie, 1895: 144, pl. 21, figs. 1-6.
- HOMOTYPIC SYNONYM: *Lithothamnion nodulosum* f. *typicum* Foslie, 1905a: 62, *nom. inval.*

- 181 LECTOTYPE: TRH C5-2999, 18.vii.1894, leg. M. H. Foslie (designated by Woelkerling,
  1993: 158).
- 783 TYPE LOCALITY: Brækstad (now Brekstad), Trondheimsfjorden, Norway (Woelkerling
  784 *et al.*, 2005: 414).
- *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 786 *Comment*: We sequenced one of the three rhodoliths comprising the lectotype from
- 787 Brekstad, Norway, and the sequence was identical to *L. soriferum*. The specimen sequenced
- is preserved in a round, red box; it is illustrated in Printz (1929, pl. 25, fig. 2) and not in pl.
- 789 21 as marked on the box (see Woelkerling *et al.*, 2005: 414). The lectotype is narrowed
- herein to the sequenced specimen. Based on morpho-anatomy, Athanasiadis (2016: 228)
- 191 listed *L. nodulosum* as *Incertae Sedis*; DNA sequence data have confirmed the placement of
- the species in synonymy with *L. soriferum*.
- 793
- *Lithothamnion sonderi* f. *sublaevigatum* Foslie, 1905a: 24.
- 795 HOLOTYPE: TRH B15-2426, 21.vii.1902, leg. M. H. Foslie.
- 796 TYPE LOCALITY: The islet in front of the lighthouse, Røvær, Norway (Woelkerling *et al.*
- 797 2005: 323).
- 798 *Holotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 799 Comment: The partial rbcL sequence of the holotype is identical over its entire length to the
- 800 corresponding type sequences of *L. soriferum*. This is the only type specimen of a synonym
- 801 of *L. soriferum* that is an encrusting, epilithic coralline; all others are rhodoliths. The
- 802 holotype material comprised six epilithic crusts, with the piece illustrated in Printz (1929,
- 803 pl. 4, fig. 8) in a separate box (Woelkerling, 1993: 211). We sequenced the specimen

804 located separately within a blue, round box that corresponds to the piece illustrated in plate
805 4, figure 8 (Printz (1929).

806

807 *Lithothamnion soriferum* f. *divaricatum* Foslie, 1891: 41, pl. 3, fig. 2.

808 LECTOTYPE: TRH C11-3161, 5.viii.1882, leg. M. H. Foslie (designated by Woelkerling,

809 1993: 79).

810 TYPE LOCALITY: Tromsø, Norway.

811 *Lectotype DNA sequence: rbcL-293* (bp 1172-1464), GenBank XXXXXXXX.

812 Comment: Woelkerling et al. (2005: 437) noted that there was one specimen in the

813 lectotype collection with a green tag: "Lith. Mon. pl. 20, f. 9" (corresponding to Printz

814 1929, pl. 20, fig. 9), and another specimen with the annotation, "Specimen matches Foslie

815 1891a, pl. 3, fig. 2, lower left". Previously, Woelkerling (1993: 79) designated as lectotype

816 element the collection containing this latter specimen because no further type material had

817 been found at that time. The specimen selected for sequencing was the specimen with the

green tag "Lith. Mon. pl. 20, f. 9" (corresponding to Printz, 1929, pl. 20, fig. 9) among the

819 five rhodoliths contained in the lectotype collection. The lectotype is narrowed herein to the

820 sequenced specimen. Based on morpho-anatomy, Athanasiadis (2016: 228) listed

821 L.soriferum f. divaricatum as Incertae Sedis; DNA sequence data have confirmed the

822 placement of the taxon in *L. soriferum*.

823

*Lithothamnion uncinatum* Foslie, 1895: 154, pl. 19, figs. 11-14.

825 HOMOTYPIC SYNONYMS: Lithothamnion calcareum f. uncinatum (Foslie) Foslie,

826 1897: 9; *Lithothamnion norvegicum* f. *uncinatum* (Foslie) Foslie, 1900: 13.

- 827 HOLOTYPE: TRH C3-2998, 1890, leg. unknown.
- 828 TYPE LOCALITY: Kragerø, Norway.
- 829 Holotype DNA sequence: rbcL-293 (bp 1172-1464), GenBank XXXXXXXX.
- 830 *Comment*: According to Woelkerling (1993: 231), the holotype collection comprises
- 831 several rhodolith specimens illustrated in Foslie (1895, pl. 19, figs. 11-14). The specimen
- depicted in Foslie (1895, pl. 19, fig. 11) was sequenced, and is identical to *L. soriferum*.
- 833 Athanasiadis (2016: 231) listed this name as a synonym of *Lithothamnion norvegicum*,
- 834 which he considers *Incertae Sedis* (see below).
- 835
- 836 *Historical collection*:
- 837 UPS A648806, Mestervik, Tromsø, Norway, 20.ix.1890, leg. Foslie. DNA sequence: rbcL-
- 838 263 (bp 1172-1434), GenBank XXXXXXX (Supplementary table S1).
- 839
- 840 Lithothamnion tophiforme (Esper) Unger, 1858: 21, pl. 5, fig. 14.
- 841 BASIONYM: Millepora polymorpha f. tophiformis Esper, 1789: pl. XV [Millepora]
- 842 NEOTYPE: TRH C12-3179, no date, no habitat data, leg. C. Ryberg (designated by Adey,
- 843 1970c).
- 844 TYPE LOCALITY: Julianehaab, Greenland.
- 845 *Neotype DNA sequence: rbc*L-263 (bp 1172-1464), GenBank XXXXXXXX.
- 846 *Comment*: We sequenced the neotype designated by Adey *et al.* (2005), and the GenBank
- sequences used in Adey *et al.* (2015b) are in agreement with the neotype sequence.
- 848
- 849 HETEROTYPIC SYNONYMS:

