Imported diphyllbothriasis in Switzerland: molecular methods to define a clinical case of *Diphyllbothrium* infection as *Diphyllbothrium dendriticum*, August 2010

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Following a first clinical case of infection by *Diphyllbothrium dendriticum* in Switzerland in 2006, we report a second case in the country. The species was identified by molecular methods. In the Swiss, French and Italian subalpine regions, human diphyllbothriasis has seen a comeback since the late 1980's, and *Diphyllbothrium latum* is usually considered the causative agent of the disease. In addition, several locally acquired and imported clinical infections due to allochthonous *Diphyllbothrium* species have been documented in the last years. Due to the colonisation potential of these parasites and their probably underestimated presence in the human population, there is a need for discriminating them at the medical laboratory level. Because the morphological characters are very similar among the different taxa, a correct identification requires the use of molecular methods. Molecular identification would improve diagnosis and help monitor the distribution of *Diphyllbothrium* species in Europe.

Introduction

Diphyllbothriasis is due to tapeworms of the genus *Diphyllbothrium* and is acquired through the consumption of raw or poorly cooked freshwater fish. In distinct stages of their life cycle, *Diphyllbothrium* spp. occupy different host species. The eggs present in the water are ingested by small crustaceans, such as copepods, constituting the first intermediate hosts in which they develop to a larval stage. As crustaceans are predated by fish, fish become the second intermediate hosts where *Diphyllbothrium* larvae further develop. The definite hosts are fish eating mammals (including humans) or birds, where *Diphyllbothrium* matures into an adult stage.

Human diphyllbothriasis is often a mild illness: about half of the patients are asymptomatic, while the others mostly suffer from minor discomforts, such as diarrhoea, abdominal pain and digestive troubles. Rare cases of megaloblastic anaemia associated with

vitamin B12 deficiency have been reported in the literature [1].

Globally, the incidence of human diphyllbothriasis has decreased in the last 20 years, particularly in northern and eastern Europe [2]. Nevertheless, in some Swiss, French and Italian subalpine regions the disease has seen a comeback, as shown by the more than 530 cases reported since 1987 from around lakes Lemman (Geneva), Morat, Bienne, Maggiore, Lario (Como), Iseo and Garda [2]. In these areas, the species *Diphyllbothrium latum* is considered the causative agent of diphyllbothriasis. However, infections due to allochthonous species (*D. nihonkaiense*, *D. dendriticum*) have been recently documented [3-7]. A first clinical case due to *D. dendriticum* was diagnosed in 2006 in Switzerland, raising the question of potential transmission to susceptible intermediate hosts present in the local environment [4]. In this report, we describe a second case of symptomatic infection by *D. dendriticum* in Switzerland. Confirmation of the species was done by molecular identification.

A four year-old boy expelled tapeworm segments in stool, 12 days after returning from a 15-day holiday travel in Singapore and Bali in August 2010. He had been suffering from abdominal pains (cramps) and loose stools since his return to Switzerland. Standard laboratory procedures held in the clinical laboratory (Dianalabs) led to the identification of *Diphyllbothrium* spp., based upon the presence of typical operculated eggs in segments. However, because of the unusual shape of a tapeworm proglottid (longer than wide as if stretched, with an excentred uterus) and the possible Asian origin of infection, the specimen was preserved in 70° ethanol and sent to the Institute of Parasitology in Bern and to the Cantonal Institute of Microbiology in Bellinzona, where it was identified as *D. dendriticum* by molecular methods.

Mebendazole was first administered to the patient, with no curative effect, as confirmed by the persisting presence of eggs in stool after three weeks. The patient was then re-medicated with praziquantel and recovered promptly. No parasites were found upon stool testing six weeks after praziquantel therapy.

The patient's family did not present with symptoms and underwent no further investigation, except for the seven year-old patient's sister who was checked for intestinal parasites but found negative.

Methods

The faecal specimen was processed by standard sedimentation technique [8] to concentrate putatively present *Diphylobothrium* eggs and subsequently assess these by light microscopy. A segment of proglottids of approximately 5 cm length was processed for staining with lacto-acetic carmine according to Rukhadze and Blajin [9].

