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Diversity and bioactivity of cultured aquatic fungi from the High Arctic region

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Abstract This study assessed the diversity and α -glycosidase inhibitory activity of cultured fungi isolated from four aquatic environments (stream, pond, glacial ice, and estuary) in the Ny-Ålesund region (Svalbard, Norway, High Arctic). A total of 134 fungal isolates were obtained from 13 water samples. Based on morphological characteristics and sequence analyses of the nuclear ribosomal DNA internal transcribed spacer region, these fungal isolates were identified as belonging to 47 species, with 26 belonging to the Ascomycota, 20 to the Basidiomycota, and one to the Zygomycota. The most frequently detected fungal species were *Vishniacozyma* sp. 2, *Cadophora* sp. 2, *Phenoliferia* sp. 1, *Dioszegia* sp. 2, and *Mortierella* sp.; these species occurred in 10, eight, seven, six, and five of the samples, respectively. Among the 134 fungal isolates, 17 isolates of 15 species displayed high α -glycosidase inhibitory activity in culture. The results suggest that diverse and distinct populations of cultured fungi are present in Arctic aquatic environments, and they include taxa that are potential sources of bioactive molecules that may be used as prototype drugs for medicinal proposals.

Keywords aquatic habitats, bioactivity, diversity, environmental mycology, High Arctic

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1 Introduction

Average Arctic temperatures have increased at almost twice the global average rate over the past 100 years (Solomon et al., 2007). Some resulting environmental changes are of great concern, such as retreating glaciers and the shrinkage and disappearance of lakes and ponds (Overpeck et al., 1997; Smith et al., 2005; Jepsen et al., 2013). Arctic aquatic environments (e.g., glaciers, lakes, and ponds) are unique habitats that may harbor many undiscovered and previously discovered species of microorganisms (e.g., fungi, algae, and bacteria), which may possess unique physiological characteristics. In particular, Arctic fungi have been described as a promising source of bioactive

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natural compounds (Frisvad, 2008; Chen et al., 2013). Hence, there is an urgent need to improve baseline information regarding the fungal diversity and bioactive substances in these habitats.

Currently, approximately 3000 fungal species have been reported from aquatic environments worldwide (Shearer et al., 2007), while only a few studies have been conducted on cultured aquatic fungal species in polar regions, including the Arctic (Kovayasi et al., 1967; Booth et al., 1971; Butinar et al., 2007; Gulis et al., 2012) and Antarctica (Ellis-Evans, 1985, 1996; Goncalves et al., 2012). Recently, the diversity and composition of aquatic fungal communities in the Arctic was revealed by high-throughput sequencing (Zhang et al., 2016), which can enhance the characterization of fungal diversity compared with culture methods. Thus, obtaining cultured fungal isolates may play a key role in the search for bioactive compounds. To the best of our knowledge, the bioactivity of cultured aquatic fungi in the Arctic has not been investigated previously.

Previous studies have proven that different bioactive fungal isolates found in polar regions represent a unique source of newly discovered prototype molecules that can be used in drug discovery studies (Santiago et al., 2012; Godinho et al., 2013). Additionally, fungi have been proven to be a source of α -glycosidase inhibitors (Kim et al., 2004; Kang et al., 2013; Kumar et al., 2014; Rivera-Chávez et al., 2013, 2015), which reduce the impact of carbohydrates on blood sugar and are regarded as effective anti-diabetic drugs used for the treatment of diabetes mellitus type 2 (adult-onset diabetes) (van de Laar et al., 2005), a very serious metabolic disorder that is characterized by high blood sugar. Therefore, the aims of this study were to clarify the diversity of cultured aquatic fungi in aquatic habitats near Ny-Ålesund and to assess their α-glycosidase inhibitory activity.

2 Materials and methods

2.1 Study sites and sample collection

The study area is located in the Ny-Ålesund region (78°55'N, 11°56'E), which is entirely within the High Arctic zone and has a high number of glaciers, lakes, ponds, and glacial streams. Sampling was performed in July 2013. Water samples were collected directly into Labplas (Labplas Inc., TWIRL'EM sterile sampling bags Sainte-Julie, QC, Canada). Then, the microbial samples were collected by filtering 1 L of the water samples. Microbial biomass was trapped onto 47-mm diameter, 0.2-µm pore size membrane filters (Pall Corporation, Port Washington, NY, USA). The membrane filters were placed in sterile centrifuge tubes at -20°C at the Chinese Yellow River Station and then flown to our laboratory in Beijing, China. The locations and environmental types of the 13 water samples are shown in Figure 1 and Table 1.



Figure 1 Detailed view of the four aquatic environments. a, stream; b, pond; c, glacial ice; d, estuary.

2.2 Isolation of aquatic fungi

The membrane filters were placed on solid medium in 9-cm diameter Petri dishes. The solid medium contained potato dextrose agar (PDA), tetracycline (50 mg \cdot L⁻¹), and

streptomycin sulfate (50 mg·L⁻¹). The Petri dishes were sealed by parafilm, incubated for 1 month at 12°C, and examined periodically. When fungal colonies developed, they were transferred to PDA slants. Subcultures were also incubated on PDA.

	environments	In the Ny-Ale	sund region
Code	Coordinates	Altitude/m	Environment type
R1	78°55′18.25″N; 11°56′01.42″E	16	Stream
R2	78°54′46.23″N; 11°50′39.70″E	90	Stream
R3	78°57′47.21″N; 11°35′09.67″ E	25	Stream
R4	78°54′42.97″N; 11°58′30.76″E	37	Stream
R6	78°53′53.08″N; 12°09′36.08″E	43	Stream
R7	78°58′05.39″N; 11°34′15.17″E	9	Stream
L1	78°55′06.39″N; 11°56′36.11″E	18	Pond
L2	78°57′53.95″N; 11°35′05.65″E	18	Pond
L3	78°57′59.59″N; 12°04′06.91″E	45	Pond
S 1	78°55′41.53″N; 11°54′23.05″E	0	Estuary
S2	78°54′14.35″N; 12°09′29.32″E	0	Estuary
X1	78°53′09.78″N; 12°10′31.12″E	171	Glacial ice
X2	78°54′00.48″N; 12°09′31.96″E	29	Glacial ice

 Table 1
 Location and type of the sampled aquatic environments in the Ny-Ålesund region

2.3 DNA extraction, amplification, and sequencing

Genomic DNA was extracted using a modified cetyltrimethylammonium bromide (CTAB) method (Cubero et al., 1999). The internal transcribed spacer (ITS) region between the small-subunit and large-subunit ribosomal genes DNA was polymerase chain reaction (PCR)-amplified with the primers ITS1F and ITS4 (White et al., 1990). The ITS region was amplified using the following PCR procedure: 95°C for 3 min, followed by 37 cycles of 94°C for 30 s, 52°C for 30 s, and 72°C for 30 s, followed by a final extension at 72°C for 10 min. The PCR products were purified and sequenced with the same primers by Sangon Biotech Co., Ltd. (Beijing, China). The ITS sequence data in this study were deposited in GenBank under the accession nos. KR698797 to KR698930.