- 850 Lithothamnion alcicorne Kjellman, 1883: 121, pl. 5, figs. 1-8.
- 851 HOMOTYPIC SYNONYMS:
- Lithothamnion soriferum f. alcicorne Foslie, 1891: 41, pl. 3, fig. 4.
- Lithothamnion tophiforme f. alcicorne Foslie, 1895: 147 (as 'alcicornis').
- LECTOTYPE: TRH C13-3203 (designated herein), 5.viii.1882, leg. M. H. Foslie.
- 855 TYPE LOCALITY: Tromsø, Norway.
- 856 *Lectotype DNA sequence: rbc*L-293 (bp 1172-1464), GenBank XXXXXXXX.
- 857 *Comment*: The *rbc*L sequence obtained is identical to the neotype sequence of *L*.
- tophiforme. Kjellman (1883: 122) stated that this species was from Tromsø and that Foslie
- was the collector. Woelkerling *et al.* (2005: 443) called TRH C13-3203 syntype material.
- 860 The specimen selected for sequencing and designated herein as lectotype was illustrated by
- Printz (1929, pl. 19, fig. 11). Athanasiadis (2016: 221) cited this species as *Incertae Sedis*;
- 862 DNA sequence data have confirmed the placement of the species in synonymy with *L*.
- 863 tophiforme.
- 864
- 865 *Historical collection:*
- TRH C3-2948, Viprandsund, Haugesund, Norway, 17.v.1897, leg. M. H. Foslie.
- 867 DNA sequence: rbcL-293 sequence (bp 1172-1464), GenBank XXXXXXX; identical to
- 868 the neotype sequence of *L. tophiforme*.
- 869 *Comment:* This collection is topotype material of *Lithothamnion norvegicum* (Areschoug)
- 870 Kjellman (1883: 122, basionym: Lithothamnion calcareum var. norvegicum Areschoug,
- 871 1875: 4). The box contains a large collection of rhodoliths but the specimen sequenced is
- 872 located separately within a small box marked "Prep. 522".

873	Areschoug (1875:4) cited Wittrock as the collector L. calcareum var. norvegicum,
874	but he did not illustrate material, nor indicate a type. We have been unable to locate any
875	material in either L or UPS where other Areschoug material is located. Eleven collections
876	of Foslie's in TRH are listed under L. norvegicum (Woelkerling et al. 2005: 407-408), but
877	only the sequenced specimen is topotype material. A collection by Wittrock is in TRH (C3-
878	2985) and illustrated in Printz (1929, pl. 16, figs. 37-38) but the date of collection and
879	habitat data are lacking, making it difficult to know if this is type material. Woelkerling &
880	Verheij (1995: 67) noted syntype material is present in L. Both the material in TRH and in
881	L needs to be sequenced to understand the correct application of L. norvegicum.
882	
883	"Lithothamnion" species and infraspecific taxa that belong in Phymatolithon
884	
885	
005	Phymatolithon calcareum (Pallas) W.H.Adey & McKibbin
886	Phymatolithon calcareum (Pallas) W.H.Adey & McKibbin Lithothamnion squarrulosum f. palmatifida Foslie, 1899: 6.
886	Lithothamnion squarrulosum f. palmatifida Foslie, 1899: 6.
886 887	Lithothamnion squarrulosum f. palmatifida Foslie, 1899: 6. LECTOTYPE: TRH C1-2892, 12.v.1893, leg. L. K. Rosenvinge; designated by
886 887 888	<i>Lithothamnion squarrulosum f. palmatifida</i> Foslie, 1899: 6. LECTOTYPE: TRH C1-2892, 12.v.1893, leg. L. K. Rosenvinge; designated by Woelkerling <i>et al.</i> (2005: 398).
886 887 888 889	Lithothamnion squarrulosum f. palmatifida Foslie, 1899: 6. LECTOTYPE: TRH C1-2892, 12.v.1893, leg. L. K. Rosenvinge; designated by Woelkerling <i>et al.</i> (2005: 398). TYPE LOCALITY: Fladen, østl. (= Eastern) Kattegat, Denmark (Woelkerling <i>et al.</i> 2005:
886 887 888 889 890	Lithothamnion squarrulosum f. palmatifida Foslie, 1899: 6. LECTOTYPE: TRH C1-2892, 12.v.1893, leg. L. K. Rosenvinge; designated by Woelkerling <i>et al.</i> (2005: 398). TYPE LOCALITY: Fladen, østl. (= Eastern) Kattegat, Denmark (Woelkerling <i>et al.</i> 2005: 398).

