# Investigating the Clitellata (Annelida) of Icelandic springs with alternative barcodes

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DNA barcoding is an invaluable tool to identify clitellates, regardless of life stage or cryptic morphology. However, as COI (the standard barcode for animals) is relatively long (658 bp), sequencing it requires DNA of high quality. When DNA is fragmented due to degradation, alternative barcodes of shorter length present an option to obtain genetic material. We attempted to sequence 187 clitellates sampled from springs in Iceland. However, the material had been stored at room temperature for two years, and DNA of the worms had degraded, and only three COI sequences were produced (i.e., <2%) success rate). Using two alternative barcodes of 16S (one ca. 320 bp, the other ca. 70 bp long) we increased the number of sequenced specimens to 51. Comparisons of the 16S sequences showed that even the short 70 bp fragment contained enough genetic variation to separate all clitellate species in the material. Combined with morphological examinations we recognized a total of 23 species, where at least 8 are new records for Iceland, some belonging to genera new for Iceland: Cernosvitoviella and Pristina. All the new taxa are included in an updated species list of Icelandic Clitellata. The material revealed some stygophilic species previously known to inhabit springs, but true stygobionts, which are restricted to groundwater habitats, were not found. Our study shows that short 16S fragments can be obtained from DNA too degraded to be used in traditional COI barcoding, and contain enough genetic variation to separate closely related clitellate species.

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### INTRODUCTION

The clitellate fauna ("oligochaetes" and leeches) of Iceland was early on studied by Černosvitov (1929, 1931, 1936) and Nielsen & Christensen (1959), but also more specifically reviewed by Bruun (1938a, 1938b) for Hirudinea, Backlund (1949) for Lumbricidae, Hrabě (1952) for Lumbriculidae and Naididae (including the former Tubificidae), Christensen (1962) and Nurminen (1973) for Enchytraeidae, and Erséus (1976) for marine Enchytraeidae and Naididae. Since then, a few

additional species have been reported in scattered publications on either pure taxonomy or more general biological issues, such as parasitology (for marine leeches) or ecology; for references, see updated species list for Iceland below. The previous studies report clitellates mainly from soils, lakes, rivers and seashores, but not from freshwater springs, which are the focus of this study.

Springs represent ecotones between groundwater and surface water and give rise to specialized invertebrate communities. On the European mainland, groundwater clitellates are rather well known, with many species endemic to various regions (e.g., Sambugar *et al.* 1999; Giani *et al.* 2001, 2011; Achurra & Rodriguez 2008; Bojková *et al.* 2011; Martin *et al.* 2015). Groundwater and spring invertebrate communities in Iceland were recently investigated by Govoni *et al.* (2018) and Kreiling *et al.* (2018), but these studies focused on insects and crustaceans and the clitellate diversity in Icelandic springs has until now been largely unknown.

In the present study, we examined clitellates collected as a part of a survey on invertebrate fauna in freshwater springs around Iceland. With the intent to save time, we decided to identify the material primarily using molecular data rather than by traditional morphological examination; and to our knowledge, there have not been any published studies of clitellates from Iceland containing genetic sequences, to this date. DNA barcoding (e.g., Hebert et al. 2003, 2004) would allow us to identify juvenile specimens and possible cryptic species. We would then corroborate the identity of the successfully sequenced specimens also by morphological observations. Although this procedure did not work exactly as first intended, the aim to present all the identified species from the springs will still be achieved in this paper. We will also provide an updated list of all species of Clitellata known from Iceland.

# MATERIAL AND METHODS

Worms were collected from 31 springs during the summer of 2015 as a part of a broader study on spring invertebrates in Iceland (Kreiling *et al.* in prep.). A Surber sampler (0.093 m<sup>2</sup>) with 63  $\mu$ m mesh size was used for collection of clitellates in the benthic substrate of the spring, and electrobugging (Lento & Morin 2014, Kreiling *et al.* 2018) was used for collection of invertebrates in the spring source. The clitellates were stored in 96 % ethanol at room temperature (~20°C) for about two years before further processing. As described below, identification to species level was unsuccessful for a part of the collection, and in the end, the results of the study were based on material from only 19 of the freshwater springs (Figure 1; Table 1).

The clitellate specimens were first examined under a stereomicroscope and the amputated posterior ends of 187 specimens were used for DNA extraction using the QuickExtract DNA Extraction Solution 1.0 (Epicentre, Madison, WI, USA), following the manufacturer's instructions.

The original idea was to barcode all selected specimens using the standard animal barcode COI (cytochrome c oxidase subunit I). However, the DNA proved to have deteriorated considerably (probably due to prolonged storage at room temperature, possibly in too low alcohol concentration), as we were unable to obtain COI sequences for a vast majority of the



Figure 1. Map of the 19 freshwater springs in Iceland from which clitellata have been collected and successfully sequenced in this study.

Table I. List of the sampling locations with habitat description and some abiotic measurements. Spring names refer either to the name of the water body (stream or lake), or to the surrounding area, or the closest farm. Limnocrene (L) springs form pools of standing water, whereas rheocrene (R) springs originate streams. Elev. = Elevation, O.S. = Oxygen Saturation.