2.4 Identification and phylogeny

The isolated fungi were identified based on their phylogenetic position and sequence similarity. Sequence data were used to investigate phylogenetic relationships using Molecular Evolutionary Genetics Analysis (MEGA) software v. 6.0. Phylogenetic trees were constructed using the neighbor-joining algorithm with bootstrap values calculated from 1000 replicate runs. A maximum composite likelihood model was used to estimate evolutionary distance. For sequence similarity determination, nucleotide Basic Local Alignment Search Tool (BLASTn) searches were performed in the GenBank databases to find the most closely related fungal species. In addition, the identification of all the isolates was confirmed by their phenotypic characteristics, including colony aspect (texture and color) and growth rate.

2.5 Statistical analyses

To evaluate the similarity of the cultured fungal species from the four aquatic environments, a Venn diagram of shared and unique fungal species was made using Venny 2.0 (http://bioinfogp.cnb.csic.es/tools/venny/). The fungal diversity in the four aquatic environments was evaluated using the Shannon's diversity index (H'). Furthermore, a heatmap analysis was performed to visualize all the detected fungal species and to compare their abundances among the four aquatic habitats using R 3.1.1 statistical software.

2.6 Fungal cultivation and preparation of extracts for α-glycosidase activity assays

The fungal isolates were inoculated into 25 mL of liquid medium (comprising 0.5% soymeal, 1.0% corn steep liquor, 2.0% glucose, 1.0% dextrin, 0.5% of peptone, and 0.2% of $(NH_4)_2SO_4$) in 100-mL flasks. The flasks were incubated at 16°C for 15 d, and then the cultured materials and fermentation liquor from each flask were respectively transferred to 50-mL vials containing an equal volume of acetone. After a 24-h incubation at room temperature, the organic phase was decanted, and the solvent was removed using a vacuum centrifuge at 35°C. An aliquot of each dried extract was dissolved in dimethyl sulfoxide to prepare a 100 mg·mL⁻¹ stock solution, which was stored at 4°C.

2.7 Assay for α-glycosidase inhibitory activity

The α -glycosidase inhibitory activity of the fungal isolates was determined according to the method described by Huang et al. (2004). α -glycosidase was extracted from the upper duodenum of female Kunning mice (22.0-25.0 g, 12 weeks old, specificpathogen-free) (National Institutes for Food and Drug Control, Beijing, Certificate No. SCXK jing 2009-0017), and 4-nitrobenzene-α-D-glucopyranoside was used as the substrate to detect a-glycosidase activity. All animal procedures in this study were approved by the Animal Ethical Experimentation Committee of Beijing Union University according to the requirements of the National Act on the Use of Experimental Animals (China). The screening was based on measuring the absorbance of p-nitrophenol at 405 nm, which is generated by the reaction of 4-nitrobenzene- α -D-glucopyranoside and α -glycosidase. Acarbose, which is an oral α -glycosidase inhibitor, was used as a positive control. The formula used to calculate the inhibition rate is:

 $[(X-Z)/(Y-Z)] \times 100\%$,

where X is the absorbance value in the presence of the

sample; Y is the absorbance value in the presence of acarbose (control); and Z is the absorbance value in the absence of the sample and acarbose (blank). All assays were performed in triplicate.

3 Results

3.1 Isolation and identification of aquatic fungi

A total of 134 fungal isolates were obtained from the 13

water samples (Table S1). All the isolates were identified to the species level via a phylogenetic analysis (Figure 2) and by their sequence identity (Table 2). For example, fungal isolate I14F-01251 showed the highest sequence similarity (99%) with *Cadophora luteo-olivacea* (NR111149), and it clustered with *Cadophora luteo-olivacea* in one group with a high bootstrap value (100) in the phylogenetic tree (Figure 2); therefore, isolate I14F-01251 was identified as a *Cadophora* species.

 Table 2
 Molecular identification of 47 fungal species isolated from the aquatic environments in Ny-Ålesund: maximum identity was determined using BLASTn searches of the ITS region

No.	Representative	GenBank	Close related species	Number of bp	Identification
1	Isolate code	accession number	(GenBank accession number)	(Maximum identity)	
1	114F-01218	KR698814	Cadophora melinii (JN689950)	462/484 (95%)	Cadophora sp. 1
2	114F-01251	KR698847	Cadophora luteo-olivacea (NR111149)	540/543 (99%)	Cadophora sp. 2
3	I14F-01232	KR698828	Cadophora viticola (HQ661097)	503/506 (99%)	Cadophora sp. 3
4	I14F-01273	KR698869	Cystobasidium laryngis (KY103134)	530/534 (99%)	Cystobasidium sp. 1
5	I14F-01313	KR698909	Cystobasidium psychroaquaticum (KY103148)	533/535 (99%)	Cystobasidium sp. 2
6	I14F-01216	KR698812	Dioszegia rishirensis (KY103358)	425/432 (98%)	Dioszegia sp. 1
7	I14F-01299	KR698895	Dioszegia fristingensis (NR136970)	420/434 (97%)	Dioszegia sp. 2
8	I14F-01324	KR698920	Dioszegia aurantiaca (KY103346)	422/435 (97%)	Dioszegia sp. 3
9	I14F-01208	KR698804	Glaciozyma martinii (NR132821)	556/571 (99%)	Glaciozyma sp. 1
10	I14F-01331	KR698927	Glaciozyma watsonii (KY103477)	593/595 (99%)	Glaciozyma sp. 2
11	I14F-01284	KR698880	Glarea lozoyensis (NR137138)	432/447 (97%)	Glarea sp.
12	I14F-01322	KR698918	Gremmeniella balsamea (NR137589)	434/470 (92%)	Godroniaceae sp.
13	I14F-01263	KR698859	Goffeauzyma gilvescens (NR073228)	588/588 (100%)	Goffeauzyma sp.
14	I14F-01206	KR698802	Glarea lozoyensis (NR137138)	406/451 (90%)	Helotiales sp. 1
15	I14F-01242	KR698838	Coleophoma paracylindrospora (KU728491)	416/483 (86%)	Helotiales sp. 2
16	I14F-01249	KR698845	Hymenoscyphus ohakune (NR137109)	450/514 (88%)	Helotiales sp. 3
17	I14F-01303	KR698899	Glarea lozoyensis (NR137138)	418/460 (91%)	Helotiales sp. 4
18	I14F-01332	KR698928	Variosporium delicatum (KF730848)	450/478 (94%)	Helotiales sp. 5
19	I14F-01269	KR698865	Holtermanniella festucosa (KY102693)	516/517 (99%)	Holtermanniella sp.
20	I14F-01288	KR698884	Acremonium persicinum (NR131260)	472/538 (88%)	Hypocreales sp.
21	I14F-01217	KR698813	Keissleriella trichophoricola (KJ869113)	459/466 (98%)	Keissleriella sp.
22	I14F-01257	KR698853	Mycosymbioces mycenaphila (NR137807)	433/481 (90%)	Leotiomycetes sp.
23	I14F-01226	KR698822	Leptosphaeria veronicae (JF740255)	452/461 (98%)	Leptosphaeria sp. 1
24	I14F-01307	KR698903	Leptosphaeria sclerotioides (JF740193)	504/523 (99%)	Leptosphaeria sp. 2
25	I14F-01281	KR698877	Leucosporidiella fragaria (NR073287)	567/579 (98%)	Leucosporidiella sp. 1
26	I14F-01283	KR698879	Leucosporidiella muscorum (NR073286)	536/552 (97%)	<i>Leucosporidiella</i> sp. 2
27	I14F-01256	KR698852	Mortierella antarctica (NR111580)	588/593 (99%)	<i>Mortierella</i> sp.
28	I14F-01228	KR698824	Mrakia frigida (NR111044)	585/591 (99%)	Mrakia sp. 1
29	I14F-01259	KR698855	Mrakia gelida (KY104295)	556/557 (99%)	<i>Mrakia</i> sp. 2
30	I14F-01289	KR698885	Mrakia fridida (NR111044)	589/615 (96%)	Mrakia sp. 3
31	I14F-01312	KR698908	<i>Mycoarthris cf. coralline</i> (KC485428)	389/396 (98%)	<i>Mycoarthris</i> sp.
32	I14F-01219	KR698815	Paraleptosphaeria macrospora (JF740238)	479/484 (99%)	Paraleptosphaeria sp.
33	I14F-01261	KR698857	Phenoliferia glacialis (KY104503)	553/556 (99%)	Phenoliferia sp. 1
34	I14F-01287	KR698883	Phenoliferia psychrophenolica (KY104504)	538/562 (96%)	Phenoliferia sp. 2
35	I14F-01265	KR698861	Phoma boeremae (NR135962)	475/478 (99%)	Phoma sp.