unwritten green label was selected for DNA sequencing. The *rbc*L sequence differed by 2

- bp from the neotype sequence of *P. calcareum*, while the *psbA* sequences were identical
- 896 (neotype specimen BM000712373, Supplementary table S3).
- 897
- 898 *Phymatolithon scabriusculum* (Foslie) V.Peña, P.W.Gabrielson & Hughey comb. nov.
- 899 BASIONYM: Lithothamnion scabriusculum Foslie, 1895: 170.
- 900 HOLOTYPE: TRH C20-3502, 5-10 fathoms, 2.viii.1887, leg. M. H. Foslie.
- 901 *Type locality*: Kjelmø (now Kjelmøya), Finnmark, Norway (Woelkerling *et al.*, 2005: 493).
- 902 DNA sequence: rbcL-293 (bp 1172-1464), GenBank XXXXXXXX.
- 903 Comment: The largest fragment preserved in a small box marked '520' (further information
- 904 in Woelkerling et al. 2005: 493) was selected for DNA sequencing. Identical rbcL
- sequences were obtained independently by VP and by PWG/JRH, which differed by 1-3 bp
- from the GenBank sequences of *P. rugulosum* (vouchers US 170942 and BM000659095)
- 907 collected in Gulf of Maine and Helgoland, Germany, respectively (Adey et al. 2015b). By
- 908 the rule of priority (Art. 11.4 of the ICN, Turland *et al.*, 2018), this name must be adopted
- 909 for *P. rugulosum*.
- 910
- 911 HETEROTYPIC SYNONYM:
- 912 *Phymatolithon rugulosum* W.H.Adey, 1964: 381, figs. 15-20, 27-29, 35-36, 39-44, 51-64.
- 913 HOLOTYPE: Adey 61-41A-3 in MICH, 2.xi.1961, 3-5 m depth, leg. W. H. Adey.
- 914 TYPE LOCALITY: Merchant Island, East Penobscot Bay, Maine.

- 916 Phymatolithon squamulosum (Foslie) W.H.Adey, Hernandez-Kantun &
- 917 **P.W.Gabrielson**

- 918 BASIONYM: Lithothamnion squamulosum Foslie, 1895:183.
- 919 HOLOTYPE: TRH B5-1962, vii. 1894, leg. P. Boye.
- 920 TYPE LOCALITY: Sogn, Sulen (now Sula), indre (= inner), Stensund (now Steinsund),
- 921 Norway (Woelkerling *et al.* 2005: 265).
- 922 Holotype DNA sequences: psbA and rbcL-293 (bp 1172-1464), GenBank XXXXXXXX
- 923 and XXXXXXX, respectively.
- 924 *Comment*: The collection comprises a larger box with two smaller boxes, one square and
- 925 one round (Woelkerling *et al.*, 2005: 265). The specimen in the rounded box with the
- annotation "Prep. 65" was sequenced. Recently this taxon was transferred to *Phymatolithon*
- 927 according to the molecular data obtained from an isotype preserved in BM (BM000044670,
- Box 434, Adey *et al.*, 2018). The DNA sequencing of the holotype confirms this result; the

929 *rbc*L sequences of both type collections were identical.

930

931 *Recent collections*:

Norway: Krøttøya (Troms) and Averøya. Intertidal to subtidal (9 m depth), encrusting on
cobble, pebbles and shells, and on hard substrata. Specimens with uniporate (gametangial
or carposporangial) and multiporate conceptacles. *DNA sequences: psbA* (Supplementary
table S2).

936

937 N.B. We have not dealt with *Lithothamnion sonderi* Hauck in this paper as we have not

- 938 sequenced the type material, even though the species is thought to be widespread in
- 939 crustose forms at low light levels in the NE Atlantic, from Nordland (Norway) to North
- 940 Spain (Chamberlain & Irvine, 1994).

941

## 942 **Discussion**

Since DNA sequences from the first type specimens of geniculate (Gabrielson *et al.*, 2011)

and non-geniculate (Sissini et al., 2014) corallines were published, it has become

945 increasingly clear that the primary method to unequivocally apply a historical name is to

obtain DNA sequences from the type material to compare them to sequences from other

947 historical or more recently collected specimens (Martone *et al.*, 2012; Hind *et al.*, 2014a;

948 2014b; 2015; Hernández-Kantún et al., 2015a; 2016; Richards et al., 2017; 2018;

949 Gabrielson et al., 2018; 2019; Peña et al., 2018; Jeong et al., 2020; Maneveldt et al., 2020;

950 Puckree-Padua et al., 2020). Herein, we have applied that methodology to the numerous

951 species and infraspecific names of Arctic and Subarctic non-geniculate corallines published

952 by Kjellman (1883; 1889) and later by Foslie (1891; 1895; 1896; 1899; 1900; 1905a, b;

953 1908) and others.

With the exception of the *Lithothamnion tophiforme* (Unger, 1858), first published as *Millepora polymorpha* var. *tophiformis* Esper (1789), the oldest names applicable to

Arctic and Subarctic non-geniculate corallines are those of Kjellman (1877; 1883; 1889).