Spring (type)	Habitat	Latitude Longitude (WGS84)	Elev. [m]	Sampling date	Water temp. [°C]	рН	O.S. [%]
Botnar I (R) SE Iceland	Gushing, shallow spring at the edge of lava field; fine sand	63°38.707' N 018°14.749' W	36	10 July 2015	5.6	8.0	74.2
Botnar II (L) SE Iceland	Spring emerging from lava field; low primary production; fine sand	63°39.275' N 018°15.142' W	33	10 July 2015	7.4	7.9	78.2
Enni (R) NW Iceland	Spring forming a small stream on a grassy hillslope; sand	65°53.371 N 019°19.755' W	151	19 September 2015	4.5	7.3	75.9
Galtalækur (R) S Iceland	Spring forming a small stream in wooded area; high density of surrounding vegetation (grasses, shrubs and trees); gravel	64°00.453' N 019°55.148' W	128	8 July 2015	5.1	7.9	72.3
Grænavatn Norður (L) NE Iceland	Spring on lake shore; gravel and mud	65°32.905' N 016°58.908' W	291	22 July 2015	6.5	8.9	60.6
Grænavatn Suður (L) NE Iceland	Spring on lake shore; high primary production; lava rock	65°32.205' N 017°00.477' W	285	22 July 2015	4.5	9.0	63.3
Hengill IS8 (R) S Iceland	Hot spring forming a stream in geothermal area; high primary production; rock	64°03.414'N 021°18.439' W	381	13 July 2015	16.6	7.5	66.1
Herðubreiðarlindir (L) Central Highlands	Big spring in the Central Highlands, forming a deep stream; high density of surrounding vegetation (grasses and shrubs); fine sand	65°11.548' N 016°13.508' W	493	16 August 2015	5.9	6.8	65.6
Hruni (L) SE Iceland	Spring in a garden pond; mud and gravel	63°51.547' N 017°44.486' W	43	11 July 2015	3.5	7.9	75.8
Kálfaströnd (L) NE Iceland	Spring on lake shore; lava rock and sand	65°33.759' N 016°56.710' W	283	21 July 2015	5.1	9.2	54.0
Krákárbotnar (R) Central Highlands	Small, isolated spring in the Central Highlands with almost no surrounding vegetation; sand	65° 19.852' N 017°04.654' W	430	26 July 2015	8.6	8.8	69.5
Langivogur (L) NE Iceland	Hot spring at lake shore; high primary production; lava rock	65°37.012' N 016°55.000' W	286	23 July 2015	19.8	8.3	75.1
Lón (L) NE Iceland	Spring at the shore of a shallow lagoon; lava rock	66°05.785' N 016°55.514' W	6	24 July 2015	4.9	8.0	77.8
Lækjarbotnar Hol (R) S Iceland	Spring forming a small stream on meadow; high density of surrounding vegetation (grasses); lava rock and sand	63°57.422' N 020°15.892' W	78	8 July 2015	5.5	7.9	75.6
Miðhúsaskógur (L) S Iceland	Spring at the shore of shallow pond; low primary production; fine sand and lava rock	64°17.373' N 020°30.706' W	184	8 July 2015	2.4	9.3	78.1
Staðarhraun Bær (R) W Iceland	Spring at the edge of lava field, form- ing a small stream; lava rock and gravel	64°44.610' N 022°05.647' W	62	28 July 2015	5.1	5.3	79.0
Staðarhraun Kirkja (R) W Iceland	Spring at the edge of lava field; sand and lava rock	64°44.855' N 022°05.812' W	62	28 July 2015	4.6	5.3	79.2
Steinsstaðir (R) NW Iceland	Hot spring forming a small stream; high primary production; sand and mud	65°28.162' N 019°21.390' W	62	4 August 2015	40.24	8.47	86.3
Þverá (L) SE Iceland	Spring in shallow pond; sand	63°52.396' N 017°49.199' W	53	11 July 2015	5.1	7.5	76.1

specimens. Additionally, as most worms were tiny the DNA samples as such were small. DNA is known to degrade faster when stored at room temperature rather than in a freezer (Vink et al. 2005), but old fragmented DNA can still be amplified by using primers that target a shorter gene fragment (Hajibabaei et al. 2006). There is an alternative reverse primer developed for naidid clitellates (COI-E, Bely & Wray 2004), but as it targets the same sequence length as the primers we used (Folmer et al. 1994) we reasoned it would not handle fragmented DNA better and thus did not use it. A number of universal primers have been developed targeting shorter fragments of the COI gene, to manage degraded DNA mainly in metabarcoding studies (Meusiner et al. 2008, Leray et al. 2013). However, said primers have only been tested for a limited number of annelids and the universality of some has been questioned (Arif et al. 2011). As an alternative, mitochondrial 16S rDNA has been suggested as a more favourable option, particularly for metabarcoding, since this gene's highly conserved sites make it easy to develop universal primers (Deagle et al. 2014). As we have an interest in the potential for metabarcoding of clitellates we decided to try two shorter gene fragments of 16S.

PCRs were run for all specimens using Red Taq DNA Polymerase Master Mix (VWR, Haasrode, Belgium) in 25  $\mu$ L reactions, with the three primer pairs in Table 2. The PCR products were examined using electrophoresis on an agarose gel and the successful samples were purified using exonuclease I and FastAP thermosensitive alkaline phosphatase. The purified products were sequenced by MWG Eurofins Operon (Edersberg, Germany), and the resulting trace files were assembled in Geneious 6.1.8 (Drummond *et al.* 2011). The successful sequences were compared to data in the Barcoding of Life Database (BOLD) and GenBank (NCBI), in order to recognize barcode clusters representing putative species, and to genetically identify the specimens to species.

We also compared the sequences of the two 16S barcodes (one ca. 320 bp, the other ca. 70 bp long), by alignment and Neighbor-Joining (NJ) analyses in Geneious. The COI and longer 16S sequences were later uploaded to Genbank, the shorter 16S sequences are provided in the text below.

