

Authentication of Iceland Moss (Cetraria islandica) by UPLC-QToF-MS chemical profiling and DNA barcoding

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**Authentication of Iceland Moss by UPLC-QToF-MS chemical profiling
and DNA barcoding**

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Abstract

The lichen *Cetraria islandica* or Iceland Moss is commonly consumed as tea, food ingredients (e.g. in soup or bread) and herbal medicines. *C. islandica*, which has two chemotypes, can be difficult to distinguish from the sister species *Cetraria ericetorum*. They are collectively referred to as the *Cetraria islandica* species complex. This study aimed to use an UPLC-QToF-MS chemical profiling together with DNA barcoding to distinguish species and chemotypes of the *C. islandica* species complex. Our results show that the two chemotypes of *C. islandica* are clearly distinguishable from each other and from *C. ericetorum* by the chemometric approach. The RPB2 barcode was able to differentiate *C. islandica* from *C. ericetorum* with a barcode gap, but the widely used nrITS barcode failed. Neither of them could discriminate chemotypes of *C. islandica*. In conclusion, this integrative approach involving chemical profiling and DNA barcoding could be applied for authentication of Iceland Moss materials.

Keywords: *Cetraria islandica*, *Cetraria ericetorum*, DNA barcoding, chemical profiling, authentication

1 **1. Introduction**

2
3 Taxonomically, *Cetraria islandica* or Iceland Moss is not a moss species
4 but a lichen taxon and the classification is based on the symbiotic fungal
5 partner (Parmeliaceae, ascomycete) (Ingólfssdóttir, 2000). This lichen is
6 consumed in Iceland as tea, food ingredients (e.g. in milk soups and bread)
7 and herbal medicines (Xu et al., 2016). Considerable morphological and
8 chemical variations have been found among Icelandic populations of *C.*
9 *islandica* and two chemotypes have been reported (Kristinsson, 1969). They
10 consist of the fumarprotocetraric acid (FA)-producing and FA-deficient
11 races, where the latter chemotype has exclusively been found in Iceland
12 (Kristinsson, 1969). Traditional use seems to favor the FA-deficient
13 chemotype of *C. islandica*, which is believed to be less bitter (Kristinsson,
14 1968). Furthermore, the species boundaries between *C. islandica* and its
15 sibling species *Cetraria ericetorum* are still ambiguous: *C. ericetorum* has
16 similar morphology to certain morphotypes of *C. islandica* and it is reported
17 to be FA-deficient. Together *C. islandica* and *C. ericetorum* are collectively
18 called the *Cetraria islandica* species complex (Kristinsson, 1969; Thell,
19 Stenroos, & Myllys, 2000). Their chemical profiles, particularly of the *C.*
20 *islandica* FA-deficient chemotype and *C. ericetorum*, have not been
21 thoroughly investigated for food safety, and an accurate identification
22 method for these lichen materials is needed.

23
24 Chemical profiling or fingerprinting, in particular when using an untargeted
25 approach, can directly detect chemical hazards and contaminants in food or
26 herbal materials, with the limitation that their species sources cannot be
27 determined (de Boer, Ichim, & Newmaster, 2015). Chemometric analysis
28 using complex metabolite datasets has shown great potential in the
29 inspection of food adulteration as well as in the characterization of markers
30 for adulteration detection (Cubero-Leon, Peñalver, & Maquet, 2014).

31 Untargeted chemical profiling is especially useful in the distinction of
32 closely-related plant species, where certain genetic markers may not be
33 informative (Messina, Callahan, Walsh, Hoebee, & Green, 2014). That
34 approach has been successfully applied to the lichen *Ramalina siliquosa*
35 complex using liquid chromatography-mass spectrometry (LC-MS) (Parrot,
36 Jan, Baert, Guyot, & Tomasi, 2013).

37
38 Recently, DNA barcoding has emerged as an effective tool in the
39 identification of plant and animal materials using defined species-specific
40 DNA markers. It has found wide application in the authentication and
41 traceability of food materials (Galimberti et al., 2013). This approach has
42 been extended to the authentication of multiple ingredients samples using a
43 more advanced DNA metabarcoding approach, which involves next
44 generation sequencing (Staats et al., 2016). Practically, DNA barcoding has
45 been applied for authentication of fungi-based dietary products (Raja, Baker,
46 Little, & Oberlies, 2017). Furthermore, identification of lichenized fungi
47 using DNA barcoding has been successfully performed using the fungal
48 nuclear ribosomal internal transcribed spacer region (nrITS) (Kelly et al.,
49 2011), which has been proposed as the universal DNA barcode for fungi
50 (Schoch et al., 2012).

51
52 Two DNA barcodes, nrITS and RPB2 (the second largest subunit of
53 ribosomal polymerase II) were selected for this study on the *C. islandica*
54 lichen materials. Although the widely used nrITS region is known to
55 provide a sufficient amount of variation to distinguish between most fungal
56 species and is represented by many reference sequences in public databases,
57 some drawbacks for DNA barcoding and especially DNA metabarcoding
58 have been reported (Větrovský, Kolařík, Žifčáková, Zelenka, & Baldrian,
59 2016). While nrITS can identify species, its multi-copy nature of the ITS
60 region may render problems with relative quantification of species in mixed

61 samples. In other cases, intra-individual polymorphism including multiple
62 functional genes, putative pseudo genes or recombinants hamper
63 identifications (Mark, Cornejo, Keller, & Flück, 2016). The single-copy
64 RPB2 gene has been proposed as an alternative to the nrITS region, which
65 could overcome some of these challenges (Větrovský et al., 2016). The
66 performance of RPB2 will be compared to that of nrITS in our study.

67

68 The overall aim of this study was to explore the usefulness of an
69 authentication approach for *Cetraria islandica* species complex using
70 UPLC-QToF-MS chemical profiling and DNA barcoding. Specific
71 objectives were: Firstly, to distinguish chemotypes of the species complex
72 by comparing their UPLC-QToF-MS chemical profiles using chemometric
73 data analysis, and secondly to compare the discriminatory power of RPB2
74 and nrITS barcodes for the taxa of the *C. islandica* species complex.

75

76 **2. Materials and Methods**

77

78 2.1 Sampling and chemotype identification

79

80 *Cetraria islandica* (L.) Ach. (English: Iceland Moss; Icelandic: fjallagrös)

81 and *Cetraria ericetorum* Opiz specimens were collected in Iceland.

82 Authentic specimens of *C. islandica* are provide by lichenologists from the

83 Icelandic Institute of Natural History, Akureyri, Iceland (AMNH). Collected

84 voucher specimens are deposited at the AMNH herbarium. Intraspecific

85 morphological variation of Iceland Moss (*C. islandica*) as well as the

86 interspecific similarity between the two species (*C. islandica* and *C.*

87 *ericetorum*) can be seen in Figure 1. Additional *C. ericetorum* type

88 specimens were kindly provided by Dr. Stefan Ekman, Museum of

89 Evolution, Uppsala University, Uppsala, Sweden. The chemotypes of *C.*

90 *islandica* and *C. ericetorum* were tested using a conventional spot testing

91 method (Kristinsson, 1969). Briefly, a small fragment was cut from a thallus
92 with a blade, and drops of *p*-phenyldiamine (PD) solution (ca. 2% in
93 ethanol) were added to lichen medulla on a white paper with a glass
94 capillary tube. Then the medullary color reactions were visualized under a
95 stereoscope. Specimens showing a red medullary color after spot testing
96 were assigned as PD+, while the ones without color change as PD-. PD spot
97 testing was carried out on fragments from three different parts of the thallus,
98 to make sure the chemotype. Fragments were discarded immediately after
99 testing. Voucher information and gene accession numbers are provided in
100 Online Resource (see Table S1). In total, 30 specimens of PD+ *C. islandica*,
101 15 specimens of PD- *C. islandica* and 18 specimens of uniformly PD- *C.*
102 *ericetorum* were identified.