957 By sequencing type material, Adey et al. (2015b) confirmed the application of two of

958 Kjellman's names, L. compactum Kjellman (1883) to Clathromorphum, and L. loculosum

959 Kjellman (1889) to Neopolyporolithon, and showed that a third, Lithothamnion foecundum

960 Kjellman (1883), currently placed in Leptophytum W.H.Adey, does not belong in that

961 genus, but its generic position remains unresolved. Lectotype material of the oldest name,

962 Lithophyllum arcticum Kjellman (1877), collected at Uddebay, Novaya Zemlya, Russia,

963 was shown to be an earlier available name for *Neopolyporolithon loculosum* (Kjellman)

	964	W.H.Adey, P.W.Gabrielson.	G.P.Johnson & Hernández-Kantún	, namely N. arc	ticum
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965 (Kjellman) P.W.Gabrielson, S.C.Lindstrom & Hughey (Gabrielson et al., 2019).

966 Lithothamnion flavescens Kjellman (1883) was trasferred to Leptophytum by

- 967 Athananasiadis (2016), but this was based on morpho-anatomy, which has been shown to
- 968 be problematic in correctly placing non-geniculate species in a genus (Hind *et al.*, 2016;
- 2018; Gabrielson et al., 2019). Of the remaining six Kjellman names, five are treated

970 herein, L. alcicorne, L. durum, L. glaciale, L. intermedium and L. soriferum. Attempts to

amplify L. ungeri Kjellman (1883) were unsuccessful. Thirty-five type specimens of

972 Lithothamnion species and infraspecific taxa described by Foslie were sequenced. Below

973 we discuss the systematics of the Arctic and Subarctic *Lithothamnion* species that we

974 recognize, including *L. lemoineae*, and one of the species of *Clathromorphum*, *C.* 

975 circumscriptum, for which we found an additional synonym. Distributions, habits and

- habitat data for each species are updated, including range maps for the *Lithothamnion*
- 977 species.

978 *Clathromorphum circumscriptum.* DNA sequencing confirmed two earlier proposed
979 heterotypic synonyms for this species, *L. durum* and *L. evanescens*, the first proposed by
980 Foslie (1900) and the second by Lebednik (1977). Added to these is *L. coalescens*, which
981 had been considered a synonym of *C. compactum* by Foslie (1905a) and recently by

982 Athanasiadis (2016).

The habit and habitat of *C. circumscriptum* are provided by Adey *et al.* (2015b), and the sequenced specimens confirm this information. The species is reported to be circum Arctic ranging south to the Subarctic in both the Atlantic and Pacific Oceans based on morpho-anatomy and its distinct habit (Adey, 1965; Adey *et al.*, 2013; 2015a, b). In the

987 NW Atlantic Subarctic, C. circumscriptum is generally more abundant at depths of less 988 than 10 m on moderately exposed rocky shores (Steneck, 1978; Adey & Hayek, 2011). 989 DNA sequences confirm its presence in Greenland, but material from the Russian Arctic 990 and NW Pacific Ocean has not been sequenced. DNA sequences from the western Gulf of 991 Alaska are from specimens collected in mid (GenBank MT732997) and high intertidal 992 pools (MT733001). Other mid pool (MT732990) and low intertidal collections 993 (MT732992, MT732993, MT732996) from this area represent an undescribed species of 994 Clathromorphum. Records based on morpho-anatomy from SE Alaska may also represent 995 an undescribed species. 996 Lithothamnion glaciale. There is no doubt that the lectotype specimen designated 997 herein is the one illustrated by Kjellman (1883, pls. 2, 3), despite having listed multiple 998 syntype localities. The partial *rbc*L sequence from the lecotype differs by 1 bp from all 999 other specimens that thus far have been sequenced, including one from Spitzbergen. DNA 1000 sequencing also shows that eighteen specific and infraspecific Lithothamnion taxa later 1001 named by Foslie are heterotypic synonyms of L. glaciale. Lithothamnion intermedium, 1002 described in the same publication as L. glaciale (Kjellman, 1883), was listed most recently 1003 by Athanasiadis (2016) as *Incertae Sedis*. Kjellman provided three syntype localities for L. 1004 intermedium, but he himself only collected the specimen at Karlsøy (Carlsö); specimens 1005 from the other two localities (Tromsø and Vadsø) were collected by Foslie. In UPS only 1006 two specimens could be located, one from Karlsøy, collected by Kjellman and with a label 1007 in his handwriting, and the other collected by Foslie from Mestervik, Tromsø. The 1008 Kjellman specimen from Karlsøy we have designated as the lectotype; its sequence differs 1009 by 1 bp from the sequence of the L. glaciale lecotype. By DNA sequence the specimen

collected by Foslie is *L. soriferum. Lithothamnion glaciale* and *L. intermedium* were
published at the same time and are the same species so either name can be used for this
species. We selected *L. glaciale* because of its long-standing use by the coralline research
community and because the lectotype material is homotypic, whereas *L. intermedium*mostly has been ignored.