The vouchers (anterior ends) of 181 specimens, i.e., excluding most earthworms (Lumbricidae), were stained in paracarmine, dehydrated in xylene and mounted in Canada balsam on microscope slides following Erséus (1994). The vouchers of specimens identified to species, or in some cases generic, level were deposited in the Swedish Museum of Natural History (SMNH) and they are listed in Table 3. Reference specimens were also included to aid in the identification of two species with only immature specimens, whose barcodes could not be matched with anything in the public databases: CE19501 Fridericia dura, collected in Kristiansund, Möre og Romsdal, Norway, 63.1258 N, 7.7352 E by Christer Erséus, 13 Aug 2013, and CE22027 Marionina cf. argentea, collected in Nedrehus, Maurangerfjorden, Kvinnherad, Hordaland, Norway, 60.1295 N, 6.3146 E by Christer Erséus & Mårten Klinth, 14 May 2014, both deposited in the University Museum of Bergen (ZMBN) (Table 3).

## RESULTS

It soon became apparent that the DNA of the samples had degraded substantially, as we obtained successful COI sequences from only three of the 187 selected worms. They were genetically identified as *Bimastos rubidus* (Savigny, 1826) *sensu lato, Cernosvitoviella pusilla* Nurminen, 1973, and *Chaetogaster* cf. *diastrophus* (Gruithuisen, 1828), respectively (Table 3).

The longer (320 bp) of the two 16S barcodes was more successful than the COI barcode, but we still only got results for 54 specimens. Moreover, after examination of the microscope slides, the morphology did not agree with the DNA results for eight of these 54 worms, most likely due to DNA contamination, leaving only 46 individuals confidently identified by both DNA and morphology (Table 3 specifies how each specimen was identified).

Target	Primers	PCR program	Reference
COI 658 bp	LCO1490 (forward) GGTCAACAAATCATAAAGATATTGG HCO2198 (reverse) TAAACTTCAGGGTGACCAAAAAATCA	95°C 5 min, (35 cycles of 95°C 40 s, 45°C 45 s 72°C 1 min), 72°C 8 min	(Folmer <i>et al.</i> 1994)
16S ca. 320 bp	Ann16SF (forward) GCGGTATCCTGACCGTRCWAAGGTA Ann16SR (reverse) TCCTAAGCCAACATCGAGGTGCCAA	95°C 5 min, (35 cycles of 95°C 30 s, 50°C 30 s 72°C 1 min), 72°C 8 min	(Sjölin et al. 2005)
16S ca. 70 bp	ewD (forward) ATTCGGTTGGGGCGACC ewE (reverse) CTGTTATCCCTAAGGTAGCTT	95°C 5 min, (35 cycles of 95°C 30 s, 58°C 30 s 72°C 10 s), 72°C 5 min	(Bienert et al. 2012)

Table 2. Primers and PCR programs used to sequence COI and 16S.

Table 3. List of Icelandic specimens used in this study with specimen ID's (identification numbers), identification method (B = BOLD; G = Genbank; M = morphology; R = based on match to other non-Icelandic reference material, presented at the end of the table, and with sampling sites specified in material and methods), the spring in which they were collected, Genbank accession numbers and museum voucher ID's. We only deposited the longer 16S fragment. Sequences from the shorter 16S fragment are presented in the text of the Results. More detailed description of the springs in Table 1.

Taxon	Specimen	Spring locality	Genbank acc. no.			Voucher ID	
(Identification method)	ID		COI	16S (320 bp)	16S (70 bp)		
Enchytraeidae							
Cernosvitoviella aggtelekiensis Dózsa-Farkas, 1970 (GM)	CE30974	Hruni		MK837025		SMNH 176517	
<i>Cernosvitoviella</i> cf. <i>minor</i> Dózsa-Farkas, 1990 (GM)	CE31592	Hruni			Sequence in Results	SMNH 176518	
<i>Cernosvitoviella pusilla</i> Nurminen, 1973 (M)	CE30979	Botnar II		MK837026	(Overlapping)	SMNH 176519	
<i>Cernosvitoviella pusilla</i> Nurminen, 1973 (B)	CE31607	Staðarhraun Bær	MK837024	MK837027	(Overlapping)	SMNH 176520	
Cognettia varisetosa (Martinsson, Rota & Erséus, 2015) (M)	CE30958	Enni				SMNH 176522	
Enchytraeus buchholzi 1 Vejdovský, 1879 (M)	CE30973	Hruni		MK837028	(Overlapping)	SMNH 176523	
Enchytraeus buchholzi 1 Vejdovský, 1879	CE31564	Botnar I			Sequence in Results	SMNH 176524	
Enchytraeus buchholzi 2 Vejdovský, 1879 (M)	CE31504	Langivogur		MK837029		SMNH 176525	
<i>Fridericia dura</i> (Eisen, 1879) (R)	CE30963	Krákárbotnar		MK837030	(Overlapping)	SMNH 176526	
Henlea perpusilla Friend, 1911 (G)	CE30978	Botnar II		MK837031	(Overlapping)	SMNH 176527	
Lumbricillus arenarius (Michaelsen, 1889) (GM)	CE31573	Herðubreiðarlindir		MK837032		SMNH 176528	
Lumbricillus arenarius (Michaelsen, 1889) (GM)	CE31575	Herðubreiðarlindir		MK837033		SMNH 176529	
Lumbricillus arenarius (Michaelsen, 1889) (GM)	CE31577	Herðubreiðarlindir		MK837034	(Overlapping)	SMNH 176530	
Marionina cf. argentea (Michaelsen, 1889) (R)	CE31590	Þverá		MK837035		SMNH 176531	
Marionina sp.	CE31579	Herðubreiðarlindir		MK837036		SMNH 176532	
Marionina sp.	CE31580	Herðubreiðarlindir		MK837037		SMNH 176533	
Marionina sp.	CE31583	Þverá		MK837038		SMNH 176534	
Marionina sp.	CE31587	Þverá		MK837039		SMNH 176535	
Marionina sp.	CE31589	Þverá		MK837040		SMNH 176536	
Marionina sp.	CE31603	Grænavatn Norður		MK837041		SMNH 176537	
Mesenchytraeus cf. armatus (Levinsen, 1884) (GM)	CE30954	Miðhúsaskógur		MK837042	(Overlapping)	SMNH 176538	
Mesenchytraeus cf. armatus (Levinsen, 1884) (GM)	CE30968	Grænavatn Suður			Sequence in Results	SMNH 176539	
Mesenchytraeus cf. armatus (Levinsen, 1884) (GM)	CE30972	Hruni		MK837043	(Overlapping)	SMNH 176540	