103

104 2.2 Chemical profiling

105

106 2.2.1 LC-MS analysis

107

108 Air-dried lichen thallus (ca. 20 mg) was weighed and ground into powders
109 under liquid nitrogen. Powdered lichen materials were macerated with
110 acetone under shaking in ambient temperature for 2 h. The extraction was
111 repeated twice. Extracts were combined and evaporated with nitrogen gas
112 flow. Dried residues were then solubilized in HPLC-grade acetonitrile
113 (ACN), diluted into 0.1 mg/mL and filtered (pore size 0.45 µm; GE
114 healthcare, UK) before analyses by Waters ACQUITY UPLC™ (Waters
115 Corporation, Milford, MA, USA) coupled to Waters Q-ToF SYNAPT G1
116 mass spectrometer (Waters MS Technologies, Manchester, UK).

117

118 The UPLC system was equipped with a binary solvent delivery system and
119 autosampler. Chromatographic separation of lichen compounds was
120 conducted on an ACQUITY UPLC BEH C18 column (2.1 mm x 100 mm,

121 1.7 μm ; Waters corp., Milford, MA, USA). The column oven was kept at
122 40°C and the autosampler was maintained at 6.0°C. The mobile phase
123 consisted of solvent A: H₂O with 0.1% formic acid in water and solvent B:
124 0.1% formic acid in ACN. Gradient elution was used at a flow rate of 0.40
125 mL/min as follows: 30% B, 0-1 min; linear gradient from 30% B/70% A to
126 70% B/30% A, 1-3 min; linear gradient from 70% B/30% A to 100% B, 3-9
127 min; holding at 100% B, 9-13 min; linear gradient from 100% B to 30%
128 B/70% A, 13-14 min; holding at 30% B/70% A, 14-15 min. Pooled samples
129 were used as quality control. The injection volume was 5 μL .

130

131 The Synapt G1 QToF-MS mass spectrometer was operated in negative
132 electrospray ionization mode (capillary voltage 3.0 kV, source temperature
133 120°C, desolvation temperature 400°C, cone gas flow 50 L/h, desolvation
134 nitrogen gas flow 800 L/h). Ions with mass range 50 to 1600 m/z (mass to
135 charge ratio) were scanned. All samples were analyzed in triplicates. Details
136 of UPLC-QToF-MS analysis are as previously described (Xu et al., 2017).
137 The UPLC-QToF-MS system and data acquisition were controlled by the
138 MassLynx v4.1 software (Waters Corp., Milford. USA).

139

140 2.2.2 Chemometric data analysis

141

142 MS spectra were aligned and normalized using MakerLynx v4.1. Collection
143 parameters were set as 50 counts, mass window 0.05 Da and retention time
144 window 0.2 min. Replicate percentage value was set at 50%. Normalized
145 data were introduced into SIMCA v14.1 software (Sartorius Stedim Data
146 Analytics, Umeå, Sweden) for principal component analysis (PCA). PCA
147 could provide a holistic overview of the grouping of lichen specimens,
148 which was based on chemical data from organic extracts. Compounds were
149 identified by comparing their MS/MS spectra and fragmentation patterns

150 with those from isolated pure compounds, previously published data and
151 public databases (Metlin and ChemSpider).

152

153 2.3 Molecular analysis

154

155 2.3.1 DNA extraction, PCR and sequencing

156

157 Air-dried lichen residues after acetone maceration were used for total DNA
158 extraction following the CTAB protocol (Cubero, Crespo, Fatehi, & Bridge,
159 1999). Lichen DNA extracts were stored in TE buffer (pH 8.0) at -20°C
160 until use. Polymerase chain reactions (PCRs) were performed to amplify the
161 fungal nuclear ribosomal internal transcribed spacer (nr ITS) and the second
162 largest subunit of RNA polymerase II (RPB2). Each reaction (25 µL)
163 contained 1×standard Taq reaction buffer, 200 µM dNTPs, 0.2 µM forward
164 and reverse primer, 1.25 units of Taq DNA polymerase (New England
165 Biolabs), 1 µL DNA template, and PCR-grade water. The fungi-specific
166 primers used for the amplification of nr ITS region were: ITS1F (5'-
167 CTTGGTCATTTAGAGGAAGTAA-3') (Gardes & Bruns, 1993) and ITS4
168 (5'-TCC CCGCTTATTGATATGC-3') (White, Bruns, Lee, & Taylor,
169 1990), while the primers for the RPB2 regions were: RPB2-6F (5'-
170 TGGGGKWTGGTYTGYCCTGC-3') (Liu, Whelen, & Hall, 1999) and
171 fRPB2-7cR (5'-CCCATRGCTTGYTTRCCCAT-3') (Liu et al., 1999). PCR
172 amplification was carried out in a Perkin-Elmer Gene Amp PCR system
173 9700 thermal cycler. The PCR cycling conditions for nrITS were: initial
174 denaturation at 94°C for 3 min, 34 cycles of 94°C for 40 s, 54°C for 40 s,
175 68°C for 1 min, then final extension at 68°C for 5 min before cooling down
176 to 4°C. A touchdown PCR program was used for RPB2 region: 94°C for 4
177 min, followed by 6 cycles of 94°C for 1min, 55-50°C (decrease 1°C per
178 cycle) for 1min and 68°C for 1min, then 32 cycles of 94°C for 1min, 50°C
179 for 1min and 68°C for 1min, and final extension at 68°C for 7min, before

180 cooling down at 4°C. Amplicons were visualized in 1.3% agarose gel (gel
181 picture refers to Online Resource Figure S1), purified using ExoSAP
182 (Fermentas) and sent for Sanger sequencing by Marogen Inc. using the same
183 set of primers as used in PCRs.

184

185 2.3.2 DNA barcoding analysis

186

187 The discriminatory power of the nrITS and RPB2 barcodes was assessed
188 according to the monophyly criterion and the DNA barcoding gap concept
189 (Meyer & Paulay, 2005). The sequences representing each barcode were
190 aligned using MAFFT v7.215 (Kato & Standley, 2013) with default
191 parameters and trimmed if necessary. Phylogenetic trees using the Genbank
192 sequences of *C. sepincola* (accession number KC990137.1 for nrITS) as
193 outgroups were built with RAxML v. 8.0.26 (Stamatakis, 2014) with 100
194 rapid bootstrap replicates under the GTRGAMMA model. The DNA
195 barcode gap analysis was conducted on *C. islandica* and its sister species *C.*
196 *ericetorum* using the R package SPIDER (Brown et al., 2012), using the
197 best-fitting substitution models to measure pair-wise distances. Best-fitting
198 substitution models for each region (TIM2+G for nrITS; TIM2+I for RPB2)
199 were chosen by using the Aikake Information Criterion (AIC) in
200 jModelTest 2 (Darriba, Taboada, Doallo, & Posada, 2012). Because the
201 TIM2+G and TIM2+I models were not available in the R package APE
202 (Paradis, Claude, & Strimmer, 2004), which was used to calculate pairwise
203 distances, the next best-fitting model for both alignments, TrN+G, was used.

204

205 The number of false positive and false negative identifications along a DNA
206 divergence threshold were calculated and plotted using the R package
207 SPIDER (Brown et al., 2012). Additionally, the minimum interspecific and
208 maximum intraspecific divergence for each sequence was calculated and
209 plotted.