					Continued table
No.	Representative isolate code	GenBank accession number	Close related species (GenBank accession number)	Number of bp (Maximum identity)	Identification
36	I14F-01238	KR698834	Darksidea delta (NR137075)	410/475 (86%)	Pleosporales sp.
37	I14F-01227	KR698823	Pseudeurotium bakeri (NR145345)	494/499 (99%)	Pseudeurotium sp.
38	I14F-01233	KR698829	Pseudogymnoascus destructans (NR111838)	492/499 (99%)	Pseudogymnoascus sp.
39	I14F-01201	KR698797	Taphrina carnea (AF492083)	566/569 (99%)	Taphrina sp.
40	I14F-01220	KR698816	Tetracladium sp. (AJ890435)	494/514 (96%)	Tetracladium sp.
41	I14F-01244	KR698840	Tolypocladium cylindrosporum (AJ303055)	481/486 (99%)	Tolypocladium sp.
42	I14F-01280	KR698876	Genolevuria amylolytica (NR137810)	369/434 (85%)	Tremellales sp.
43	I14F-01234	KR698830	Varicosporium delicatum (DQ202516)	460/486 (95%)	Varicosporium sp.
44	I14F-01221	KR698817	Venturia tremulae (EU035438)	508/521(98%)	Venturia sp.
45	I14F-01237	KR698833	Vishniacozyma tephrensis (NR144812)	472/473 (99%)	Vishniacozyma sp. 1
46	I14F-01266	KR698862	Vishniacozyma victoriae (NR073260)	465/466 (99%)	Vishniacozyma sp. 2
47	I14F-01311	KR698907	Vishniacozyma carnescens (KY105817)	477/480 (99%)	Vishniacozyma sp. 3

Altogether, 134 fungal isolates were identified belonging to 47 species. These isolated strains belong to 27 genera from the phyla Ascomycota (17 genera), Basidiomycota (nine genera), and Zygomycota (one genus). The Ascomycota included five known orders: Helotiales (13 species), Hypocreales (two species), Pleosporales (six species), Taphrinales (one species), and Venturiales (one species). The Basidiomycota included four known orders: Cystobasidiales (two species), Cystofilobasidiales (three species), Sporidiobolales (four species), and Tremellales (nine species). The Zygomycota included only one known order: Mortierellales (one species). In addition, the most frequently detected species in the 13 water samples were Vishniacozyma sp. 2, Cadophora sp. 2, Phenoliferia sp. 1, Dioszegia sp. 2, and Mortierella sp., which were isolated from ten, eight, seven, six, and five of the samples. respectively (Table S1).

BLASTn searches in GenBank also showed that matching sequences with high similarity (\geq 97%) were derived from fungi found in various habitats (Table S1), including plant roots, wood, soil, glacial sediment, glacial water, snow, streams, cryoconite holes, and lake sediment, within the Arctic (e.g., Svalbard), Antarctica, and alpine regions (e.g., Tibetan Plateau).

3.2 Dissimilarity of fungal diversity among the different aquatic habitats

Twenty-eight, 18, 12, and 16 fungal species were isolated from the stream, pond, glacial ice, and estuary, respectively (Table S2). The Shannon diversity indices were highest for the stream habitat (4.34), lowest for the ice habitat (3.22), and intermediate for the pond (3.92) and estuary (3.93) (Table S2). A Venn diagram demonstrated that the fungal species differed among the four aquatic habitats. The number of habitat-specific species ranged from three (the estuary) to 13 (the stream) (Figure 3a). Only three species were shared by the four aquatic habitats. The number of shared species among the different aquatic habitats was low; eight, five, and five species were shared between the stream and the pond, the pond and the glacial ice, and the estuary and the glacial ice, respectively.

To better understand the dissimilarity of fungal species among the different aquatic habitats, a heatmap analysis was also performed (Figure 3b). Only three fungal species coexisted in all four aquatic habitats, and these species were Dioszegia sp. 2, Mrakia sp. 2, and Vishniacozyma sp. 2. Thirteen species (e.g., Venturia sp. and Taphrina sp.) were found only in the stream. Nine species (e.g., Hypocreales sp. and Glarea sp.) were found only in the pond, and four species (i.e., Helotiales sp. 4, Mycoarthris sp., Vishniacozyma sp. 3, and Mrakia sp. 3) were found only in the glacial ice. Three species were found only in the estuary, including Helotiales sp. 5, Dioszegia sp. 3, and Godroniaceae sp. (Figure 3b). The dominant fungal species were Cadophora sp. 2 in the stream, Vishniacozyma sp. 2 in the pond, Phenoliferia sp. 1 in the glacial ice, and Dioszegia sp. 2, Glaciozyma sp. 2, and Vishniacozyma sp. 2 in the estuary (Figure 3b and Table S2).