#### Table 3. Continued.

Taxon (Identification method)	Specimen ID	Spring locality	COL	Genbank acc. no 16S (320 bp)	16S (70 bp)	Voucher ID	
Mesenchytraeus cf. armatus (Levinsen, 1884) (GM)	CE30986	Botnar II	001	MK837045	(Overlapping)	SMNH 176542	
Lumbricidae							
Aporrectodea caliginosa (Savigny, 1826) (G)	CE30987	Staðarhraun Bær		MK837046	(Overlapping)	SMNH 176543	
Bimastos rubidus s. lat. (Savigny, 1826) (GB)	CE30982	Botnar II	MK837022		Sequence in Results	SMNH 176544	
Dendrobaena octaedra (Savigny, 1826) (G)	CE30975	Steinsstaðir			Sequence in Results	SMNH 176545	
Dendrobaena octaedra (Savigny, 1826) (G)	CE31506	Galtalækur		MK837047		SMNH 176546	
Eiseniella tetraedra (Savigny, 1826) (G)	CE30950	Langivogur		MK837048	(Overlapping)	SMNH 176547	
Naididae							
Chaetogaster cf. diastrophus (Gruithuisen, 1828) (G)	CE31491	Staðarhraun Kirkja	MK837023	MK837049		SMNH 176548	
Chaetogaster sp. = langi? (M)	CE31604	Grænavatn Norður		MK837050		SMNH 176549	
<i>Nais communis/variabilis</i> spe- cies complex, morphotype A3 (Envall <i>et al.</i> 2012) (G)	CE30951	Langivogur		MK837051		SMNH 176550	
Nais elinguis Müller, 1773 (GM)	CE30948	Lón		MK837052		SMNH 176551	
Nais elinguis Müller, 1773 (GM)	CE30949	Lón		MK837053		SMNH 176552	
Nais elinguis Müller, 1773 (GM)	CE30967	Lækjarbotnar Hol		MK837054		SMNH 176553	
Nais elinguis Müller, 1773 (GM)	CE30971	Þverá		MK837055		SMNH 176554	
Nais elinguis Müller, 1773 (GM)	CE30980	Botnar II		MK837056		SMNH 176555	
Nais elinguis Müller, 1773 (GM)	CE30981	Botnar II		MK837057		SMNH 176556	
Nais elinguis Müller, 1773 (GM)	CE30983	Botnar II		MK837058		SMNH 176557	
Nais elinguis Müller, 1773 (GM)	CE30984	Botnar II		MK837059		SMNH 176558	
Nais elinguis Müller, 1773 (GM)	CE30985	Botnar II		MK837060		SMNH 176559	
Nais elinguis Müller, 1773 (GM)	CE31480	Kálfaströnd		MK837061		SMNH 176560	
Nais elinguis Müller, 1773 (GM)	CE31493	Staðarhraun Kirkja		MK837062		SMNH 176561	
Nais elinguis Müller, 1773 (GM)	CE31581	Lækjarbotnar Hol		MK837063		SMNH 176562	
Nais elinguis Müller, 1773 (GM)	CE31582	Grænavatn Suður		MK837064		SMNH 176563	
Nais elinguis Müller, 1773 (GM)	CE31605	Staðarhraun Bær		MK837065		SMNH 176564	
Nais elinguis Müller, 1773 (GM)	CE31606	Staðarhraun Bær		MK837066		SMNH 176565	
Nais elinguis Müller, 1773 (GM)	CE31619	Staðarhraun Bær		MK837068		SMNH 176567	

Table 3. Continued.

Taxon	Specimen	Spring locality		Genbank acc. no.		Voucher ID
(Identification method)	ID		COI	16S (320 bp)	16S (70 bp)	
Nais elinguis Müller, 1773 (GM)	CE31620	Staðarhraun Bær		MK837069		SMNH 176568
Pristina foreli (Piguet, 1907) (M)	CE30943- 45	Hengill IS8				SMNH 176569
<i>Tubifex</i> cf. <i>tubifex</i> (Müller, 1774) (G)	CE31560	Botnar I		MK837070		SMNH 176570
Uncinais uncinata (Ørsted, 1842) (M)	CE31593	Hruni				SMNH 176571
Non-Icelandic reference material						
<i>Fridericia dura</i> (Eisen, 1879) (M)	CE19501	Norway	MN395701	MN394410 (478 bp)		ZMBN 110172
Marionina cf. argentea (Michaelsen, 1889) (M)	CE22027	Norway	MN395702	MN394411 (474 bp)		ZMBN 110740

The shorter 16S barcode (70 bp) only produced 17 successful sequences (Table 3), mostly from specimens already successfully barcoded with the longer 16S fragment, increasing the total number of DNA-barcoded (but non-contaminated) specimens to 51 (27 % of the original 187 specimens). The sequences of the five specimens that were successfully sequenced only for the shorter 16S fragment (i.e., five sequences not overlapping with our longer 16S uploaded on Genbank) are presented here (note that the sequence for CE31592 is incomplete):