210

211 3. Results and Discussion

212

213 3.1 Chemical profiling and chemometric data analysis

214

215 Chemical profiling of *C. islandica* organic extracts has been carried out
216 before using high performance liquid chromatography-ultraviolet detection
217 (HPLC-UV) (Fernández-Moriano, Divakar, Crespo, & Gómez-Serranillos,
218 2015; Gudjónsdóttir & Ingólfssdóttir, 1997). Those previously used methods
219 lack separation efficiency and sensitivity, and thus may underestimate the
220 chemical diversity of *C. islandica*, e.g. (+)-roccellaric acid **6** was found in *C.*
221 *islandica* using a fluorous tag-catch and release approach (Horhant, Lamer,
222 Boustie, Uriac, & Gouault, 2007), but was previously overlooked due to
223 poor chromatographic separation. Additionally, HPLC-UV (Fernández-
224 Moriano et al., 2015) analysis using high UV wavelengths may overlook the
225 content of those aliphatic lichen acids (i.e. compounds **5-8**), which are poor
226 UV absorbants. The current UPLC method achieved the separation of
227 compound **6** from its analogues **7** and **8**. A list of detected compounds is
228 provided in Table S2, and the structures of major lichen secondary
229 metabolites from the *C. islandica* species complex are illustrated in Figure 2.

230

231 Compounds **1-8** were identified in our study by comparing their molecular
232 masses, fragmentation pathway and chromatographic properties with
233 reference data as well as authentic standards isolated in previous studies
234 (Bessadóttir et al., 2014; Gudjónsdóttir & Ingólfssdóttir, 1997). MS² spectra
235 of each compound and their fragmentation patterns are provided in Online
236 Resource (see Figures S2-S4). MS chromatograms (Figure 3) show that (+)-
237 protolichesterinic acid **7** and its derivatives (i.e. **6** and **8**) are the dominant
238 compounds in organic extracts detected in negative ion mode, followed by
239 minor components, such as protocetraric acid **1** and fumarprotocetraric acid

240 3. The stereochemical diversity of (+)-protolichesterinic acid **7** seems to be
241 largely underestimated before, since two compounds (i.e. **7A** and **7B**) were
242 detected with the same molecular formula, molecular ions and
243 fragmentation patterns (Figure S4) with compound **7** (Table S2 and Figure
244 3). In *C. ericetorum*, an additional unknown compound **6A** in the peak
245 eluting out at 5.77 min (Figure 3c) was detected having the same mass to
246 charge ratio as well as fragmentation pattern as (+)-roccellaric acid **6**, which
247 suggests that **6A** could be a stereoisomer of compound **6** (Figure S4). Up to
248 now, only one stereochemical form of roccellaric acid has been reported in
249 nature, namely (+)-roccellaric acid **6** in *C. islandica* (Horhant et al., 2007).
250 Three additional stereochemical forms have been synthesized by Mulzer et
251 al. (Mulzer, Salimi, & Hartl, 1993). Minor compounds **1** and **3** eluted quite
252 early ($t_R = 2.49$ and 2.96 min, respectively) under the chromatographic
253 conditions used, reflecting that they are more water-soluble than compounds
254 **5-8**. Fumarprotocetraric acid **3** (50 mg) is reported to be moderately soluble
255 in 30 mL phosphate buffer at pH 7.4 (Syers, 1969), while the solubility of
256 compound **3** is low (1 mg/L) in 90% acetonitrile with 1% phosphoric acid
257 (Gudjónsdóttir & Ingólfssdóttir, 1997). Thus, polarity and pH of the
258 extraction solvent can be expected to have considerable influence on the
259 extraction efficiency of these lichen acids (i.e. compounds **1-4**).

260

261 As a conventional diagnostic tool, the PD spot test was used to check the
262 chemotype and the presence of compound **3** in *C. islandica* (Kristinsson,
263 1969). From LC-MS chromatograms shown in Figure 3, the red color
264 reaction by PD spot testing was found to correlate with the presence of
265 compounds **1** and **3**, while these compounds were absent in the PD- *C.*
266 *islandica* chemotype and *C. ericetorum*. The presence of the aliphatic lichen
267 acids (i.e. compounds **5-8**) did not result in a red color reaction. The co-
268 occurrence of compounds **1** and **3** in organic extracts of *C. islandica* has been
269 found in literature (Fernández-Moriano et al., 2015).

270
271 The two-component PCA score plot (Figure 4) of the UPLC-QToF-MS data
272 provides the visualization of how different chemical groups relate to each
273 other. Three chemical groups were formed based on their chemical profiles,
274 representing the PD- (CI PD-) and PD+ chemotypes (CI PD+) of *C.*
275 *islandica* and *C. ericetorum* (CE). The first component explains 43.6%
276 chemical variations, mainly interspecific differences between CE and CI.
277 The secondary component accounts for 10.4% variations, mainly
278 intraspecific differences between CI PD+ and CI PD-. PCA is a useful tool
279 in summarizing metabolite data and revealing groupings of food ingredients
280 from different biological origins (Azilawati, Hashim, Jamilah, & Amin,
281 2015; Cubero-Leon et al., 2014). From the Figure 4, the lichen *C. islandica*
282 shows high intraspecific chemical variations in Iceland with two
283 chemotypes recognized as reported before (Kristinsson, 1969), while
284 Icelandic *C. ericetorum* shows relatively less variation, even when
285 compared with non-Icelandic *C. ericetorum* specimens. This could partly be
286 explained by the limited distribution of CE, resulting in less variation. CE
287 has a restricted geographic distribution in north and east Iceland, whereas CI
288 has a rather wide distribution around Iceland (Thell & Moberg, 2011).

289

290 3.2 DNA barcoding

291

292 Sizes of PCR products were ca. ~600-900 bp for fungal nrITS and ca. ~900
293 bp for fungal RPB2 (Online Resource Figure S1). The variation of the
294 fungal nrITS PCR products was due to the presence of a group I intron
295 sequence in the longer amplicons, and absence in the short ones. In total 97
296 new sequence were obtained, including 48 for RPB2 and 49 for fungal
297 nrITS. PCRs of a few old herbarium reference specimens were not
298 successful (Online Resource Table S1).

299

300 The phylogenetic tree based on the ITS region (Figure 5a) shows that *C.*
301 *ericetorum* is paraphyletic and nested within the *C. islandica* clade, whereas
302 both *C. ericetorum* and *C. islandica* are monophyletic in the RPB2 tree
303 (Figure 5b). Therefore the phylogenetic analyses support RPB2 as a barcode
304 with power to discriminate between the two species of *Cetraria*, but shows
305 that the nrITS barcode does not discriminate the species under study. The
306 pairwise distance analysis supports this interpretation, revealing that RPB2
307 is able to distinguish the two taxa, whereas the nrITS region always yields
308 false positive or negative identifications (Figure 6). There are other cases
309 where the nrITS region fails to discriminate lichenized fungi (Kelly et al.,
310 2011; Pino-Bodas, Martín, Burgaz, & Lumbsch, 2013), though the nrITS
311 region revealed interspecific barcoding gaps among most species of genera
312 *Melanelia* and *Montanelia* (Leavitt et al., 2014; Pino-Bodas et al., 2013).
313 The failure of the nrITS region in this case might be attributed to
314 intragenomic polymorphism in the ITS region, a phenomenon described in
315 other lichen-forming fungi (Kelly et al., 2011; Mark et al., 2016). It has also
316 been reported that the nrITS region alone is not suitable to estimate the
317 phylogenetic relationships within the *C. islandica* group (Thell et al., 2000).
318
319 The limited application of RPB2 as a barcoding region has been explained
320 by difficult PCR amplification and sequencing (Schoch et al., 2012).
321 Specimens stored for over 3 years have shown problems in PCR
322 amplification (Kelly et al., 2011). We have, however, not encountered a
323 PCR amplification problem for either locus even with specimens which are
324 15 years old. Successful PCR amplification of the RPB2 region using even
325 older specimens (*Cladonia* sp.) of about 30 years has also been recorded
326 (Pino-Bodas et al., 2013). Age-dependent problems with PCR amplification
327 may be taxon-specific, as well as influenced by the DNA extraction method
328 in use. We noted that the sequence alignment of RPB2 is much simpler. In
329 contrast to the hypervariability of the nrITS region, RPB2 is also

330 recommended as an alternative marker for phylogenetic analysis (Větrovský
331 et al., 2016). We therefore reject the nrITS region and propose the RPB2
332 region as an efficient DNA barcode for testing medicinal products
333 containing Iceland Moss, at least in terms of discriminating between *C.*
334 *ericetorum* and *C. islandica*.