Genomic DNA from about 50 mg of proglottid tissue was extracted with the DNeasy Blood and Tissue Kit (Qiagen). Polymerase chain reaction (PCR) was performed using the Taq PCR Master Mix Kit (Qiagen) with primers targeting a region of the 5.8S ribosomal RNA (5.8S rRNA) comprising internal transcribed spacers (ITS) 1 and 2 [10], the 18S ribosomal RNA (18S rRNA) [11] and the cytochrome c oxidase subunit 1 gene (cox1) [3,12] sequences. The amplification of all targets was carried out under the following conditions: 5 min at 94 °C, 35 cycles consisting of 30 s at 94 °C, 40 s at 45 °C, 1 min at 72 °C, and a final extension step of 10 min at 72 °C. Amplicons were visualised by electrophoresis in a 0.8% agarose gel containing ethidium bromide, and purified through Sephadex G-50 columns (GE Healthcare). DNA was quantified with a ND-100 Spectrophotometer (NanoDrop Technologies Inc.). Sequencing was performed with the BigDye Terminator Cycle Sequencing Ready Reaction kit (Applied Biosystems), according to the provider's recommendations. Samples were purified by osmosis with 0.025 µm nitrocellulose filters (Millipore) in tris ethylenediamine-tetraacetic acid (TE) buffer pH 8 for two hours. Eight µl of purified solution were placed in 0.5 ml Genetic Analyzer Sample Tubes with 12 µl Hi-Di Formamide (Applied Biosystems). Samples were then loaded in an automated sequencing system (ABI PRISM 310 Genetic Analyzer; Perkin Elmer).

Sequence electropherograms were corrected by using the software EditSeq (DNASTAR Inc.). Their identity was first checked by basic local alignment search tool (BLAST) [13]. Sequence fragments of 657 and 375 nucleotides in length, derived from the PCRs targeting the ITS1-5.8SrRNA and cox1 genes were then respectively compared to representative ITS1-5.8SrRNA or cox1 sequences from different *Diphylobothrium* spp. by pairwise and multiple alignments using ClustalW [14] with the software Molecular Evolutionary Genetics Analysis (MEGA) version 4.0 [15]. Phylogenetic trees

(neighbour-joining method; Kimura-2 parameters; bootstrap test for 500 replicates) were subsequently inferred from the alignments.

Results

Microscopical analyses of the coprological sediment revealed the presence of oval-shape unembryonated eggs (mean size: 49 x 64 µm; range: 48.5–52.5 x 62.5–70 µm), characterised by the presence of a hardly visible operculum and a small knob at the abopercular end (Figure 1).

Microscopical analyses of the stained proglottids revealed the presence of only one set of reproductive organs per proglottid (Figure 2). The central uterine structure showed several rosette-shaped loops. Morphological criteria matched to those described

FIGURE 1

Diphylobothrium dendriticum eggs recovered from a patient stool, Switzerland, 2010

White arrow: operculum; grey arrow: knob.

The pictures were taken under 400x magnification and the mean size of the eggs was 49 x 64 µm.

FIGURE 2

Diphyllbothrium dendriticum segment recovered from a patient stool, Switzerland, 2010

A. *Diphyllbothrium dendriticum* segment.

B. Enlargement of an individual proglottid where the central uterine structure forms several rosette-shaped loops.

for *Diphyllbothrium* spp. [16], while a species-specific identification was not possible based upon morphology.

The nucleotide sequences of the ITS1-5.8S rRNA, 18S rRNA and *cox1* derived from the *Diphyllbothrium* affecting the patient were deposited in public databases and are available in the European Molecular Biology Laboratory (EMBL), GenBank and DNA Data Bank of Japan (DDJB) databases under accession numbers HQ682065, HQ682066 and HQ682067. Due to technical problems, the sequence of ITS2 was not fully obtained and therefore not used in the analysis.

The results of BLAST search showed that all the sequenced targets respectively reached $\geq 99\%$ identity (highest scores) at the nucleotide level with *D. dendriticum* homologous reference sequences of respective GenBank accession numbers FM204787 (ITS1-5.8S rRNA), DQ768164, DQ181945 (18S rRNA) and AM412738 (*cox1*). The 18S rRNA sequence also showed 99% identity with those of *D. ditremum* (GenBank accession numbers: DQ768165, DQ181944) and *D. latum* (GenBank accession number: DQ316795).

The position of the sample sequences of ITS1-5.8S rRNA and *cox1* regions in the phylogenetic trees confirmed the *Diphyllbothrium* spp. affecting the patient as *D. dendriticum* (Nitsch, 1824) (Figures 3 and 4). The phylogenetic tree built on the basis of 18S rRNA sequences is not shown, because this target is not useful for the discrimination of *D. dendriticum* from other species [2].