#### CE31592 Cernosvitoviella cf. minor

TTGGGGCGACCAAGGAAAAATCATCCTTAATAAAAA AGACATAC; CE31564 Enchytraeus buchholzi 1 ATTCGGTTGGGGCGACCCAGGATAAATCATCCTGTAA AAAATAGACAAATATGTCAACCATATGAACCTAGTTA GATCACAGATCAAGCTACCTTAGGGATAACAGA; CE30968 Mesenchytraeus cf. armatus TATTCGGTTGGGGGCGACCATGGATAAATCATCCATAA TTTATAAGACAAACTAGTCATTAATAGATCCTTTTAAG

ATCACAGAATCAAGCTACCTTAGGGATAACAGA;

CE30982 Dendrodrilus rubidus

ATTCGGTTGGGGCGACCAGGGAAATAATAAATCATCC CTCATTATAAGATAAATAAATCTCTCCCATGACCCTTG AAAAAGATCAAAAAACCAAGCTACCTTAGGGATAAC AGA:

CE30975 Dendrobaena octaedra

TATTCGGTTGGGGCGACCAGGGAAATTAATAATCATC CCTTAGTCAAAGATTTATTAATCTATAAATAAGACCCT ACTAAGATCTAAAGAACAAGCTACCTTAGGGATAACA GA.

Some worms were thus successfully sequenced only for one or two of the three barcode markers.

In the NJ analyses, both the 320 bp (Figure 2) and 70 bp (Figure 3) 16S barcodes clustered specimens of the same

species, and clearly separated the recognized species from each other.

Among the 51 DNA-barcoded individuals we identified 20 different species, at least six of which are new records for Iceland (Table 4): *Cernosvitoviella aggtelekiensis* Dózsa-Farkas, 1970, *C. cf. minor* Dózsa-Farkas, 1990, *C. pusilla, Fridericia dura* (Eisen, 1879), *Mesenchytraeus* cf. *armatus* (Levinsen, 1884), and *Chaetogaster* cf. *diastrophus*. Among the barcoded worms, we also found a small specimen of *Chaetogaster*, which is possibly *Ch. langi* Bretscher, 1896 (previously known from Iceland; Hrabě 1952), but not yet confidently identified. Its 16S barcode (320 bp) matches a species also found in Sweden, Norway and the Azores (Klinth & Erséus, unpublished data). The species referred to as *Marionina* sp. could also potentially be new to Iceland.

In the barcoded material, most species were represented by a single or a few specimens only, except Nais elinguis Müller, 1773, for which we obtained 16S (320 bp) sequences from 18 individuals (Table 3). Some of the other species reported here (Table 3) belong to complexes of closely related, possibly cryptic, species: For the time being, they are identified as the closest name-bearing morpho-species, but may in the future be recognized and described as separate taxa. These taxa are: Enchytraeus buchholzi Vejdovský, 1879 (for which we found two separate species matching the general E. buchholzi morphology, "buchholzi 1" and "2"), Ch cf. diastrophus, Marionina cf. argentea (Michaelsen, 1889), Mesenchytraeus cf. armatus, and Nais communis/variabilis (Piguet, 1906; i.e., we found here morphotype A3 sensu Envall et al. 2012). Moreover, there is still some uncertainty whether the earthworm Bimastos rubidus (Savigny, 1826) should include B. subrubicundus (Eisen, 1874), and B. tenuis (Eisen, 1874), all three of which have been reported from Iceland.

The material that did not produce any molecular data was examined based on morphology and could in most cases only be determined to genus level; such specimens will not be further



Figure 2. Neighbor-Joining tree for the longer (320 bp) 16S barcode of the 46 successfully barcoded specimens and two reference barcodes (Ref.). Scale bar shows 10 % genetic distance under the Jukes-Cantor model. Note that the NJ tree is a poor estimation of actual phylogenetic relationships.



Figure 3. Neighbor-Joining tree for the shorter (70 bp) 16S barcode of all 51 16S-barcoded specimens and two reference barcodes (Ref.). The alignment is 79 bp long and consists of 17 barcodes sequenced using the specific primers of this shorter fragment (see Table 2), and 36 barcodes from the corresponding section in the longer (320 bp) fragment. Scale bar shows 10 % genetic distance under the Jukes-Cantor model. Note that the NJ tree is a poor estimation of actual phylogenetic relationships.

treated here (this is why not all originally sampled springs are shown in Figure 1 and Table 1). However, we did identify a few additional species in our spring material based on morphology alone. One being *Cognettia varisetosa* (Martinsson, Rota & Erséus, 2015), earlier regarded as *C. glandulosa* (previously recorded from Iceland); *C. varisetosa* is thus, at least nominally, a new record for Iceland. We also found specimens of *Pristina foreli* (Piguet, 1907), which represents a genus (*Pristina Ehrenberg*, 1828) never recorded in Iceland before. Finally, we found *Uncinais uncinata* (Ørsted, 1842), a taxon already known from the country (see Table 4).

In total, we identified 23 species, of which at least 8 are new records to Iceland. These identified species were collected from 19 of the 31 sites sampled.

## DISCUSSION

#### Clitellates of the Icelandic springs

The species found in the Icelandic springs are a mixture of Lumbricidae, Enchytraeidae and Naididae. Earthworms (Lumbricidae) are mostly terrestrial, but among our four species found, *Eiseniella tetraedra* is a characteristic inhabitant of running water or wet soils, and common also in caves and springs, in the Western Palaearctic (Sims & Gerard 1985). The other three are terrestrial worms "accidently found in water" (Timm 2009, p. 188).