3.3 Assay of α-glycosidase inhibitory activity

All the 134 aquatic fungal isolates were grown using liquid medium to obtain crude cell-free extracts, which were then screened for α -glycosidase inhibitory activity. Seventeen extracts of the fungal isolates (12.7%), which belonged to 15 species, displayed α -glycosidase inhibitory activity (Table 3). Among these 17 fungal isolates, Helotiales sp. 4 (114F-01296), *Venturia* sp. (I14F-01221), *Dioszegia* sp. 2 (I14F-01229), *Mortierella* sp. (I14F-01243), and *Mrakia* sp. 2 (I14F-01250) had high levels of α -glycosidase inhibitory activity.

a ..

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	94 Hebitales sp. 1 114F-01236 (KR69880) 99 Glarea sp. 114F-01238 (KR69880) 99 Glarea sp. 114F-01238 (KR69880) 90 Glarea sp. 114F-01238 (KR69880) 91 Glarea sp. 114F-01218 (KR69880) 91 Glarea sp. 114F-01218 (KR69880) 91 Glarea sp. 114F-01218 (KR698820) 100 Cadophora sp. 3 [114F-01232 (KR698820) 100 Cadophora sp. 3 [114F-01232 (KR698820) 100 Cadophora sp. 114F-01232 (KR698820) 100 Fetadalum sp. 114F-01232 (KR698823) 100 Pseudogymnoacus sp. 114F-01233 (KR698823) 100 Pseudogymnoacus destructarus (NR11383) 99 Miccosptime sp. 114F-01232 (KR698823) 100 Pseudogymnoacus destructarus (NR11383) 100 A lectionize sp. 114F-01232 (KR698820) 100 A cortonice mericine with Kr115300 100 A cortonice sp. 114F-01228 (KR698820)
	100 Mirakia gp.el/104-01239 (KR698085) 100 Mirakia gelida (KY104295) ▲ Mirakia gp.3 114F-01289 (KR698885)
L	92 92 100 Goffeauzyma sp. 114F-01263 (KR698859) 100 A Vishniacozyma sp. 1 14F-01237 (KR698833)
	Visiniacozyma sp. 3 114F-01311 (KR698907)
	95 Vishniacozyma camescens (KY105817) 91 ▲ Vishniacozyma sp. 2 14F-01266 (KR698862) 97 Vishniacozyma victoriae (NR073260)
	▲ Holtermanniella sp. 114F-01269 (KR698865) 100 ¹ Holtermanniella festucosa (KY102693)
	100 ¹ Cystobasidium psychroaquaticum (FN868153) 100 ▲ Glaciozyma sp. 1 14F-01208 (KR69804) 100 ⊂ Glaciozyma sp.tini (KR133821)
	64 Glaciozyma sp. 2 114F-01331 (KR698927) 100 ¹ Glaciozyma watsonii (KY103477)
	Phenoliferia psychrophila (KY104504) 877 Phenoliferia glacialis (KY104503) 100 ▲ Phenoliferia sp. 1 14F-01261 (KR698857)
	Phenoliferia sp. 2 114F-01287 (KR698833) 100 ← Leucosporidiella sp. 1 14F-01281 (KR698877) Leucosporidiella sp. 1 14F-01281 (KR698877) 100 ← Leucosporidiella sp. 2 114F-01283 (KR698879)

Figure 2 Phylogenetic tree of the 47 cultured fungal species (based on a representative isolate of each species) and other related fungal species based on the ITS (ITS1-5.8S-ITS2) region sequences of ribosomal DNA. The tree was constructed with the neighbor-joining method. Bootstrap support values are indicated for major nodes having values ≥ 60 .

Table 3The α -glycosidase inhibitory activity of 17 fungal
isolates from the aquatic environments in the
Ny-Ålesund region

Ny-Alesun	u region	
Fungal species	Isolate no.	α-glycosidase inhibitory activity/% [#]
Cadophora sp. 1	I14F-01218	64.1 ± 0.02
Dioszegia sp. 2	I14F-01229	66.1 ± 0.01
Dioszegia sp. 2	I14F-01253	65.0 ± 0.01
Dioszegia sp. 2	I14F-01264	65.6 ± 0.02
Helotiales sp. 2	I14F-01242	64.4 ± 0.01
Helotiales sp. 3	I14F-01249	64.8 ± 0.01
Helotiales sp. 4	I14F-01296	67.1 ± 0.02
Hypocreales sp.	I14F-01288	63.1 ± 0.02
Keissleriella sp.	I14F-01217	57.4 ± 0.12
Leotiomycetes sp.	I14F-01257	62.8 ± 0.02
Leptosphaeria sp. 1	I14F-01226	65.5 ± 0.01
Tetracladium sp.	I14F-01220	63.7 ± 0.01
Mortierella sp.	I14F-01243	66.1 ± 0.01
Mrakia sp. 2	I14F-01250	66.1 ± 0.01
Phenoliferia sp. 1	I14F-01230	65.8 ± 0.01
Phenoliferia sp. 2	I14F-01287	63.7 ± 0.04
Venturia sp.	I14F-01221	66.1 ± 0.01

Notes: # Values are means of three replicate \pm S.D.

4 Discussion

Among the 27 fungal genera detected in this study, many have been reported previously from aquatic habitats in polar regions, including Vishniacozyma and Phenoliferia in the Arctic (Butinar et al., 2007), and Cadophora, Vishniacozyma, Mortierella, Phoma, Pseudeurotium, and Phenoliferia in Antarctica (Ellis-Evans, 1985, 1996; Goncalves et al., 2012). Some of the fungal genera were found previously in non-aquatic habitats in the Arctic. For example, the genera Acremonium, Morteriella, Tolypocladium, and Varicosporium were also isolated from Arctic soil (Singh et al., 2012; Hafizah et al., 2013). In previous studies, a Cadophora sp. was isolated from a plant in Antarctica (Rosa et al., 2010), and a Tolypocladium sp. was isolated from Arctic soil in Svalbard (Hafizah et al., 2013), while a Phoma sp. was detected in Arctic plant roots (Botnen et al., 2013). Additionally, several fungal species, including species of Cadophora, Vishniacozyma, Mrakia, Mortierella, and Phenoliferia (as shown in Table S1), can even be found in Antarctica. These results suggest that some of the fungal species found in this study may originate from non-aquatic habitats in the Arctic ecosystem. It also suggests that the presence of specific psychrophilic and psychrotolerant fungi varies across different habitats within polar areas. However, the cultured fungal isolates obtained in this study represent a small portion of all the fungal species present in Arctic aquatic environments. In a previous study, we assessed the



Figure 3 Venn diagram displaying the number of fungal species common among the different aquatic habitats of the Ny-Ålesund region (**a**); heatmap diagram showing the distribution and abundance of the 47 fungal species among the four aquatic habitats (**b**).

diversity of aquatic fungal communities in the same region using high-throughput pyrosequencing, and we detected 641 operational taxonomic units, which belonged to 90 known genera (Zhang et al., 2016).