335

336 Although the RPB2 region is effective for discriminating between species,
337 chemotypes of *C. islandica* are not discriminated (Figure 5). In Figure 5b, *C.*
338 *islandica* specimens from Iceland show two strongly supported clades, I and
339 II. All of the *C. islandica* PD- chemotype specimens fall into clade I, but are
340 interspersed with PD+ chemotypes, while clade II contains exclusively PD+
341 *C. islandica* specimens. Some lichen chemotypes have been shown to be
342 monophyletic (Fehrer, Slavíková-Bayerová, & Orange, 2008), but they can
343 also be not (Lutsak, Fernández-Mendoza, Nadyeina, Şenkardeşler, &
344 Printzen, 2017).

345

346 Domestically, *Cetraria islandica* is sold as whole lichen-thalli food
347 ingredients or tea in Iceland. Accurate identification is generally not
348 difficult for taxonomic experts, but it may prove intractable to identify
349 powdered lichen materials, which lack morphological or sometimes
350 chemical characters. DNA barcoding as outlined here could substantially
351 facilitate identification by comparing new sequence data with reference data
352 generated from expertly identified voucher specimens.

353

354 The current study focused on the authentication of natural lichen materials
355 without downstream processing. In case of highly processed herbal
356 materials where DNA may undergo considerable degradation, an alternative
357 method could be double gene targeting PCR, which amplifies selected
358 shorter regions (e.g. 70-150 bp) (Hossain et al., 2016, 2017).

359

360 3.3 The advantage of the integrative approach for authentication of
361 Iceland Moss
362
363 Lichen material authentication generally operates at the species level and as
364 shown in this study DNA barcoding using RPB2, is an efficient method for
365 species identification in the *C. islandica* species complex. The advantage of
366 DNA barcoding is in identification of raw plant materials, sources of
367 contaminants and species composition (de Boer et al., 2015), which is
368 beyond the scope of chemical analysis. Generally, chemical profiling of
369 lichen secondary metabolites do not have the independent role in
370 identification/authentication of lichen species (Lumbsch, 1998). First, the
371 utility of metabolite data in lichen identification varies among lichen
372 taxa/populations. Our results (Table S2 and Figure 4) have demonstrated the
373 utility of chemical profiling in discriminating species (*C. islandica* and *C.*
374 *ericetorum*) and chemotypes (PD+ and PD-) in the *Cetraria islandica*
375 species complex. However, chemical profiling may have limited utility in
376 species discrimination where remarkable chemical variations (e.g. different
377 in major lichen compounds) are present, such as the lichen *Ramalina*
378 *siliquosa* (Lumbsch, 1998; Parrot et al., 2013). Such a huge variation may
379 pose a challenge in lichen identification: how much chemical variation is
380 allowed to define a species? To address this problem, it has been suggested
381 that chemical characters be correlated with other characters, preferentially
382 genetic sequence data (Lumbsch, 1998). The correlation between
383 phylogenetic relationship and chemotyping (i.e. PD+ and PD-) was
384 investigated in our study (Figure 5b).
385
386 Although chemical profiling does not have an independent role in lichen
387 species identification/Iceland Moss authentication, it is indispensable for the
388 quality control of marker or health-beneficial components. It can provide
389 both qualitative and quantitative information on phytochemical composition

390 during extraction and downstream processing. Coupled to chemometric
391 tools, chemical profiling could also be used in differentiation of intraspecific
392 chemical variants, which is superior to DNA barcoding.

393

394 In conclusion, this study highlights the integrative use of chemical profiling
395 and DNA barcoding for the authentication of Iceland Moss. The members of
396 *Cetraria islandica* species complex were easily characterized using
397 chemometric tools. Furthermore, DNA barcodes were compared and the
398 locus RPB2 proved to be superior to nrITS in distinguishing species of *C.*
399 *islandica* species complex. Our study shows how chemical profiling and
400 DNA barcoding can be used to differentiate chemical variants and species in
401 the complex, and suggests the use of this integrated approach for accurate
402 characterization of this closely related taxa as well as other plant materials
403 used for human consumption.

404

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406

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415

416 **Conflicts of interest:** none

417

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592

593 Figure captions
594

595 **Figure 1.** Morphological variation and similarity of Iceland
596 Moss (*Cetraria islandica*) chemotypes and its sibling species
597 *Cetraria ericetorum*. *C. islandica* shows considerable
598 morphological variation, from wide (**a** and **c**) to narrow thallus
599 (**b** and **d**). Two chemotypes were identified in *C. islandica*
600 specimens using *p*-phenyldiamine (PD) spot testing/staining,
601 including PD+ (red medullary color after staining; **a** and **b**) and
602 PD- (no red medullary color after staining; **c** and **d**). The lichen
603 *C. ericetorum* (**e**) is uniformly PD- and has narrow thallus.
604 Scale: 1 cm.

605

606 **Figure 2.** Chemical structures of major lichen secondary
607 metabolites detected in the *Cetraria islandica* species complex.
608 Compounds include protocetraric acid **1**, succinprotocetraric
609 acid **2**, fumarprotocetraric acid **3**, virensic acid **4**,
610 nephrosterinic acid **5**, (+)-roccellaric acid **6**, (+)-
611 protolichesterinic acid **7** and (+)-lichesterinic acid **8**. Minor
612 compounds refer to Table S2.

613

614
615 **Figure 3.** MS chromatograms of the PD+ (**a**) and PD- (**b**)
616 chemotypes of *Cetraria islandica* and PD- *C. ericetorum* (**c**)
617 and thallus color reaction by PD staining of PD+ (**d**) and PD- (**e**)
618 chemotypes of *Cetraria islandica* and PD- *C. ericetorum* (**f**).

619 Major secondary metabolites are labelled corresponding to
620 structures **1-8** in Figure 2. Compounds **7A** and **7B** were
621 tentatively identified as stereoisomers of (+)-protolichesterinic
622 acid **7**, and **6A** a stereoisomer of (+)-roccellaric acid **6**. Scale =
623 0.5 mm.

624

625 **Figure 4.** PCA plot giving an overview of metabolite data and
626 indicative grouping of species and chemotypes in the *Cetraria*
627 *islandica* species complex. Three chemical groups include PD-
628 chemotype (CI PD-), *C. islandica* PD+ chemotype (CI PD+)
629 and *C. ericetorum* (CE). Authentic herbarium specimens were
630 marked as dark green (CI PD+), dark red (CI PD-) and grey
631 (CE).

632

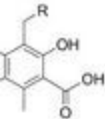
633 **Figure 5.** Maximum likelihood (ML) trees of the *Cetraria*
634 *islandica* species complex reconstructed using barcode markers.
635 (a) ML tree reconstructed using the nrITS barcode, with *C.*
636 *ericetorum* specimens marked in red; (b) ML tree using the
637 RPB2 marker, where tree well-supported clades were identified:
638 I, II and III. The PD+ chemotype is labelled with a red dot after
639 each specimen. Bootstrap values > 70 are shown above
640 branches in both trees.

641

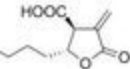
642 **Figure 6.** Barcoding gap analysis of *Cetraria islandica* species
643 complex for each marker. (a) Number of false positive and

644 false negative identifications along a threshold from 0.1% – 4%
645 DNA divergence; **(b)** Evaluation of inter- vs. intraspecific
646 divergence. The distances for each gene were calculated
647 according to the best model of evolution. Samples that are in
648 the top-left half of the plot have a greater minimum
649 interspecific than maximum intraspecific divergence and
650 exhibit a barcode gap.