1015 The habit and habitat of L. glaciale were described by Adey (1966a) and Adey et al. 1016 (2005) based primarily on NW Atlantic material identified by morpho-anatomy. 1017 Specimens ranged from epilithic crusts to free living rhodoliths. More recently, encrusting 1018 epilithic forms of L. glaciale have been reported to be very common from the low intertidal 1019 to the photic limit (Adey & Hayek, 2011). Most collections, however, are from the low 1020 intertidal to a depth of  $\sim 15$  m, which is more a reflection of collection limits than the 1021 species' true vertical distribution. Numerous studies in the past two decades have 1022 documented the presence of rhodoliths throughout the North Atlantic and Arctic at depths of ~3 to 50 m, and while most studies have assumed L. glaciale as the forming species, 1023 1024 little to no corroborative DNA sequencing work has been carried out as part of these studies 1025 (Halfar et al., 2000; Blake and Maggs, 2003; Kamenos & Law, 2010; Gagnon et al., 2012; 1026 Teichert et al., 2012; 2014; Adey et al., 2015a; Millar & Gagnon, 2018; Schoenrock et al., 1027 2018b; Bélanger & Gagnon, 2020; Teed et al., 2020). Although this assumption is 1028 legitimate given the ubiquity of L. glaciale in both oceans, morphological deviations from 1029 the norm in a few L. glaciale rhodoliths from Newfoundland and Labrador suggests that 1030 rhodoliths may also include other species of corallines such as L. tophiforme and C.

1031 compactum (D. Bélanger & P. Gagnon, unpublished data). In Norway, Anglés d'Auriac et

al. (2019) reported plurispecific maerl mainly composed of *L. glaciale* and associated
species of *Phymatolithon* and *Lithophylllum*.

1034 We confirm by DNA sequencing the presence of L. glaciale throughout the North 1035 Atlantic (Fig. 2). We have not confirmed many of the Arctic Ocean reports. The 1036 observation of uniporate conceptacles (mostly carposporangial) and multiporate 1037 tetra/bisporangial conceptacles in our collections from Norway, confirmed the common 1038 occurrence of gametophytes and tetra/bisporophytes for this species, as is suggested in the 1039 literature (e.g., Chamberlain & Irvine, 1994). 1040 In the North Pacific this species was first reported by Saunders (1901: 442) based 1041 on specimens collected in Prince William Sound and further west at Kukak Bay and the 1042 Shumagin Islands and identified by Kjellman. We have not confirmed by DNA sequencing 1043 the identity of these specimens. Recently, however, Bringloe & Saunders (2019) reported 1044 this species from Nome, Alaska, USA (Bering Strait), and we have sequenced specimens 1045 from Malcolm Island, central British Columbia, Canada south to Monterey County, 1046 California, USA (Fig. 2). In the NE Pacific, only encrusting epilithic specimens were 1047 found, although they may completely cover pebbles so that they appear to be rhodoliths. 1048 We cannot confirm reports based on morpho-anatomy from Japan and Arctic Russia, but 1049 likely L. glaciale is present in those areas as well. Reports of this species from any tropical 1050 and warm temperate regions as found on AlgaeBase (Guiry & Guiry, 2020) are highly 1051 improbable. Reports from the Subantarctic (Heydrich, 1900; Lemoine, 1913) need to be 1052 confirmed.

1053 *Lithothamnion lemoineae*. Melbourne *et al.* (2017) provided a *psbA* sequence from 1054 the holotype of *L. lemoineae* from the NW Atlantic (Maine, USA) to unequivocally link

DNA sequences from field-collected material to the name. Based on DNA sequenced specimens, we expand the habit of *L. lemoineae* from encrusting epilithic to epiphytic, to epizoic on shells, and as rhodoliths. The species can also occur in the intertidal, as well as in the shallow subtidal to 12 m depth. In the NW Atlantic Subarctic, *L. lemoineae* often dominates rock on exposed shores (Adey & Hayek, 2011), particularly at 10-15 m depth (R. Steneck, *pers. obs.*).

1061 Melbourne et al. (2017) found that specimens from England, thought to be L. 1062 lemoineae based on morpho-anatomy (Chamberlain & Irvine, 1994), were not that species 1063 based on DNA sequencing. However, based on DNA sequenced material, we corroborated 1064 a recent record of L. lemoineae from the same region in the NE Atlantic (Svalbard, 1065 Norway; Hofmann & Heesch, 2018), and likewise Bringloe & Saunders (2019) have the 1066 first confirmed record from the NE Pacific at Nome, Alaska, USA. Here, we confirm its 1067 occurrence on Kodiak Island, Gulf of Alaska (GenBank MT733005), where it occurred on 1068 low intertidal bedrock. The report of this species from the NW Pacific by Lee (2008) needs 1069 to be verified by DNA sequencing. We also confirm the species distribution in the NW 1070 Atlantic from Labrador, Canada to Maine, USA (Fig. 3). 1071 Lithothamnion soriferum. We located in UPS three specimens of L. soriferum in an 1072 envelope with a label in Kjellman's handwriting and from one of the cited localities, Carlsö 1073 (now Karlsøy). These appear to have been illustrated and published by Kjellman (1883, pl. 1074 1, figs. 1-4) with some artistic license. This is similar to what we found for L. durum