Discussion

At the time of tapeworm evacuation, the patient was known to have had a meal in a Japanese restaurant in Singapore (on day 4 of his journey) where the family had consumed various dishes (fish, chicken). According to the recollection of the family members, the meal did not contain raw fish. Retrospective investigations revealed that the patient regularly consumed fish, e.g. smoked salmon with pasta at home. This salmon was always bought in the same department store in France and was of the same brand. According to the product information, it belonged to the species *Salmo salar*, was farmed in Norway, smoked in France and guaranteed 'never frozen'. The homemade sauce was made by dropping slices of smoked salmon into boiling cream

FIGURE 3

Neighbour-joining tree based on 5.8S ribosomal RNA internal transcribed spacer 1 sequences, to determine the *Diphyllbothrium* species affecting a patient, Switzerland, 2010

The tree is based on 5.8S ribosomal RNA internal transcribed spacer 1 (5.8S rRNA-ITS1) sequence fragments of 657 bp. The 'sample' refers to the sequence of the unknown *Diphyllbothrium* spp. affecting the patient. GenBank accession numbers of all the sequences used to construct the tree are indicated. On the tree nodes, Kimura-2 parameters bootstrap values >50 for 500 replicates are shown.

FIGURE 4

Neighbour-joining tree based on cytochrome c oxidase subunit 1 sequences, to determine the *Diphyllobothrium* species affecting a patient, Switzerland, 2010

The tree is based on cytochrome c oxidase subunit 1 (cox1) sequence fragments of 375 bp. The 'sample' refers to the sequence of the unknown *Diphyllobothrium* spp. affecting the patient. GenBank accession numbers of all the sequences used to construct the tree are indicated. On the tree nodes, Kimura-2 parameters bootstrap values >50 for 500 replicates are shown.

and leaving to cook for one minute before serving. According to his mother, the patient enjoyed picking pieces of uncooked fish during the preparation of the meal. Short before travelling to south-east Asia, the patient consumed poorly cooked perch fillets (*Perca fluviatilis*) fished in the Lemman lake in a restaurant near home. He also ate sushi in Switzerland one week before tapeworm segments were noticed.

Aquatic birds (especially Laridae) are usually the definitive hosts of *D. dendriticum*, while humans are only occasionally infected. According to literature, the incubation period of *D. dendriticum* in herring gulls (*Larus argentatus*) lasts from five to 20 days [17]. In humans, *Diphyllobothrium* plerocercoids generally develop into mature adults in two to six weeks [2].

Based on these observations, the source of the patient's infection might be either salmon from Norway, unknown fish from Asia or perch from Switzerland. However, because the presence of *D. dendriticum* in perch has not been documented so far, it is unlikely that the perch meal was the source of infection. The sushi meal in Switzerland can be excluded because the symptoms were already present at that time.

In the last two decades, diphyllobothriasis has shown a recrudescence in a number of European countries, especially in the subalpine lakes region [2]. The use of molecular methods also showed the presence of allochthonous *D. nihonkaiense* and *D. dendriticum* in France, Switzerland, Finland and the Czech Republic (Table). Except for two cases of *D. dendriticum* diagnosed in Switzerland [4] and the Czech Republic [18]

that were probably acquired abroad (Norway, Alaska or Canada), most of the documented infections were locally acquired, attributed to imported salmons.

Susceptible intermediate hosts for *D. dendriticum* such as copepods (*Eudiaptomus* and *Cyclops* species [19]) and fish (*Oncorhynchus mykiss*, *Salmo trutta*, *Coregonus clupeaformis*, *C. albula*, *C. lavaterus* [19-21]) are known to be present in Switzerland. This would theoretically allow the introduction and the autochthonous transmission of the parasite. This colonisation potential emphasises the need for correct identification of *Diphyllobothrium* species involved in clinical cases of infection.

Interestingly, in the two Swiss cases due to *D. dendriticum*, the molecular investigation was undertaken because of the unusual shape of some segments.

The transmission of *Diphyllobothrium* plerocercoids may be prevented by freezing fish at -20 °C for one to seven days, depending on its thickness. The Swiss law [22] provides that it is forbidden to sell any kind of fresh fish products (both local and imported) intended to be consumed raw or semi-raw, unless they have been stored at a temperature not higher than -20 °C for a least 24 hours. However, it has been demonstrated that this rule is not always followed properly [23], and of course there is no control of food bought abroad by individuals for personal use. Therefore cooking at 55 °C or more, for at least 5 min, remains the most reliable way to prevent transmission of parasitic worms possibly present in fresh fish [2].