All Enchytraeidae (13 species) and Naididae (7 species) in the studied springs are known also from continental Europe. This conclusion is largely based on molecular data, as we were able to compare the 16S barcodes of the Icelandic specimens with the corresponding barcodes of <400 species of Enchytraeidae and Naididae from Sweden and Norway (Erséus and Klinth, unpubl.). This enabled us to identify certain (cryptic) forms within the species complexes of some traditional morphospecies (i.e., *Cernosvitoviella minor* s. lat., *Enchytraeus buchholzi* s. lat., *Marionina argentea* s. lat., *Chaetogaster diastrophus* s. lat., *Tubifex tubifex* s. lat.) and one small, yet unidentified *Chaetogaster* species. However, proper binominal names of these cryptic species are not yet established.

*Fridericia dura* (Enchytraeidae) is typically terrestrial (Dózsa-Farkas 2019), but was found outside its normal habitat in this study. The remaining enchytraeids and all naidids are normally restricted to aquatic or semi-aquatic habitats (Timm 2009; Schmelz & Collado 2010; Klinth *et al.* 2017b), and they appear as a somewhat impoverished assemblage of the clitellates typical of streams, rivers, lakes and ponds in other parts of Northern Europe.

Springs are windows into the stygofauna, i.e., stygofaunal species are categorized as those restricted to groundwater (stygobites), those inhabiting both surface and ground waters, or preferring a transition zone of these habitats (stygophiles), and those accidentally or occasionally present in groundwater (stygoxenes). No Icelandic clitellates so far known are stygobites. For instance, there are no records of species of the genera typically containing stygobitic (often endemic) taxa in continental Europe, such as Trichodrilus Claparède, 1862 (Lumbriculidae), Aberrantidrilus Martin, 2015, Aktedrilus Knöllner, 1935, Gianius Erséus, 1992, Protuberodrilus Giani & Martinez-Ansemil, 1979, Rhyacodrilus Bretscher, 1901, and Troglodrilus Juget et al., 2006 (all Naididae). However, five meiobenthic species found in the present study (Cernosvitoviella aggtelekiensis, C. pusilla, C. cf. minor, Marionina cf. argentea and Pristina foreli) are associated with surface waters as well as springs and groundwater in Norway and Sweden, and (when in springs) often in various combinations with each other (Erséus & Klinth, unpubl.). These taxa may be regarded as stygophiles, and their small size may be advantageous in springs, where nutrient levels are often low. Moreover, three other taxa (Lumbricillus arenarius, Marionina sp. and Nais elinguis) are normally associated with marine, intertidal habitats. Lumbricillus arenarius is also known from a spring in Northern Svalbard (Klinth et al. 2017b), and Nais elinguis is well known from both springs and coastal streams (e.g., Timm 2009), but the unidentified, possibly new species of Marionina was earlier collected only in marine habitats in Norway and Sweden (Erséus & Klinth, unpubl.).

Enchytraeus buchholzi s. lat., a species complex generally associated with "not too acidic" soils (Schmelz and Collado, 2010), sometimes occurs in freshwater (Timm 2009). The two genetically distinct forms of E. buchholzi found in our study are common in wet soils, including springs, in mainland Scandinavia (Erséus & Klinth, unpubl.). They thus appear to be more aquatic than other members of the complex. As for *Tubifex tubifex* s. lat., most of the cryptic species studied by us (Erséus & Klinth), including the one from the (Icelandic) Botnar I spring, are occasionally found in springs of other parts of Northern Europe. To conclude, we consider our recorded Lumbricidae spp. (possibly excepting the somewhat "stygophilic" Eiseniella tetraedra), Enchytraeus buchholzi 1 & 2, Henlea perpusilla, Mesenchytraeus cf. armatus, T. cf. tubifex, Cognettia varisetosa, Chaetogaster spp. and Uncinais uncinata as stygoxenes.

#### Clitellata of Iceland, an updated species list

In Table 4, >90 taxa of Clitellata reported from Iceland to date are listed. The exact number of species is not yet known, considering that several taxa are species complexes. The present study has added eight binominal species new for Iceland, plus one unidentified *Marionina* sp. that may be new to science, and the small unidentified *Chaetogaster* sp., which if not a new species is possibly *C. langi. Cernosvitoviella* and *Pristina* are genera that have not been reported from Iceland before.

#### Barcodes and species identification

This study shows that when traditional (COI) barcoding fails due to DNA degradation, at least part of the material may be identified by targeting a shorter gene fragment (i.e., another barcode). The problem is to decide how short a barcode can be and still be species-specific enough for secure species identification. In theory, when degraded DNA is fragmented into ever-smaller pieces, the smaller the target sequence selected the higher the yield of successful sequences, but at the cost of less genetic information for distinguishing species. In our case, the longer of the two 16S fragments (320 bp), produced significantly more sequences than COI (46 compared to 3), and revealed enough genetic variation to separate closely related species (Figure 2). For some of our taxa, however, species separation was based on only one or a few substitutions in the 16S fragment. It is therefore important to note that we refer to these similar 16S sequences as belonging to separate species, on the basis of other genetic information of other individuals of the same species, mainly from the more variable markers ITS (Internal Transcribed Spacer region) and COI (Erséus and Klinth, unpubl.). To be able to use a short gene fragment such as our 320-bp 16S to identify species it is clear that a large library with multiple sequences from all potential species, representing both inter- and intraspecific variability, is required.