- 1075 (Kjellman, 1889) and L. glaciale (Kjellman, 1883), and in all of these examples the
- 1076 specimens are convincing matches to the illustrations. DNA sequences from type
- 1077 specimens of five species and infraspecific taxa later named by Foslie are all exact matches

1078	to the lectotype specimens of L. soriferum. The recently described L. erinaceum
1079	(Melbourne et al., 2017) also is a heterotypic synonym of L. soriferum. Since being
1080	published, L. soriferum has had a checkered history, sometimes being recognized as a
1081	distinct species (Foslie, 1905; Zinova, 1955) but mostly it has been considered a synonym
1082	of L. tophiforme (Foslie, 1895; Lund, 1959; Jaasund, 1965; Lee, 1969; Adey, 1970a;
1083	Vinogradova, 2010). Interestingly, Adey et al. (2005), in a detailed examination of L.
1084	tophiforme, did not mention L. soriferum, and, most recently, Athanasiadis (2016) treated
1085	the species as Incertae Sedis.
1086	Based on DNA sequences, L. soriferum is a distinct species and occurs as an
1087	epilithic, epiphytic or epizoic (specimen on a worm tube) crust or as a free-living rhodolith.
1088	Specimens are primarily subtidal to 27 m depth, but we have one occurrence in the low
1089	intertidal as an epilithic crust. In contrast to L. glaciale, gametangial plants have not been
1090	observed in any of the collections. Only multiporate tetra/bisporangial conceptacles were
1091	observed, as it is indicated in the type collections of two heterotypic synonym (L. breviaxe
1092	and L. granii, Woelkerling et al., 2005) and in the literature (Melbourne et al., 2017).
1093	The species is widespread in the central and eastern North Atlantic (Greenland,
1094	Iceland, UK, Norway), but there is no evidence of its occurrence in the NW Atlantic; in the
1095	NE Pacific there are sporadic records from the Aleutian Islands (Robinson et al., 2017) and
1096	Prince William Sound (Konar et al., 2006), Alaska, USA south to Gwaii Haanas, British
1097	Columbia, Canada (Fig. 4). For having such a widespread distribution, this species appears
1098	uncommon compared to L. glaciale. In the first report of L. soriferum from the NE Pacific
1099	(Konar et al., 2006), this species was misidentified as Phymatolithon calcareum based on
1100	morpho-anatomy. This appears to be so because the cell types in the cross-section image

1101 (Konar *et al.*, 2006, fig. 3B) were misidentified. The figure clearly shows flared epithallial
1102 cells that are not characteristic of any *Phymatolithon* species, but that are found in
1103 *Lithothamnion* species. All reports of *P. calcareum* outside boreal NE Atlantic and the
1104 Mediterranean Sea waters based on morpho-anatomy are doubtful and need to be confirmed
1105 by DNA sequences.

1106 Lithothamnion tophiforme. The sequence of the neotype specimen, TRH C13-3203, 1107 designated by Adey et al. (2005), confirms the identity of the specimen used by Adey et al. 1108 (2015b) to represent this species. Adey (1970a) reported this species to be both encrusting, 1109 especially on shells or shell fragments, and also as free-living rhodoliths. All of the 1110 sequenced specimens to date have been rhodoliths. Adey et al. (2005) considered L. 1111 tophiforme an Arctic species, but also stated that its abundance in the high Arctic is 1112 unknown, a situation that remains unchanged. They also noted that it is found only in 1113 colder waters below 10 m depth and at temperatures below 10° C, and the sequenced 1114 specimens confirm this pattern as all were collected below 15 m depth. All of the recently 1115 collected sequenced specimens are from the NW Atlantic, from Newfoundland, Canada 1116 northward (Fig. 5). The neotype specimen is the only confirmed specimen from Greenland, 1117 and only two historical specimens from the 19th century are from Norway, where the 1118 species was not found recently in an extensive collection effort presented herein. 1119 In the NW Atlantic Subarctic, L. tophiforme rhodoliths commonly co-occur with L. 1120 glaciale rhodoliths, while dominating at depths >25 m (Adey et al., 2015a). In rhodolith 1121 beds where both species occur, thalli of L. tophiforme and L. glaciale sometimes merge, 1122 forming plurispecific rhodoliths with a characteristic color mosaic where L. tophiforme 1123 generally presents a more brownish-orangy color than L. glaciale (D. Bélanger & P.

Gagnon, unpublished data). Our sequenced specimens of *L. tophiforme* presented a variety
of shapes and sizes, from a few centimetre-long twig-like thalli, to large (>10 cm across)
branching spheroidal rhodoliths. A few specimens presented a distinct growth form with
fanned-shaped branches. The phenotypic plasticity of *L. tophiforme* highlights the
importance of DNA sequencing for identification.