The fungal communities from the different aquatic habitats (stream, pond, glacial ice, and estuary) in this High Arctic region differed in their composition, suggesting that aquatic fungal communities are selective in their habitat preference. Additionally, the highest level of fungal diversity was detected in the stream habitat, while the lowest level was detected in the glacial ice habitat, and intermediate levels were detected in the pond and estuary habitats. These data are consistent with previous findings regarding the variability of aquatic fungal communities from different habitats (stream, pond, glacial ice, and estuary), as revealed using high-throughput sequencing (Zhang et al., 2016). This is likely a result of the differences in the physicochemical properties of the respective habitats in this region, including temperature, salinity, and conductivity (Zhang et al., 2016).

The ability of Arctic and Antarctic fungi to survive in extreme environmental conditions suggests that they may possess unique metabolic pathways that allow them to generate specific or novel compounds with potential medicinal activity that could be used to develop new drugs (Santiago et al., 2012). Many fungal species isolated from Antarctic habitats have been shown to have potential medicinal activity and bioactive compounds (Santiago et al., 2012; Godinho et al., 2013). However, until now, the potential medicinal activity of Arctic fungi and their ability to produce bioactive compounds have been poorly understood.

Our results showed that the cell-free extracts of 15 fungal species have α -glycosidase inhibitory activity. This is the first time that Arctic fungi have been shown to have α -glycosidase inhibitory activity, suggesting that they produce bioactive secondary metabolites that are responsible for the observed α -glycosidase inhibitory activity. Some previous studies reported that only a few fungal species isolated from non-Arctic habitats are a source of α-glycosidase inhibitors (van de Laar F A et al., 2005; Kang et al., 2013; Kumar et al., 2014; Rivera-Chávez et al., 2013, 2015). Kang et al. (2013) found that extracellular α -glucosidase inhibitory activities were very low (2.1%-3.1%), or even undetectable, for 34 fungal supernatants, whereas higher intracellular α -glucosidase inhibitory activities were detected (20.6%-48.3%) in cell-free extracts. In particular, a new α -glucosidase inhibitory peptide was purified and identified from a cell-free extract of Aspergillus orvzae. Kim and Nho (2004) isolated an α -glucosidase inhibitor (SKG-3) from the fruiting bodies of Ganoderma lucidum, and they showed that SKG-3 is a highly specific and effective reversible inhibitor of α -glucosidase. The identification of 15 fungal species with α -glycosidase inhibitory activity in the present study substantially increases the potential number of bioactive secondary metabolites that might be used to develop new drugs.

In summary, this study showed that aquatic environments in the Arctic shelter rich, diverse, and complex fungal communities. Currently, many aspects of the taxonomy, ecology, and potential biotechnological applications of aquatic fungi from the Arctic remain poorly documented. Our results reinforce the need for further studies of aquatic fungal communities in the Arctic, as well as their phylogenetic relationships with species that occur in other cold regions (e.g., Antarctica or the Tibetan Plateau). In addition, we found that many aquatic fungal isolates had α -glycosidase inhibitory activity, which may provide new insights into the potential biotechnological applications of such fungi in Arctic aquatic environments

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Sample code	Isolate code	Identification	GenBank accession number	Closest related species (GenBank accession number)	Number of bp (Maximum identify)	Reported habitat
R1	I14F- 01201	<i>Taphrina</i> sp.	KR698797	Taphrina carnea (AF492083)	566/569 (99%)	Unreported
	I14F- 01202	Cadophora sp. 2	KR698798	<i>Cadophora cf. luteo-olivacea</i> (KC180677)	497/504 (99%)	Hair root of plant in Argentina
	I14F- 01203	Cadophora sp. 2	KR698799	<i>Cadophora cf. luteo-olivacea</i> (KC180677)	492/501 (98%)	Hair root of plant in Argentina
	I14F- 01204	Cadophora sp. 2	KR698800	Cadophora luteo-olivacea (KF053558)	504/504 (100%)	Wood in Antarctica
	I14F- 01205	Cadophora sp. 2	KR698801	Cadophora cf. luteo-olivacea (KC180677)	492/501 (98%)	Hair root of plant in Argentina
	I14F- 01206	Helotiales sp. 1	KR698802	Helotiales sp. (JQ715700)	471/476(99%)	Soil in Antarctica
	I14F- 01207	Cadophora sp. 2	KR698803	Cadophora cf. luteo-olivacea (KC180677)	497/504 (99%)	Hair root of plant in Argentina
R2	I14F- 01208	Glaciozyma sp. 1	KR698804	<i>Glaciozyma martinii</i> (HQ432816)	556/557 (99%)	Sediment from glacier in Italy
	I14F- 01209	Cadophora sp. 2	KR698805	Cadophora luteo-olivacea (KF053558)	504/504 (100%)	Wood in Antarctica
	I14F- 01210	Mortierella sp.	KR698806	Mortierella antarctica (JX975843)	598/599 (99%)	Soil in North Victoria Land, Antarctica
	I14F- 01211	Vishniacozyma sp. 2	KR698807	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01212	Phenoliferia sp. 1	KR698808	Phenoliferia glacialis (JQ857032)	541/542 (99%)	King George Island, Antarctica
	I14F- 01213	Vishniacozyma sp. 2	KR698809	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
R3	I14F- 0214	Vishniacozyma sp. 2	KR698810	Vishniacozyma victoriae (KP299253)	436/437 (99%)	Antarctic snow
	I14F- 01215	Vishniacozyma sp. 2	KR698811	Vishniacozyma victoriae (KP299253)	436/437 (99%)	Antarctic snow
	I14F- 01216	Dioszegia sp. 1	KR698812	Dioszegia rishirensis (AB545810)	418/424(99%)	Soil from island in Japan
	I14F- 01217	<i>Keissleriella</i> sp.	KR698813	Keissleriella trichophoricola (KJ869113)	459/466 (98%)	Plant in Italy
	I14F- 01218	Cadophora sp. 1	KR698814	Cadophora melinii (JN689950)	462/484 (95%)	Wood tissues of mechanical wounds in silver birch
	I14F- 01219	Paraleptosphaeria sp.	KR698815	Leptosphaeria sp. (KJ542322)	492/492 (100%)	Root of alpine plant in China
	I14F- 01220	<i>Tetracladium</i> sp.	KR698816	Tetracladium sp. (JX029123)	485/495 (98%)	Qinghai-Tibet Plateau
	I14F- 01221	Venturia sp.	KR698817	Venturia sp. (AB916509)	499/502 (99%)	Bird feather in Norway
	I14F- 01222	<i>Goffeauzyma</i> sp. 2	KR698818	Goffeauzyma victoriae (NR073228)	586/586 (100%)	Unreported
	I14F- 01223	<i>Goffeauzyma</i> sp. 2	KR698819	Goffeauzyma victoriae (NR073228)	587/587 (100%)	Unreported
	I14F- 01224	Vishniacozyma sp. 2	KR698820	Vishniacozyma victoriae (AY040654)	436/437 (99%)	Antarctic snow
	I14F- 01225	Mrakia sp. 2	KR698821	Mrakia gelida (KM819098)	578/578 (100%)	Water from glacier in Antarctica
	I14F- 01226	Leptosphaeria sp. 1	KR698822	Leptosphaeria sclerotioides (KM280063)	481/501 (96%)	Seed in Switzerland