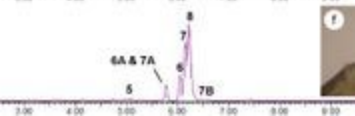
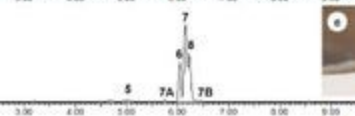
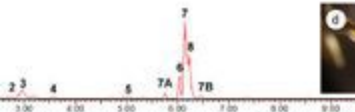
1
 H_2O_4 2
 H_3O_4 3



6

7

8



95%) = (-10.56, 8.105)

CI PD+



CI PD-



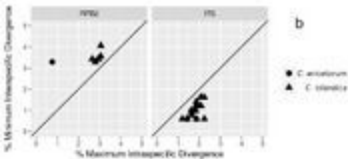
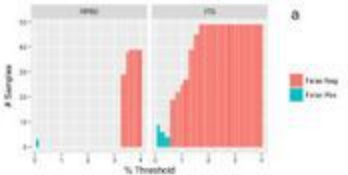
CI



-10 -8 -6 -4 -2 0 2 4 6 8

E1





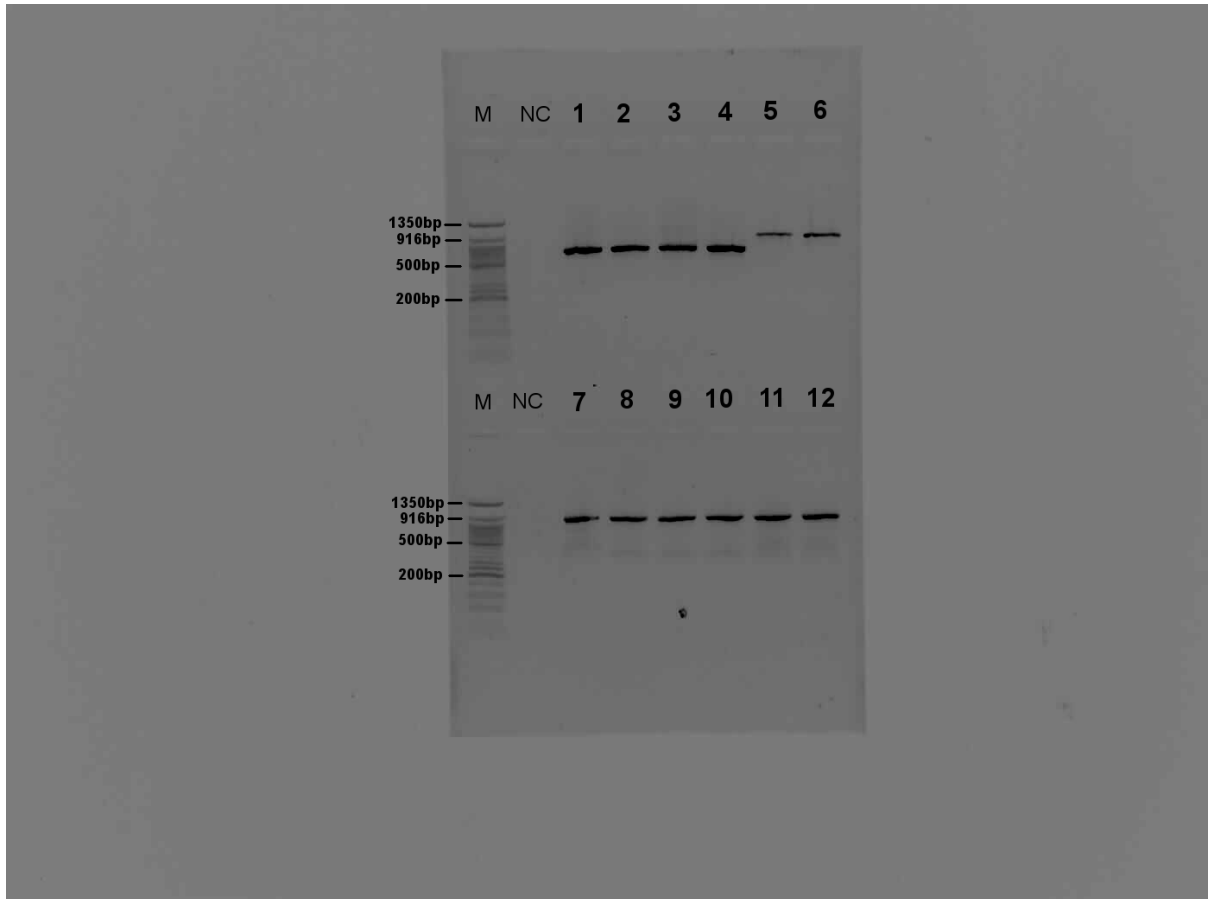


Figure S1. Agarose gel electrophoresis of PCR products from fungal nrITS and RPB2 loci. Lane 1-6: PCR products of fungal nrITS locus, ranging from 600 (intron-absent) to 900 bp (intron-present). Lane 7-12: PCR products of fungal RPB2 locus of ca. 900 bp. M: ladder. NC: negative control.

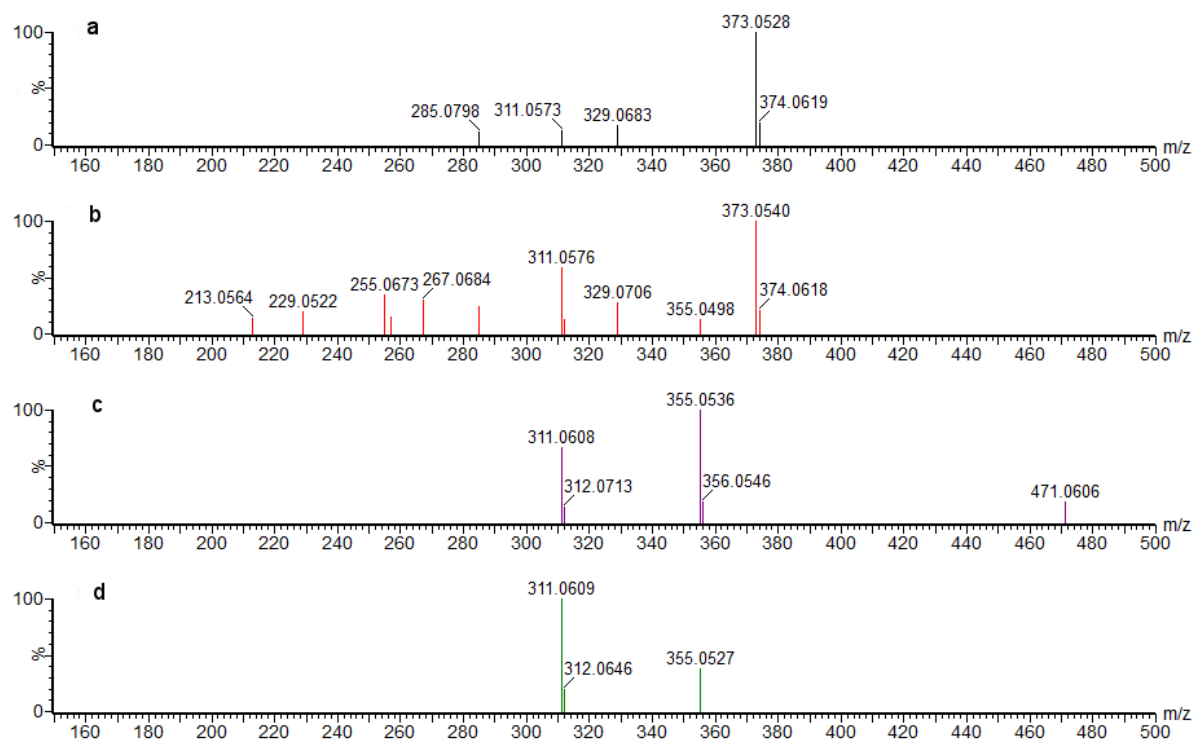


Figure S2. MS spectra of major depsidones in the PD+ *Cetraria islandica* chemotype. MS (a) and MS² (b) spectra for protocetraric acid **1**; MS (c) and MS² (d) spectra for fumarprotocetraric acid **3**.