1129

1130 Identifying non-geniculate coralline algae to species. These Arctic and Subarctic 1131 Lithothamnion species exemplify the difficulties of identifying non-geniculate coralline 1132 species using morpho-anatomy. DNA sequencing has revealed that by the first decade of 1133 the 20th Century, 30 specific and infraspecific names had been given to three species: L. 1134 glaciale, L. soriferum and L. tophiforme. Recently, based on morpho-anatomy, the vast 1135 majority of these were listed as Incertae Sedis by Athanasiadis (2016) including, for 1136 example L. alcicorne, L. breviaxe, and L. soriferum. In the over 100 years since these had 1137 been named and examined by numerous coralline morpho-anatomists, their identity could 1138 not be determined with any certainty. And there remain another 21 species or infraspecific 1139 taxa of Lithothamnion named by Foslie, 20 from Norway and one from Scotland from 1140 which we were unable to amplify DNA using PCR. It is highly unlikely, however, that any 1141 of these names would apply, due to the extensive sequencing of Norwegian specimens 1142 reported herein and the rule of priority. With the exception of *L. lemoineae*, which has 1143 never been recorded from Norway, the applied Lithothamnion names predate any of those 1144 published by Foslie.

Further complicating the naming of specimens without DNA sequencing is that all of these species can occur either as encrusting corallines attached to a substrate or as

1147	unattached rhodoliths -encrusting a core or not - sometimes occurring singly, but also in
1148	beds. Lithothamnion lemoineae previously had only been reported to occur as an epilithic
1149	crust, whereas L. soriferum (also as L. erinaceum) had not been known as an epilithic crust.
1150	In some regions, species can have restricted morphologies, for example in Norway L.
1151	glaciale is found as an epilithic crust, or as free-living maerl, whereas in the NE Pacific it
1152	has so far only been reported as an epilithic crust. And three of the four species, L. glaciale,
1153	L. lemoineae and L. soriferum, can occur from the intertidal to, at minimum, 12 m depth.
1154	The finding by DNA sequencing of four Arctic and Subarctic Lithothamnion
1155	species that had been named as multiple specific and infraspecific taxa using morpho-
1156	anatomy is clearly opposite the cryptic diversity commonly recorded in temperate
1157	corallines (e.g., Pardo et al., 2014; 2017; Hernández-Kantún et al., 2015a, b; Peña et al.,
1158	2015a, b; Richards et al., 2018; Pezzolesi et al., 2019). However, this plethora of specific
1159	and infraspecific names in the Arctic and Subarctic regions was primarily the work of
1160	Foslie, who, as illustrated in the taxonomic results, changed his mind numerous times about
1161	which taxa should be recognized and at what rank. In tropical regions, DNA sequencing of
1162	non-geniculate corallines has shown that some species are widely distributed, whereas most
1163	have local distributions (Sissini et al., 2014; Peña et al., 2014b; Hernández-Kantún et al.
1164	2016; Gabrielson et al., 2018; Maneveldt et al., 2019).
1165	In the Arctic and Subarctic additional species of Lithothamnion and
1166	Clathromorphum need to be recognized based on the DNA sequencing reported herein
1167	(Supplementary table S2). It is also likely that the Arctic and Subarctic Lithothamnion
1168	species will need to be transferred to a new genus, as the generitype of Lithothamnion, L.
1169	muelleri Lenormand ex Rosanoff, belongs in a different clade (Yeong et al., 2020).

1170 It is critical that in this time of rapid ocean warming and acidification, 1171 particularly in polar regions, that we have a firm understanding of the taxa currently present 1172 in order to document future changes in their habitats and distributions. Importantly, the 1173 biogeography of coralline algae appears especially sensitive to ocean thermogeography 1174 (Adey & Steneck, 2001; Adey & Hayek, 2011). The relevance of non-geniculate coralline 1175 algae in these regions as ecosystem engineers cannot be overstated, whether occuring as 1176 encrusting species attached primarily to rock substratum (Freiwald & Henrich, 1994; Adey 1177 et al., 2005; Adey et al., 2015a) or as free-living maerl (Pardo et al., 2014; Teichert, 2014; 1178 Teed et al., 2020). As polar seas warm and become increasingly acidified, these coralline 1179 algal species will either be forced to live at lower depths, where they will be limited by the 1180 availability of photosynthetically active radiation through the water column, or they will 1181 become extinct. DNA barcoding of organisms in these habitats, coupled with DNA 1182 sequencing of type and historical specimens, provides the foundation to document these 1183 imperiled species.

1184

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- 1199 Supplementary information
- 1200 Supplementary Note 1. Type collections & historical specimens DNA-extracted but
- 1201 unsuccessfully sequenced for *psbA* and *rbcL*.
- Supplementary Table S1. Type collections and historical specimens sequenced in thestudy.
- 1204 Supplementary Table S2. Collection and vouchering details for the recent specimens
- 1205 sequenced in the study. For each species, specimens are sorted by collection areas: NE
- 1206 Atlantic, NW Atlantic and NE Pacific.
- 1207 Supplementary Table S3. List of GenBank accession numbers used in this study with
- 1208 details of locality information and references.