Appendix Table 1Molecular identification of 134 fungal strains isolated from aquatic habitats of the Ny-Ålseund region:
maximum identification conducted using BLASTn searches of the ITS region

Continued table

Sample code	Isolate code	Identification	GenBank accession number	Closest related species (GenBank accession number)	Number of bp (Maximum identify)	Reported habitat
R4	I14F- 01227	Pseudeurotium sp.	KR698823	Pseudeurotium bakeri (KP055601)	509/512 (99%)	Crop field soil in South Korea
	I14F- 01228	Mrakia sp. 1	KR698824	Mrakia blollopis (AB908178)	573/577 (99%)	Lake sediment in Antarctica
	I14F- 01229	Dioszegia sp. 2	KR698825	Dioszegia fristingensis (EU070927)	425/432 (98%)	Phylloplane
	I14F- 01230	Phenoliferia sp. 1	KR698826	Phenoliferia glacialis (JQ857037)	543/543 (100%)	King George Island, Antarctica
	I14F- 01231	Phenoliferia sp. 1	KR698827	Phenoliferia glacialis (JQ857037)	542/543 (99%)	King George Island, Antarctica
	I14F- 01232	Cadophora sp. 3	KR698828	Cadophora viticola (HQ661097)	503/506 (99%)	Grapevine nurseries in Spain
	I14F- 01233	Pseudogymnoascus sp.	KR698829	Pseudogymnoascus pannorum (KP411562)	502/504 (99%)	Soil in caves, Canada
	I14F- 01234	Varicosporium sp.	KR698830	Varicosporium delicatum (JQ412864)	455/478 (95%)	Streams in Alaska
	I14F- 01235	Cadophora sp. 2	KR698831	Cadophora cf. luteo-olivacea (KC180677)	497/504 (99%)	Hair root of plant in Argentina
	I14F- 01236	Mortierella sp.	KR698832	Mortierella antarctica (JX975843)	590/595 (99%)	Soil in North Victoria Land, Antarctica
	I14F- 01237	Vishniacozyma sp. 1	KR698833	Vishniacozyma tephrensis (JX188132)	437/437 (100%)	Plant in USA
	I14F- 01238	Pleosporales sp.	KR698834	Uncultured fungus clone (KF297131)	496/496 (100%)	Arctic soil in USA (78.78°N, 103.55°W)
	I14F- 01309	Pseudogymnoascus sp.	KR698905	Pseudogymnoascus pannorum (KP411562)	499/504 (99%)	Soil in caves, Canada
R6	I14F- 01239	Phenoliferia sp.	KR698835	Phenoliferia glacialis (JQ857037)	542/543 (99%)	King George Island, Antarctica
	I14F- 01240	Cadophora sp. 2	KR698836	Cadophora cf. luteo-olivacea (KC180677)	496/504 (98%)	Hair root of plant in Argentina
	I14F- 01241	Paraleptosphaeria sp.	KR698837	Leptosphaeria sp. (KJ542322)	479/486 (99%)	Root of alpine plant in China
	I14F- 01242	Helotiales sp. 2	KR698838	Uncultured <i>Clathrosphaerina</i> clone (HQ212329)	479/587 (98%)	Arctic soil
	I14F- 01243	Mortierella sp.	KR698839	Mortierella antarctica (JX975843)	590/595 (99%)	Soil in North Victoria Land, Antarctica
	I14F- 01244	Tolypocladium sp.	KR698840	<i>Tolypocladium cylindrosporum</i> (FJ025173)	511/512 (99%)	Soil from alpine grassland
	I14F- 01245	Vishniacozyma sp. 2	KR698841	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01301	Mortierella sp.	KR698897	Mortierella antarctica (JX975843)	590/595 (99%)	Soil in North Victoria Land, Antarctica
R7	I14F- 01246	Cadophora sp. 2	KR698842	Cadophora cf. luteo-olivacea (KC180677)	490/501 (98%)	Hair root of plant in Argentina
	I14F- 01247	Phenoliferia sp. 1	KR698843	Phenoliferia glacialis (JQ857037)	542/543 (99%)	King George Island, Antarctica
	I14F- 01248	Vishniacozyma sp. 2	KR698844	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01249	Helotiales sp. 3	KR698845	Helotiales sp. (JX852359)	490/508 (96%)	Moss in Antarctica
	I14F- 01250	Mrakia sp. 2	KR698846	Mrakia gelida (KM819098)	578/578 (100%)	Water from glacier in Antarctica

Continued table

Sample code	Isolate code	Identification	GenBank accession number	Closest related species (GenBank accession number)	Number of bp (Maximum identify)	Reported habitat
R7	I14F- 01251	Cadophora sp. 2	KR698847	Cadophora luteo-olivacea (KF053558)	504/504 (100%)	Wood in Antarctica
	I14F- 01252	Cadophora sp. 3	KR698848	Cadophora viticola (HQ661097)	504/506 (99%)	Grapevine nurseries in Spain
	I14F- 01253	Dioszegia sp. 2	KR698849	Dioszegia fristingensis (EU070927)	426/433 (98%)	Phylloplane
	I14F- 01254	Phenoliferia sp. 1	KR698850	Phenoliferia glacialis (JQ857037)	541/542 (99%)	King George Island, Antarctica
	I14F- 01255	Tolypocladium sp.	KR698851	Tolypocladium cylindrosporum (FJ025173)	518/519 (99%)	Soils from alpine grassland
	I14F- 01256	Mortierella sp.	KR698852	<i>Mortierella antarctica</i> (JX975843)	590/595 (99%)	Soil in North Victoria Land, Antarctica
	I14F- 01302	Vishniacozyma sp. 2	KR698898	Vishniacozyma victoriae (KP299253)	432/432 (100%)	Antarctic snow
L1	I14F- 01257	Leotiomycetes sp.	KR698853	Uncultured Leotiomycetes clone (JF449892)	426/466 (91%)	Leaf litter in Austria
	I14F- 01258	Cadophora sp. 2	KR698854	Cadophora cf. luteo-olivacea (KC180677)	485/498 (97%)	Hair root of plant in Argentina
	I14F- 01259	Mrakia sp. 2	KR698855	Mrakia gelida (KM819098)	578/578 (100%)	Water from glacier in Antarctica
	I14F- 01260	Cadophora sp. 2	KR698856	Cadophora luteo-olivacea (KF053558)	504/504 (100%)	Wood in Antarctica
	I14F- 01261	Phenoliferia sp. 1	KR698857	Phenoliferia glacialis (JQ857037)	542/543 (99%)	King George Island, Antarctica
	I14F- 01262	Vishniacozyma sp. 2	KR698858	Vishniacozyma victoriae (KP299253)	437/437 (99%)	Antarctic snow
	I14F- 01263	Goffeauzyma sp. 2	KR698859	Goffeauzyma victoriae (NR073228)	588/588 (100%)	Unreported
	I14F- 01264	Dioszegia sp. 2	KR698860	Dioszegia fristingensis (EU070927)	425/432 (98%)	Phylloplane
	I14F- 01265	Phoma sp.	KR698861	Phoma herbarum (KF889057)	485/486 (99%)	Unreported
	I14F- 01266	Vishniacozyma sp. 2	KR698862	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01267	Cadophora sp. 2	KR698863	Cadophora cf. luteo-olivacea (KC180676)	497/507 (99%)	Hair root of plant in Argentina
L2	I14F- 01268	Cadophora sp. 2	KR698864	Cadophora cf. luteo-olivacea (KC180677)	491/501 (98%)	Hair root of plant in Argentina
	I14F- 01269	Holtermanniella sp.	KR698865	Holtermanniella festucosa (JX188167)	512/513 (99%)	Plant in USA
	I14F- 01270	Pseudogymnoascus sp.	KR698866	Pseudogymnoascus pannorum (KP411562)	501/502 (99%)	Soil in caves, Canada
	I14F- 01271	Vishniacozyma sp. 2	KR698867	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01272	Holtermanniella sp.	KR698868	Holtermanniella festucosa (JX188167)	512/513 (99%)	Plant in USA
	I14F- 01273	Cystobasidium sp. 1	KR698869	Cystobasidium laryngis (AF190014)	518/521 (99%)	Unreported
	I14F- 01274	Vishniacozyma sp. 2	KR698870	Vishniacozyma victoriae (KP299253)	436/437 (99%)	Antarctic snow
	I14F- 01275	Cystobasidium sp. 1	KR698871	Cystobasidium laryngis (GQ911550)	519/521 (99%)	Glacier, Apennines, Italy