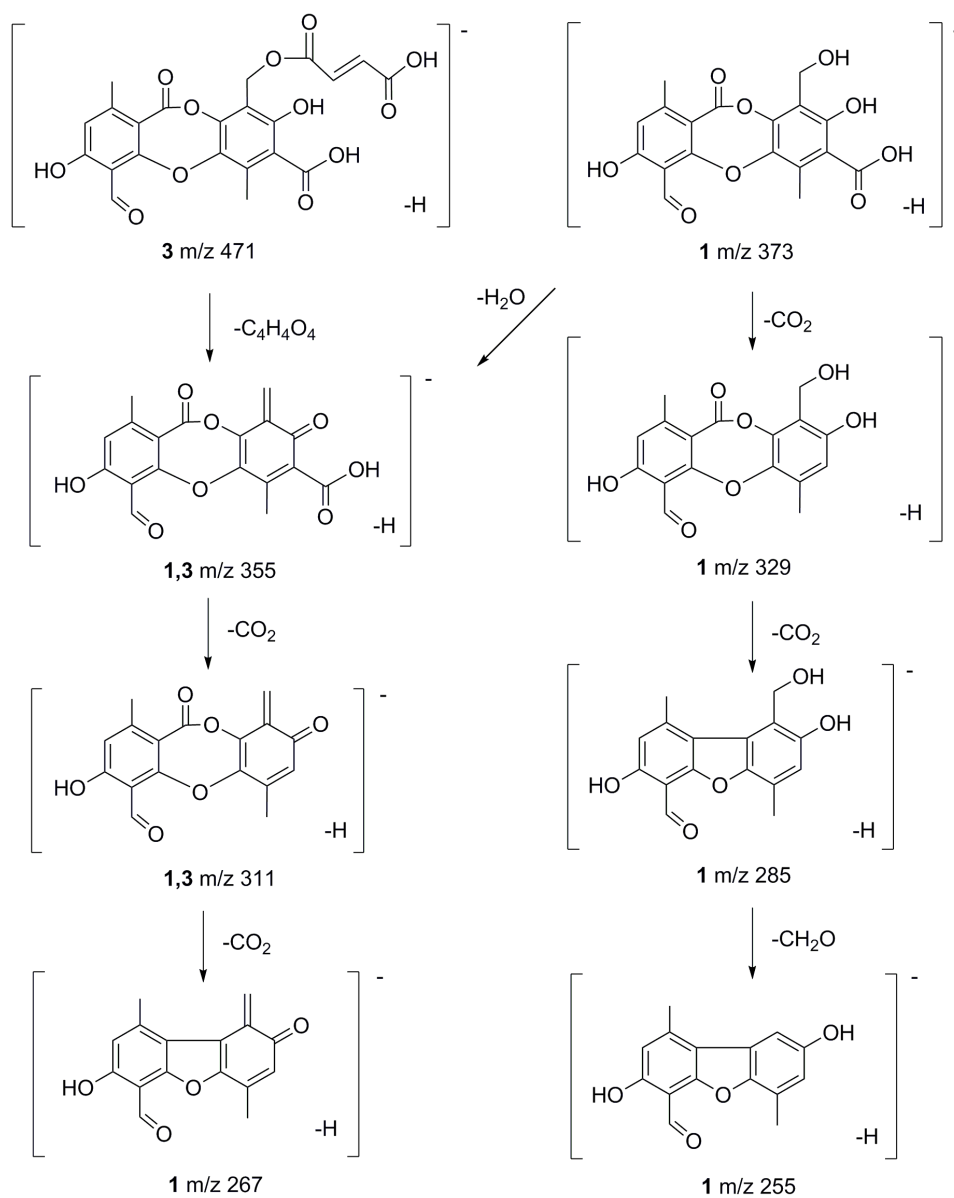


Figure S3. MS fragmentation patterns for major depsidones (protocetraric acid **1**; fumarprotocetraric acid **3**) in the PD+ *Cetraria islandica* chemotype.

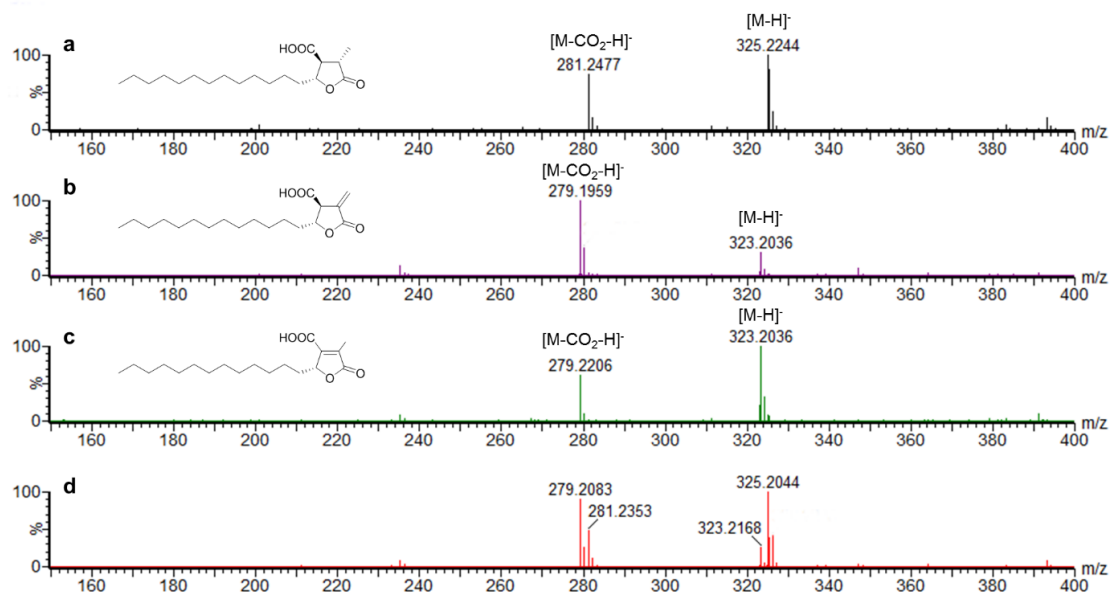


Figure S4. MS spectra for major paraconic acids in *Cetraria islandica* species complex. (a) MS spectrum of (+)-roccellaric acid **6**; (b) MS spectrum of (+)-protolichesterinic acid **7**; (c) MS spectrum of (+)-lichesterinic acid **8**. MS fragment ions of compounds **7** and **8** differed in the ratio of molecular ion $[M-H]^-$ to the decarboxylated molecular ion $[M-CO_2-H]^-$. The higher stability of lichesterinic acid molecular ion could be explained by hyperconjugation, where the electrons in the C-C bond between lactone ring and carboxylic group interacts with the unhybridized *p*-orbital in the adjacent ethylenic carbon; (d) MS spectrum of the peak eluting out at t_R 5.87 min containing two compounds **6A** and **7A** in Fig 3. They are tentatively identified as a stereoisomer (compound **6A**; m/z 325.2 and 281.2) of **6** and a stereoisomer (compound **7A**; m/z 323.2 and 279.2) of **7**, respectively.