1209 Supplementary Figure S1. Maximum Likelihood (ML) tree of COI-5P included in the

- 1210 present sudy. Bootstrap ML values >50% shown for each node. Scale bar: 0.05
- 1211 substitutions per site.
- 1212 Supplementary Figure S2. Phylogenetic tree inferred from RAxML and Bayesian
- 1213 inference of *psbA* sequences included in the present study. Bootstrap ML values >50% and
- 1214 posterior probabilities >0.50 from Bayesian inference shown for each node. Scale bar: 0.02
- 1215 substitutions per site.

1216	<b>Supplementary Figure S3</b> . Morpho-anatomy of <i>Lithothamnion glaciale</i> , UPS A-000202.
1217	Fig. A. Thallus habit and herbarium labels including label handwritten by Kjellman. Fig. B.
1218	Vertical fracture of protuberance showing radial construction and location of overgrown
1219	and buried conceptacle (black arrow). Fig. C. Vertical fracture of protuberance with
1220	conceptacle (black arrow) overgrown by a secondary hypothallus (arrowheads). Fig. D.
1221	Magnified view of secondary hypothallus (bracket) over conceptacle roof. Fig. E.
1222	Perithallus with cell fusions (white arrows). Fig. F. Perithallus (lower bracket), intercalary
1223	meristem (middle bracket, *), and a single-layered epithallus (upper bracket) of flared cells
1224	(e), one with cell roof intact (white arrow). Scale bars: Figs. B-C = 100 $\mu$ m; Figs. D-F = 10
1225	μm.
1226	Supplementary Figure S4. Morpho-anatomy of Lithothamnion soriferum, UPS A-648809.
1227	Fig. A. Thallus habit of specimens and herbarium label handwritten by Kjellman. Figs. B-
1227 1228	Fig. A. Thallus habit of specimens and herbarium label handwritten by Kjellman. Figs. B- D. Vertical fractures of protuberance showing radial construction. Fig. E. Magnified view
1228	D. Vertical fractures of protuberance showing radial construction. Fig. E. Magnified view
1228 1229	D. Vertical fractures of protuberance showing radial construction. Fig. E. Magnified view of secondary hypothallus (arrowhead, bracket) over the thallus surface of the older growth
1228 1229 1230	D. Vertical fractures of protuberance showing radial construction. Fig. E. Magnified view of secondary hypothallus (arrowhead, bracket) over the thallus surface of the older growth layer. Fig. F. Perithallus (lower bracket) with cell fusions (white arrows), intercalary
1228 1229 1230 1231	D. Vertical fractures of protuberance showing radial construction. Fig. E. Magnified view of secondary hypothallus (arrowhead, bracket) over the thallus surface of the older growth layer. Fig. F. Perithallus (lower bracket) with cell fusions (white arrows), intercalary meristem (middle bracket, *), and epithallus (upper bracket). Figs. G-H. Magnified view of
1228 1229 1230 1231 1232	D. Vertical fractures of protuberance showing radial construction. Fig. E. Magnified view of secondary hypothallus (arrowhead, bracket) over the thallus surface of the older growth layer. Fig. F. Perithallus (lower bracket) with cell fusions (white arrows), intercalary meristem (middle bracket, *), and epithallus (upper bracket). Figs. G-H. Magnified view of meristematic cells (*) and a single-layered epithallus of flared cells (e), one with cell roof
1228 1229 1230 1231 1232 1233	D. Vertical fractures of protuberance showing radial construction. Fig. E. Magnified view of secondary hypothallus (arrowhead, bracket) over the thallus surface of the older growth layer. Fig. F. Perithallus (lower bracket) with cell fusions (white arrows), intercalary meristem (middle bracket, *), and epithallus (upper bracket). Figs. G-H. Magnified view of meristematic cells (*) and a single-layered epithallus of flared cells (e), one with cell roof intact (white arrow, Fig. H). Scale bars: Fig. A = 2 cm. Figs. B-D = 100 $\mu$ m. Figs. E-H = 10

**Disclosure statement** 

1237 No potential conflict of interest was reported by the authors.

1239	Author Contributions
1240	V. Peña & P.W. Gabrielson: original concept; V. Peña, P.W. Gabrielson, L. Le Gall & J.R.
1241	Hughey: DNA sequencing types; J.L. Richards: morpho-anatomy of type specimens; V.
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- 1659 Figure legends
- 1660 **Figure 1**. Phylogenetic tree inferred from maximum likelihood (ML) and Bayesian
- 1661 inference of *rbcL* sequences included in the present study. Bootstrap ML values >60% and
- 1662 posterior probabilities >0.60 from Bayesian inference shown for each node. Scale bar: 0.05
- 1663 substitutions per site.
- 1664 **Figure 2**: Distribution map of *Lithothamnion glaciale* obtained for collections analyzed in
- 1665 the molecular studies, type collections and historical specimens.
- 1666 Figure 3: Distribution map of *Lithothamnion lemoineae* obtained for collections analyzed
- 1667 in the molecular studies, type collections and historical specimens.
- 1668 Figure 4: Distribution map of *Lithothamnion soriferum* obtained for collections analyzed in
- 1669 the molecular studies, type collections and historical specimens.
- 1670 Figure 5: Distribution map of *Lithothamnion tophiforme* obtained for collections analyzed
- 1671 in the molecular studies, type collections and historical specimens.