						Continued table
Sample code	Isolate code	Identification	GenBank accession number	Closest related species (GenBank accession number)	Number of bp (Maximum identify)	Reported habitat
	I14F- 01276	Dioszegia sp. 2	KR698872	Dioszegia fristingensis (EU070927)	427/433 (99%)	Phylloplane
	I14F- 01277	Phenoliferia sp. 1	KR698873	Phenoliferia glacialis (JQ857037)	542/543 (99%)	King George Island, Antarctica
	I14F- 01278	Phenoliferia sp. 1	KR698874	Phenoliferia glacialis (JQ857037)	542/543 (99%)	King George Island, Antarctica
	I14F- 01279	Phenoliferia sp. 2	KR698875	Phenoliferia psychrophenolica (KC333169)	548/550 (99%)	Cryoconite holes on glaciers in Svalbard
	I14F- 01307	Leptosphaeria sp. 2	KR698903	Leptosphaeria sclerotioides (EU265669)	495/501 (99%)	Plant seed in Switzerland
L3	I14F- 1280	Tremellales sp.	KR698876	Uncultured Tremellales clone (FJ553848)	432/500 (86%)	Forest soil in Canada
	I14F- 01281	Leucosporidiella sp.1	KR698877	Leucosporidiella fragaria (NR073287)	567/579 (98%)	Unreported
	I14F- 01282	Phenoliferia sp. 2	KR698878	Phenoliferia psychrophenolica (KC333169)	543/553 (98%)	Cryoconite sediment in Svalbard
	I14F- 01283	Leucosporidiella sp. 2	KR698879	Leucosporidiella muscorum (NR073286)	536/552 (97%)	Unreported
	I14F- 01284	<i>Glarea</i> sp.	KR698880	Glarea lozoyensis (NR137138)	432/447 (97%)	Unreported
	I14F- 01285	Leucosporidiella sp. 2	KR698881	Leucosporidiella muscorum (NR073286)	535/553 (97%)	Unreported
	I14F- 01286	Vishniacozyma sp. 2	KR698882	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01287	Phenoliferia sp. 2	KR698883	Phenoliferia psychrophenolica (KC333169)	552/552 (100%)	Cryoconite sediment in Svalbard
	I14F- 01288	Hypocreales sp.	KR698884	Acremonium sp. (JX171161)	525/528 (99%)	Antarctica
X1	I14F- 01289	Mrakia sp. 3	KR698885	Mrakiella cryoconiti (JX188193)	569/570 (99%)	Superficial glacial melting water from Alps
	I14F- 01290	Helotiales sp. 4	KR698886	Uncultured Helotiales clone (HQ212304)	498/499 (99%)	Arctic soil
	I14F- 01291	Phenoliferia sp. 1	KR698887	Phenoliferia glacialis (JQ857032)	541/543 (99%)	King George Island, Antarctica
	I14F- 01292	Phenoliferia sp. 1	KR698888	Phenoliferia glacialis (JQ857032)	541/543 (99%)	King George Island, Antarctica
	I14F- 01293	Phenoliferia sp. 1	KR698889	Phenoliferia glacialis (JQ857032)	541/542 (99%)	King George Island, Antarctica
	I14F- 01294	Phenoliferia sp. 1	KR698890	Phenoliferia glacialis (JQ857032)	541/543 (99%)	King George Island, Antarctica
	I14F- 01295	Helotiales sp. 4	KR698891	Uncultured Helotiales clone (HQ212304)	487/488 (99%)	Arctic soil
	I14F- 01296	Helotiales sp. 4	KR698892	Uncultured Helotiales clone (HQ212304)	479/479 (100%)	Arctic soil
	I14F- 01297	Vishniacozyma sp. 2	KR698893	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01303	Helotiales sp. 4	KR698899	Uncultured Helotiales clone (HQ212304)	504/507 (99%)	Arctic soil
	I14F- 01304	Helotiales sp. 4	KR698900	Uncultured Helotiales clone (HQ212304)	504/507 (99%)	Arctic soil
	I14F- 01305	Phenoliferia sp. 1	KR698901	Phenoliferia glacialis (JQ857032)	541/543 (99%)	King George Island, Antarctica
	I14F- 01306	Cadophora sp. 2	KR698902	Cadophora luteo-olivacea (KF053558)	504/504 (100%)	Wood in Antarctica