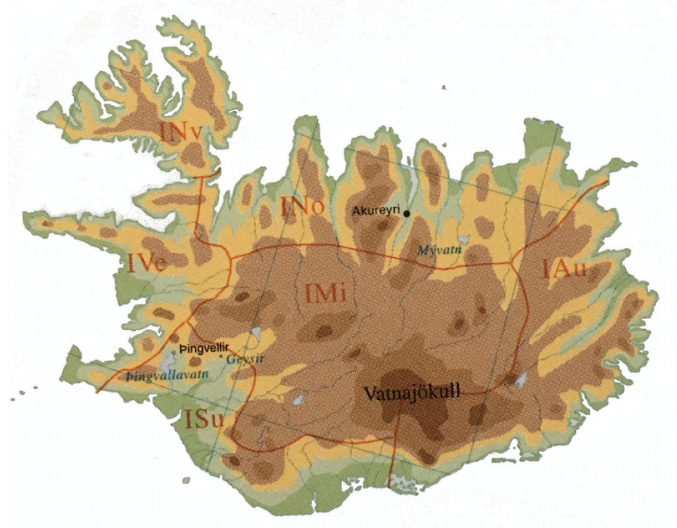
Table S1. Voucher specimens of *Cetraria islandica* species complex used in the current study, including country, collection date, voucher number, spot test results/chemotype, DNA isolate number and GenBank accession numbers.

| Country ^a | Collection date | Specimen voucher ^b | Spot test ^c | Collector | DNA Isolate | GenBank accession number | |
|---------------------------|-----------------|-------------------------------|------------------------|--------------------|-------------|--------------------------|----------|
| | | | | | | RPB2 | nrITS |
| <i>Cetraria islandica</i> | | | | | | | |
| Iceland: IVe | 21-Aug-2012 | LA31863 | PD+ | Starri Heidmarsson | CI1 | KY768945 | KY764967 |
| Iceland: IAu | 16-Aug-2012 | LA31864 | PD+ | Starri Heidmarsson | CI4 | KY768946 | KY764968 |
| Iceland: INo | 21-Aug-2012 | LA31865 | PD+ | Starri Heidmarsson | CI6 | KY768947 | KY764969 |
| Iceland: IVe | 23-Aug-2012 | LA31866 | PD+ | Starri Heidmarsson | CI7 | KY768948 | KY764970 |
| Iceland: INo | 8-Jul-2013 | LA31867 | PD- | Starri Heidmarsson | CI11 | KY768949 | KY764971 |
| Iceland: IVe | 23-Jul-2013 | LA31868 | PD+ | Starri Heidmarsson | CI12 | KY768950 | KY764972 |
| Iceland: IVe | 25-Jul-2013 | LA31869 | PD+ | Starri Heidmarsson | CI13 | KY768951 | KY764973 |
| Iceland: IVe | 25-Jul-2013 | LA31870 | PD+ | Starri Heidmarsson | CI14 | KY768952 | KY764974 |
| Iceland: INo | 16-Aug-2012 | LA31871 | PD+ | Starri Heidmarsson | CI15 | KY768953 | KY764975 |
| Iceland: ISu | 12-Jul-2013 | LA31872 | PD+ | Starri Heidmarsson | CI16 | KY768954 | KY764976 |
| Iceland: IVe | 11-Jul-2013 | LA31873 | PD+ | Starri Heidmarsson | CI17 | KY768955 | KY764977 |
| Iceland: IVe | 11-Jul-2013 | LA31874 | PD+ | Starri Heidmarsson | CI18 | KY768956 | KY764978 |
| Iceland: IVe | 23-Jul-2013 | LA31875 | PD+ | Starri Heidmarsson | CI19 | KY768957 | KY764979 |
| Iceland: IVe | 9-Aug-2013 | LA31876 | PD+ | Starri Heidmarsson | CI25 | KY768958 | KY764980 |
| Iceland: IVe | 23-Jul-2013 | LA31877 | PD+ | Starri Heidmarsson | CI26 | KY768959 | KY764981 |
| Iceland: IVe | 8-Jul-2013 | LA31878 | PD+ | Starri Heidmarsson | CI27 | KY768960 | KY764982 |
| Iceland: IVe | 8-Jul-2013 | LA31879 | PD+ | Starri Heidmarsson | CI28 | KY768961 | KY764983 |
| Iceland: IVe | 11-Jul-2013 | LA31880 | PD+ | Starri Heidmarsson | CI29 | KY768962 | KY764984 |
| Iceland: IVe | 8-Jul-2013 | LA31881 | PD+ | Starri Heidmarsson | CI30 | KY768963 | KY764985 |
| Iceland: IVe | 21-Aug-2013 | LA31882 | PD+ | Starri Heidmarsson | CI31 | KY768964 | KY764986 |
| Iceland: IVe | 8-Jul-2013 | LA31883 | PD+ | Starri Heidmarsson | CI32 | KY768965 | KY764987 |
| Iceland: INo | 24-Jul-2012 | LA31884 | PD- | Starri Heidmarsson | CI57a | KY768966 | KY764988 |
| Iceland: INo | 24-Jul-2012 | LA31885 | PD- | Starri Heidmarsson | CI57b | KY768967 | KY764989 |
| Iceland: INv | 30-Aug-2013 | LA31886 | PD+ | Starri Heidmarsson | CI58 | KY768968 | KY764990 |
| Iceland: IAu | 15-Aug-2012 | LA31887 | PD+ | Starri Heidmarsson | CI59 | KY768969 | KY764991 |

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|-----------------------------------|-------------|----------------|-----|--------------------|--------|----------|----------|
| Iceland: INv | 22-Aug-2013 | LA31888 | PD- | Starri Heidmarsson | CI60 | KY768970 | KY764992 |
| Iceland: INv | 26-Aug-2013 | LA31889 | PD- | Starri Heidmarsson | CI61 | KY768971 | KY764993 |
| Iceland: INo | 14-Aug-2012 | LA31890 | PD+ | Starri Heidmarsson | CI62 | KY768972 | KY764994 |
| Iceland: IVe | 15-Aug-2012 | LA31928 | PD+ | Starri Heidmarsson | CI63 | KY768973 | KY764995 |
| Iceland: INo | 21-Aug-2012 | LA31891 | PD+ | Starri Heidmarsson | CI64 | KY768974 | KY764996 |
| Iceland: INo | 28-Jun-2012 | LA31929 | PD- | Starri Heidmarsson | CI65 | KY768975 | KY764997 |
| Iceland: IVe | 12-Jul-2013 | LA31892 | PD+ | Starri Heidmarsson | CI66 | KY768976 | KY764998 |
| Iceland: INv | 22-Aug-2013 | LA31893 | PD- | Starri Heidmarsson | CI67 | KY768977 | KY764999 |
| Iceland: INo | 24-Jun-2012 | LA31894 | PD+ | Starri Heidmarsson | CI68 | KY768978 | KY765000 |
| Iceland: INo | 2012 | LA31895 | PD- | Starri Heidmarsson | CI69 | KY768979 | KY765001 |
| Iceland: INo | 2012 | LA31896 | PD- | Starri Heidmarsson | CI70 | KY768980 | KY765002 |
| Iceland: INo | 8-Aug-2012 | LA31897 | PD+ | Starri Heidmarsson | CI77 | KY768981 | KY765003 |
| Iceland: INo | 28-Jun-2012 | LA31898 | PD+ | Starri Heidmarsson | CI78a | KY768982 | KY765004 |
| Iceland: INo | 28-Jun-2012 | LA31899 | PD+ | Starri Heidmarsson | CI78b | KY768983 | KY765005 |
| Iceland: INo | 23-Aug-2012 | LA31927 | PD- | Starri Heidmarsson | CI87a | KY768984 | KY765006 |
| Iceland: INo | 11-Jul-2002 | LA30017 | PD- | Hordur Kristinsson | CI115 | KY768985 | KY765007 |
| Iceland: INo | 14-Aug-2012 | LA31900 | PD- | Starri Heidmarsson | CI117a | KY768986 | KY765008 |
| Iceland: INo | 10-Aug-2006 | LA31128 | PD- | Hordur Kristinsson | CI113 | - | - |
| Iceland: INo | 10-Jul-1998 | LA17549 | PD- | Hordur Kristinsson | CI36 | - | - |
| Iceland: INo | 5-Jul-1998 | LA17221 | PD- | Hordur Kristinsson | CI37 | - | - |
| <i>Cetraria ericetorum</i> | | | | | | | |
| Iceland: IAu | 10-Aug-1997 | LA18976 | PD- | Hordur Kristinsson | CE1 | - | KY765009 |
| Finland: Sodankylä | 21-Aug-2003 | NO2530 | PD- | Beata Krewicka | CE6 | KY768987 | KY765010 |
| Sweden: Uppsala | 20-May-2002 | NO23002 | PD- | Leif Tibell | CE8 | KY768988 | KY765011 |
| Sweden: Uppsala | 18-Oct-2015 | NO5626 | PD- | Stefan Ekman | CE11 | KY768989 | KY765012 |
| Iceland: IAu | 13-Jul-2014 | LA20746 | PD- | Hordur Kristinsson | CE13 | KY768990 | KY765013 |
| Iceland: INo | 29-Aug-2016 | LA31901 | PD- | Hordur Kristinsson | CE15 | KY768991 | KY765014 |
| Iceland: INo | 1-Sep-2010 | LA31538 | PD- | Hordur Kristinsson | CE16 | KY768992 | KY765015 |
| Iceland: IAu | 9-Aug-1997 | LA27354 | PD- | Hordur Kristinsson | CE2 | - | - |