Concerning the 70-bp 16S barcode, we surprisingly found that it did not produce more sequences than the 320-bp one, given the degraded DNA. Instead, it produced fewer successful barcodes. A likely explanation for this is sub-optimal binding of the primers, either due to the annealing temperature, or nucleotide variations in the primer-binding site (also indicated by a lack of bands in the post-PCR electrophoresis gel). The primers were originally designed for earthworms (Bienert *et al.* 2012), and in the present study they generally worked better for Lumbricidae than for the other families (Table 3), for which modified primers may be needed. It is possible that this very short 16S partition does not contain enough variation to delimit all closely related species of Clitellata, and yet it proved variable enough to distinguish all the 17 successfully sequenced specimens in our current material from each other (Figure 3).

We had problems with contamination in eight of our specimens; their 16S sequences (320 bp) did not match the species revealed by the morphology of the vouchers. In most cases we could attribute this to cross-contamination between samples, or possibly from the extraction lab. There were also some cases where the resulting sequences were those of human or bacterial DNA, but they were directly excluded from the counts of barcoded worms.

The integration of molecular and morphological data is particularly important in the delimitation of clitellate species (e.g., Martinsson *et al.* 2013; Klinth *et al.* 2017a). However, using DNA barcoding alone as a reliable shortcut to actual species identification has its pros and cons. In theory, clitellate barcoding is near to perfect when all species have been properly delimited. Moreover, it has the advantages of handling all life stages and even extra-organismal DNA (e.g., DNA from mucus left behind by tunnelling earthworms), and it separates cryptic species. On the other hand, this study has shown that problems occur in practice. We studied samples that suffered from Table 4. Updated checklist of clitellate species from Iceland. Previously recorded species from Iceland are presented together with the reference paper.

Species sorted by family	References
Enchytraeidae	
Achaeta unibulba Graefe, Dózsa-Farkas & Christensen 2005	Graefe et al. 2005
Bryodrilus parvus Nurminen, 1970	Nurminen 1973
Buchholzia appendiculata (Buchholz, 1862)	Christensen 1962; Nurminen 1973
Cernosvitoviella aggtelekiensis Dózsa-Farkas, 1970	This study (new record)
Cernosvitoviella cf. minor Dózsa-Farkas, 1990 (one in a species complex)	This study (new record)
Cernosvitoviella pusilla Nurminen, 1973	This study (new record)
Claparedrilus semifuscoides Klinth, Rota & Erséus, 2017 (previously reported as L. semifuscus)	Christensen 1962; Erséus 1976
<i>Cognettia glandulosa</i> (Michaelsen, 1888) previous records could have been <i>C. glandulosa</i> or <i>C. varisetosa</i> (see Martinsson, Rota & Erséus, 2015a)	Christensen 1962; Nurminen 1973
<i>Cognettia sphagnetorum</i> (Vejdovský, 1877) previous records could have been <i>C. chalupskyi</i> , <i>C. chlorophila</i> , <i>C. pseudosphagnetorum</i> or <i>C. sphagnetorum</i> (see Martinsson, Rota & Erséus, 2015b)	Christensen 1962; Nurminen 1973
<i>Cognettia varisetosa</i> (Martinsson, Rota & Erséus, 2015a) (previously a part of <i>C. glandulosa</i> )	This study (new record)
Enchytraeus albidus Henle, 1837	Christensen 1962; Erséus 1976
Enchytraeus buchholzi Vejdovský, 1879 (species complex)	Christensen 1962; Nurminen 1973; two species found in this study
Enchytraeus coronatus Nielsen & Christensen, 1959	Christensen 1962
Enchytraeus minutus Nielsen & Christensen, 1961	Nurminen 1973
Enchytraeus norvegicus Abrahamsen, 1969	Nurminen 1973
Fridericia bisetosa (Levinsen, 1884)	Christensen 1962; Nurminen 1973
Fridericia bulboides Nielsen & Christensen, 1959	Christensen 1962; Nurminen 1973
Fridericia bulbosa (Rosa, 1887)	Christensen 1962; Nurminen 1973
Fridericia callosa (Eisen, 1878)	Christensen 1962
Fridericia dura (Eisen, 1879)	This study (new record)
Fridericia galba (Hoffmeister, 1843)	Christensen 1962; Nurminen 1973
Fridericia leydigi (Vejdovský, 1877)	Nurminen 1973
Friderica maculata Issel, 1905	Christensen 1962
Fridericia perrieri (Vejdovsky)	Christensen 1962
Fridericia ratzeli (Eisen, 1872)	Christensen 1962; Nurminen 1973
Fridericia striata (Levinsen, 1884)	Christensen 1962
Grania postclitellochaeta (Knöllner, 1935)	Rota & Erséus 2003
Henlea glandulifera Nurminen, 1970	Nurminen 1973
Henlea nasuta (Eisen, 1878)	Christensen 1962
Henlea perpusilla Friend, 1911	Christensen 1962; Nurminen 1973; this study
Henlea ventriculosa (Udekem, 1854)	Christensen 1962; Nurminen 1973
Lumbricillus arenarius (Michaelsen, 1889)	Christensen 1962, this study
Lumbricillus lineatus (Müller, 1774)	Christensen 1962; Erséus 1976
Lumbricillus macrothecatus Erséus, 1976	Erséus 1976
Lumbricillus pagenstecheri (Ratzel, 1869)	Christensen 1962; Erséus 1976
Lumbricillus pumilio Stephenson, 1932	Erséus 1976
Lumbricillus reynoldsoni Backlund, 1948	Christensen 1962
Lumbricillus rivalis Levinsen, 1883 emend, Ditlevsen, 1904	Christensen 1962

Table 4. Continued.