	4	1
4	+	
	•	1

						Continued table
Sample	Isolate	Identification	GenBank accession	Closest related species	Number of bp (Maximum	Reported habitat
 	I14F-	Vishniacomma sp 2	number	Vishniacozyma victoriae	identify)	Antarctic snow
$\Lambda 2$	01298 I14F-	D:	KR090094	(KP299253) Dioszegia fristingensis	437/437 (100%)	King George Island,
	01299 1145-	Dioszegia sp. 2	KR698895	(JQ857038) Glaciozyma watsonii	434/436 (99%)	Antarctica Soil from South Victoria
	01300	Glaciozyma sp. 2	KR698896	(EU149804)	580/581 (99%)	land, Antarctica
	114F- 01308	Mrakia sp. 2	KR698904	Mrakia gelida (KM819098)	578/578 (100%)	Antarctica
	I14F- 01310	Vishniacozyma sp. 2	KR698906	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01311	Vishniacozyma sp. 3	KR698907	Vishniacozyma carnescens (KP131895)	436/437 (99%)	Cloacal swab, bird
	I14F- 01312	Mycoarthris sp.	KR698908	Uncultured <i>Mycoarthris</i> clone (JF449666)	456/470 (97%)	Beech litter in Austria
	I14F- 01313	Cystobasidium sp. 2	KR698909	<i>Cystobasidium psychroaquaticum</i> (JN572895)	518/521 (99%)	Springtail Sminthuridae feeding on fungus
	I14F- 01314	Vishniacozyma sp. 2	KR698910	Vishniacozyma victoriae (KP299253)	435/436 (99%)	Antarctic snow
	I14F- 01316	Vishniacozyma sp. 2	KR698912	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
S 1	I14F- 01315	Dioszegia sp. 2	KR698911	Dioszegia fristingensis (JQ857038)	434/435 (99%)	King George Island, Antarctica
	I14F- 01317	Vishniacozyma sp. 2	KR698913	Vishniacozyma victoriae (KP299253)	435/437 (99%)	Antarctic snow
	I14F- 01318	Mrakia sp. 1	KR698914	Mrakia blollopis (AB908178)	573/577 (99%)	Lake sediment in Antarctica
	I14F- 01319	<i>Goffeauzyma</i> sp.	KR698915	Goffeauzyma victoriae (NR073228)	588/588 (100%)	Unreported
	I14F- 01320	Dioszegia sp. 2	KR698916	Dioszegia fristingensis (EU070927)	426/433 (98%)	Phylloplane
	I14F- 01321	Vishniacozyma sp. 2	KR698917	Vishniacozyma victoriae (KP299253)	437/437 (100%)	Antarctic snow
	I14F- 01322	Godroniaceae sp.	KR698918	Godronia cassandrae (KC595271)	473/480 (99%)	Plant in Germany
	I14F- 01323	Tetracladium sp.	KR698919	Tetracladium sp. (JX029123)	490/495 (99%)	Qinghai-Tibet Plateau
	I14F- 01324	Dioszegia sp. 3	KR698920	Dioszegia aurantiaca (EU266497)	412/422 (98%)	Phylloplane
S2	I14F- 01325	Cystobasidium sp. 2	KR698921	<i>Cystobasidium psychroaquaticum</i> (JN572895)	516/521 (99%)	Springtail Sminthuridae feeding on fungus
	I14F- 01326	Mrakia sp. 2	KR698922	Mrakia gelida (KM819098)	578/578 (100%)	Water from glacier in Antarctica
	I14F- 01327	Pseudeurotium sp.	KR698923	Pseudeurotium bakeri (KP055601)	502/505 (99%)	Crop field soil in South Korea
	I14F- 01328	Mortierella sp.	KR698924	<i>Mortierella antarctica</i> (JX975843)	590/595 (99%)	Soil in North Victoria Land, Antarctica
	I14F- 01329	Tolypocladium sp.	KR698925	<i>Tolypocladium cylindrosporum</i> (FJ025173)	512/515 (99%)	Soils from alpine grassland in Qilian
	I14F- 01330	Helotiales sp. 2	KR698926	Uncultured <i>Clathrosphaerina</i> clone (HQ212329)	490/500 (98%)	Arctic soil
	I14F- 01331	Glaciozyma sp. 2	KR698927	Glaciozyma watsonii (EU149804)	581/582 (99%)	Soil from South Victoria Land, Antarctica
	I14F- 01332	Helotiales sp. 5	KR698928	Variosporium delicatum (KF730848)	450/478 (94%)	Aquatic environment
	I14F- 01333	<i>Glaciozyma</i> sp. 2	KR698929	Glaciozyma watsonii (EU149804)	581/582 (99%)	Soil from South Victoria Land, Antarctica
	I14F- 01334	Phoma sp.	KR698930	Phoma herbarum (KF889057)	471/474 (99%)	Unreported

	Stream	Pond	Ice	Estuary
Cadophora sp. 1	1	0	0	0
Cadophora sp. 2	10	4	1	0
Cadophora sp. 3	2	0	2	0
Cystobasidium sp. 1	0	2	0	0
<i>Cystobasidium</i> sp. 2	0	0	1	1
Dioszegia sp. 1	1	0	0	0
Dioszegia sp. 2	2	2	l	2
Dioszegia sp. 3	0	0	0	1
Glaciozyma sp. 1	1	0	0	0
Glaciozyma sp. 2	0	0	1	2
Giarea sp.	0	1	0	0
Godroniaceae sp.	0	0	0	1
Goffeauzyma sp.	2	l	0	l
Helotiales sp. 1	1	0	0	0
Helotiales sp. 2	1	0	0	1
Helotiales sp. 3	1	0	0	0
Helotiales sp. 4	0	0	5	0
Helotiales sp. 5	0	0	0	1
Holtermanniella sp.	0	2	0	0
Hypocreales sp.	0	1	0	0
Keissleriella sp.	1	0	0	0
Leotiomycetes sp	0	1	0	0
Lentosphaeria sp. 1	1	0	0	0
Leptosphaeria sp. 1	0	1	0	0
Lepiosphaeria sp. 2	0	1	0	0
Leucosportatetta sp. 1	0	1	0	0
Leucosportalella sp. 2	1	2	0	0
Mortierella sp.	4	0	0	1
Mrakia sp. 1	2	0	0	1
<i>Mrakia</i> sp. 2	1	1	1	1
Mrakia sp. 3	0	0	1	0
Mycoarthris sp.	0	0	1	0
Paraleptosphaeria sp.	2	0	0	0
Phenoliferia sp. 1	5	3	5	0
Phenoliferia sp. 2	0	3	0	0
Phoma sp.	0	1	0	1
Pleosporales sp.	1	0	0	0
<i>Pseudeurotium</i> sp	1	0	0	1
Pseudogymnoascus sp	3	1	0	0
Taphrina sp.	1	0	0	0
Tetracladium sp.	1	0	0	1
Tolypocladium sp.	2	0	0	1
Tremellales sp.	0	1	0	0
Varicosporium sp.	2	0	0	0
Venturia sp.	1	0	0	0
Vishniacozyma sp. 1	1	0	0	0
Vishniacozyma sp. 2	7	5	3	2
Vishniacozyma sp. 3	0	0	1	0
Number of fungal isolates	59	33	23	19
Number of fungal taxa	28	18	12	16
Shannon index (H')	4.34	3.92	3.22	3.93

Appendix Table 2 Distribution of fungal species in the four aquatic habitats from the Ny-Ålseund region: number of isolates in different aquatic habitats