Conclusion

A correct diagnosis has become crucial to evaluate the distribution of human-infecting *Diphyllobothrium* species, as well as their fish hosts, and to prevent the spread of allochthonous parasites in aquatic environments. Due to the difficulties in discriminating the different *Diphyllobothrium* taxa by morphological characters, molecular analysis has proven to be fundamental to identify these helminths at the species level. A cheap and rapid molecular test based on multiplex PCR with partial cytochrome c oxidase subunit 1 (cox1) gene, without the need of sequencing, was recently developed for the differential identification of the most common species infecting humans [24]. In case of atypical specimens of proglottids and eggs and/or specimens derived from patients who have been abroad, this test could be used to verify the parasite's identity. Improving the diagnosis of *Diphyllobothrium* parasites would help to monitor the distribution of species in Europe and trace the source of infections, an important goal at a time when eating habits are changing, fish markets are globalising and climate is changing [25].

TABLE

Case reports of allochthonous *Diphyllobothrium* infections in Europe, identified by molecular methods, 2005–2011

Year of parasite recovery	Country	Patient (age in years)	<i>Diphyllobothrium</i> species	Parasite characteristics	Clinical features	Probable source of infection	Molecular identification	Therapy	Reference
2005	Finland	Man (60)	<i>D. nihonkaiense</i>	ND	ND	ND	Cox1	ND	Data not shown
	France	Woman (44)	<i>D. nihonkaiense</i>	Egg size 41 × 60 µm	Nausea, epigastric pain, diarrhoea	Wild salmon carpaccio (<i>Oncorhynchus keta</i>) imported from Canada (Gulf of Alaska, Pacific Ocean), purchased in France	Cox1, MT-ND3	Single dose of praziquantel at 10 mg/kg.	Yera et al., 2006 [6]
	Switzerland	Woman (55)	<i>D. nihonkaiense</i>	Egg size 56.68–57.82 × 43.58–45.42 µm	Diarrhoea (probably due to another cause); proglottids passed in faeces	Raw salmon and sushi purchased in Switzerland	18S rRNA, cox1, ITS1 and 2	Praziquantel	Wicht et al., 2007 [3]
	Switzerland	Woman (52)	<i>D. nihonkaiense</i>	ND	None	Raw Pacific salmon (<i>O. keta</i>) imported from Canada or North America (Pacific Ocean), purchased in France	18S rRNA, cox1, ITS1 and 2	Single dose of praziquantel	Wicht et al., 2007 [3]
2006	Switzerland	Boy (5)	<i>D. nihonkaiense</i>	Egg size 57.5–65.0 × 40.0–42.5 µm	Mild eosinophilia (7.1%)	Pacific salmon purchased in Switzerland	ITS1, cox1, MT-ND3	Single dose of praziquantel at 12 mg/kg	Shimizu et al., 2008 [5]
	Switzerland	Woman (59)	<i>D. dendriticum</i>	Egg size 49.62–63.86 × 35.75–43.41 µm; some proglottids longer than wide, with excentred utera	Chronically relapsing courses of diarrhoea	Regular consumption of wild salmon, Japanese sushi or fish carpaccio; journeys in Canada, Alaska and Norway	18S rRNA, cox1	Single dose of praziquantel at 10 mg/kg	Wicht et al., 2008 [4]
2008	France	Woman (33)	<i>D. nihonkaiense</i>	Egg size 57 × 44 µm	Persistent, mild eosinophilia	Wild salmon carpaccio or marinade purchased in France	Cox1	Single dose of praziquantel at 15 mg/kg	Paugam et al., 2009 [7]
2010	Switzerland	Boy (4)	<i>D. dendriticum</i>	Egg size 48.5–52.5 × 62.5–70 µm; one of the proglottids longer than wide, with excentred uterus	Abdominal cramps, loose stools	Smoked salmon (<i>Salmo salar</i>) imported from Norway, purchased in France; poorly cooked perch from Switzerland; fish eaten in Asia	ITS1 and 2, 18S rRNA, cox1	Praziquantel	Present paper
2011	Czech Republic	Woman (28)	<i>D. dendriticum</i>	Egg size 49.5 × 64 µm	No symptoms reported; proglottids passed in faeces	Salmons (<i>O. tshawytscha</i> , <i>O. keta</i> , <i>O. kisutch</i> , <i>O. nerka</i> , <i>O. gorbushka</i>) and <i>Coregonus autumnalis</i> eaten in Alaska	Cox1	Single dose of praziquantel (750 mg)	Kuchta et al., 2012 [18]

Cox1: cytochrome c oxidase subunit 1; ITS: internal transcribed spacer of the 5.8S ribosomal ribonucleic acid; MT-ND3: mitochondrially encoded NADH dehydrogenase 3; NADH: reduced nicotinamide adenine dinucleotide; ND: not determined; rRNA: ribosomal ribonucleic acid.

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