Species sorted by family	References
Lumbricillus scoticus Elmhirst & Stephenson, 1926	Christensen 1962: Erséus 1976
Lumbricillus viridis (Stephenson, 1911)	Christensen 1962: Erséus 1976
Marionina argentea (Michaelsen, 1889) (species complex)	Nurminen 1973; one species found in this study
Marionina communis Nielsen & Christensen, 1959	Christensen 1962; Nurminen 1973
Marionina spicula (Leuckart, 1847)	Christensen 1962; Erséus 1976
Marionina sp.	This study (unidentified/new species?)
Mesenchytraeus cf. armatus (Levinsen, 1884) (one in a species complex)	This study (new record)
Mesenchytraeus flavus (Levinsen, 1884)	Christensen 1962; Nurminen 1973
Hirudinea	
Callobdella nodulifera Malm, 1863	Bruun 1938a
Glossiphonia complanata (Linnaeus, 1758)	Bruun 1938b; Lindegaard 1979
Helobdella stagnalis (Linnaeus, 1758)	Bruun 1938b; Lindegaard 1979
Heptacyclus scorpii (Malm, 1863)	Bruun 1938a
Johanssonia arctica (Johansson, 1899)	Perdiguero-Alonso et al. 2008
Oceanobdella microstoma (Johansson, 1896)	Bruun 1938a
Oxytonostoma typica Malm, 1863	Bruun 1938a
Platybdella anarrhichae (Diesing, 1859)	Bruun 1938a
Pontobdella muricata (Linnaeus, 1758)	Bruun 1938a
Theromyzon garjaewi (Livanow, 1903) valid species?	Bruun 1938b
Theromyzon maculosum (Rathke, 1862) valid species?	Fjeldså & Raddum 1973
Theromyzon tessulatum (Müller, 1774)	Bruun 1938b; Lindegaard 1979
Lumbricidae	
Aporrectodea caliginosa (Savigny, 1826)	Backlund 1949; Lindroth et al. 1973; this study
Aporrectodae rosea (Savigny, 1826)	Backlund 1949; Lindroth et al. 1973
Bimastos rubidus s. lat. (reported as Dendrodrilus rubidus (Savigny, 1826), Dendrodrilus subrubicundus (Eisen, 1874), and Dendrodrilus tenuis (Eisen, 1874))	Backlund 1949; Lindroth et al. 1973; this study
Dendrobaena octaedra (Savigny, 1826)	Backlund 1949; Lindroth et al. 1973; this study
Eisenia foetida (Savigny, 1826)	Backlund 1949
Eiseniella tetraedra (Savigny, 1826)	Backlund 1949; Lindroth <i>et al.</i> 1973; Lindegaard 1979; this study
Lumbricus castaneus (Savigny, 1826)	Backlund 1949; Lindroth et al. 1973
Lumbricus rubellus Hoffmeister, 1843	Backlund 1949; Lindroth et al. 1973
Lumbricus terrestris Linnaeus, 1758	Backlund 1949; Lindroth et al. 1973
Octolasium cyaneum (Savigny, 1826)	Backlund 1949
Lumbriculidae	
Lumbriculus variegatus (Müller, 1774)	Hrabě 1952; Lindegaard 1979
Stylodrilus heringianus Claparède, 1862	Hrabě 1952; Lindegaard 1979
Naididae	
Aktedrilus arcticus (Erséus, 1978)	Erséus 1978
Aulodrilus limnobius Bretscher, 1899	Hrabě 1952
Aulodrilus pluriseta (Piguet, 1906)	Hrabě 1952

#### Table 4. Continued.

Species sorted by family	References
Baltidrilus costatus (Claparède, 1863)	Erséus 1976
Chaetogaster diaphanus (Gruithuisen, 1828)	Hrabě 1952; Lindegaard 1979
Chaetogaster cf. diastrophus (Gruithuisen, 1828) (one in a species complex)	This study (new record)
Chaetogaster langi Bretscher, 1896	Hrabě 1952; Lindegaard 1979
Chaetogaster sp. (possibly Ch. langi?)	This study (Ch. langi found again?)
Christerius litoralis (Erséus, 1976)	Erséus 1976
Clitellio arenarius (Müller, 1776)	Hrabě 1952; Brinkhurst 1963; Erséus 1976
Nais communis Piguet, 1906	Hrabě 1952; Lindegaard 1979
Nais communis/variabilis species complex, morphotype A3 (see Envall et al. 2012)	This study
Nais elinguis Müller, 1773	Hrabě 1952; Erséus 1976; Lindegaard 1979; this study
Nais variabilis Piguet, 1906	Hrabě 1952
Paranais litoralis (Müller, 1784)	Erséus 1976
Pristina foreli (Piguet, 1907)	This study (new record)
Specaria josinae (Vejdovský, 1883)	Hrabě 1952
Spirosperma ferox (Eisen, 1879)	Hrabě 1952; Lindegaard 1979
Stylaria lacustris (Linnaeus, 1767)	Hrabě 1952
Thalassodrilus firmus (Erséus, 1979)	Erséus 1979
Tubifex tubifex (Müller, 1774) (species complex)	Hrabě 1952; Lindegaard 1979; one species found in this study
Tubificoides benedii (Udekem, 1855)	Brinkhurst 1963; Erséus 1976
Tubificoides kozloffi Baker, 1983	Helgason & Erséus 1987
Uncinais uncinata (Ørsted, 1842)	Hrabě 1952; Lindegaard 1979; this study

DNA deterioration, which considerably reduced the number of identified specimens. We also found evidence of DNA contamination, which would have led to the wrong conclusions, had we not compared the morphology of a specimen with the barcode sequence obtained. The conclusion is that any samples to be used for DNA analysis must be handled properly, e.g., kept at low temperature, conserved in high concentration of ethanol or DNA preserving buffers, and minimizing storage time. By doing so, the risk of low sequencing success as well as obtaining erroneous identifications due to contamination will be considerably reduced.

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