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|-------------------|-------------|----------------|-----|--------------------|------|---|---|
| Sweden: Gävleborg | 15-Jun-1997 | NO501 | PD- | Ake Agren | CE3 | - | - |
| Russian: Komi | 6-Jul-2000 | L135019 | PD- | - | CE4 | - | - |
| Canada: Quebec | 2-Jul-1999 | NO5021 | PD- | Jan-Eric Mattsson | CE5 | - | - |
| Poland: Silesia | 19-Jul-1998 | KO2101 | PD- | - | CE7 | - | - |
| Russia: Komi | 2-Jul-1997 | NO7971 | PD- | Björn Larsson | CE9 | - | - |
| Iceland: INo | 31-Jul-1996 | NO720 | PD- | Starri Heidmarsson | CE10 | - | - |
| Iceland: INo | 18-Aug-1998 | LA20809 | PD- | Hordur Kristinsson | CE12 | - | - |
| Iceland: INo | 7-Jun-1998 | LA18310 | PD- | Hordur Kristinsson | CE14 | - | - |
| Iceland: INo | 19-Aug-1998 | LA29284 | PD- | Hordur Kristinsson | CE17 | - | - |
| Iceland: INo | 10-Aug-1997 | LA18982 | PD- | Hordur Kristinsson | CE18 | - | - |

^a INo, INv, IVe, IMi, IAu and ISu refer to corresponding area in Icelandic map below;



^b Authentic herbarium specimens are marked in boldface;

^c Spot testing/chemotype identification results are reported as PD+ (medullary red color after *p*-phenylendiamine staining) and PD- (no red color after *p*-phenylendiamine staining).

Table S2. Chromatographic and MS data of metabolites tentatively identified from acetone extracts of taxa in the *Cetraria islandica* species complex.

| t_R (min) ^a | $[M-H]^-$ (m/z) ^b | Product ions (m/z) ^c | Mass error (ppm) ^d | Molecular formula | Compound ^e | Lichen ^f |
|-----------------------------|---------------------------------|--|----------------------------------|---|---|---------------------|
| 2.48 | 373.0540 | 355.0498, 329.0706, 311.0576, 285.0798 | -5.4 | C ₁₈ H ₁₄ O ₉ | Protocetraric acid 1 | CI (PD+) |
| 2.55 | 385.0650 | 341.0783, 329.2408 | - | - | Unidentified | CE |
| 2.67 | 487.0986 | 373.0665, 355.0501 , 311.0594 | - | - | Unidentified | CI (PD+) |
| 2.75 | 473.0806 | 355.0451 , 311.0550 | -0.8 | C ₂₂ H ₁₈ O ₁₂ | Succinprotocetraric acid 2 | CI (PD+) |
| 2.88 | 517.1052 | 401.0900, 369.0647, 325.0730 | 6.7 | - | Unidentified | CI (PD+) |
| 2.96 | 471.0536 | 355.0467 , 311.0581 | -1.4 | C ₂₂ H ₁₆ O ₁₂ | Fumarprotocetraric acid 3 | CI (PD+) |
| 3.07 | 489.3547 | 355.0500, 343.0474 , 311.0598, 299.0618 | - | - | Unidentified | CI (PD+) |
| 3.17 | 387.0728 | 355.0474, 343.0864, 311.0580, 299.0962 | 3.1 | C ₁₉ H ₁₆ O ₉ | Unidentified | CI (PD+) |
| 3.21 | 293.1744 | 236.1066, 221.1552 | -3.1 | C ₁₇ H ₂₆ O ₄ | Unidentified | CI, CE |
| 3.34 | 431.3405 | 355.0503, 309.1720 | 7.4 | C ₂₄ H ₄₈ O ₆ | Unidentified | CI (PD+) |
| 3.57 | 357.0607 | 313.0723, 269.0848 | -0.8 | C ₁₈ H ₁₄ O ₈ | Virensic acid 4 | CI (PD+) |
| 4.17 | 295.2257 | 277.2194, 171.1052 | -5.4 | C ₁₈ H ₃₂ O ₃ | Unidentified | CE |
| 4.71 | - | 443.3083, 279.2310 , 250.1470 | -5.0 | - | Unidentified | CI, CE |
| 4.94 | - | 297.2133, 279.2383 , 264.1647, 253.2214 | - | - | Unidentified | CI, CE |
| 5.03 | 279.2364 | 251.2069 | - | - | Unidentified | CI, CE |
| 5.09 | 295.1935 | 251.2062 | 8.8 | C ₁₇ H ₂₈ O ₄ | Nephrosterinic acid 5 | CI, CE |
| 5.44 | - | 311.2299, 281.2556 | - | - | Unidentified | CI, CE |
| 5.77 | 323.2168 | 279.2314 | -3.6 | C ₁₉ H ₃₂ O ₄ | A stereoisomer of (+)-Protolichesterinic acid 7A | CI, CE |
| 5.77 | 325.2370 | 281.2527 | -2.8 | C ₁₉ H ₃₄ O ₄ | A stereoisomer of (+)-Roccellaric acid 6A | CE |
| 6.05 | 325.2405 | 281.2511 | 8.0 | C ₁₉ H ₃₄ O ₄ | (+)-Roccellaric acid 6 | CI, CE |
| 6.17 | 323.2234 | 279.2322 | -0.7 | C ₁₉ H ₃₂ O ₄ | (+)-Protolichesterinic acid 7 | CI, CE |
| 6.23 | 323.2224 | 279.2372 | 0.6 | C ₁₉ H ₃₂ O ₄ | Lichesterinic acid 8 | CI, CE |
| 6.46 | 323.2218 | 279.2336 | 4.5 | C ₁₉ H ₃₂ O ₄ | A stereoisomer of (+)-Protolichesterinic acid 7B | CI, CE |

^a t_R means retention time;^b $[M-H]^-$ stands for deprotonated molecular ion;^c The product ion is marked in bold when it is the base peak in the MS spectrum;^d The mass error of the base peak is provided;^e Major compounds **1-8** are labelled corresponding to structures in Figure 2;^f The presence of lichen compounds in lichen taxa. CI (PD+): the PD+ chemotype of *Cetraria islandica*; CI: both PD+ and PD- chemotypes of *C. islandica*; CE: *C. ericetorum*.