

学位論文 （要約）

Molecular Systematics of the Order Phyllobothriidea
(Platyhelminthes: Cestoda) from the Coastal Seas of Japan

（日本周辺海域における吸葉目（扁形動物門：条虫綱）
の分子系統に基づく分類学的研究）

平成 26 年 12 月博士（理学）申請

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Abstract

The order Phyllobothriidea was established based on its molecular phylogeny that separates the family Phyllobothriidae from the order Tetracystida. However, many genera are left *incertae sedis* in Tetracystida, and the order Phyllobothriidea should be revised together with these taxonomically uncertain genera. In this study, the systematics of Phyllobothriidea was revised based on molecular analyses using ssrDNA and lsrDNA. As a result, 17 genera including three new genera, namely *Phyllobothrium*, *Alexandercestus*, *Bilocularia*, *Calliobothrium*, *Calyptribothrium*, *Chimaerocestos*, *Crossobothrium*, *Mitsukurinectes* n. gen., *Monorygma*, *Orygmatobothrium*, *Pelichnibothrium*, *Scyphophyllidium*, *Symcallio*, *Thysanocephalum*, *Trilocularia*, *Yamaguticestus* n. gen., and *Vertebraeovincestus* n. gen., were recognized in Phyllobothriidea. *Clistobothrium* was a junior synonym of *Pelichnibothrium*, and *Marsupiobothrium*, *Nandocestus*, *Orectolobicestus*, *Paraorygmatobothrium*, and *Ruhnkecestus* were junior synonyms of *Scyphophyllidium*. *Monorygma megacotyla* was a junior synonym of *Ph. squali*, and *Ph. squali* was transferred to *Yamaguticestus*. *Marsupiobothrium gobelinus* was transferred to *Mitsukurinectes*, and *Ph. biacetabulatum* was transferred to *Anthocephalum* of Rhinebothriidea. The sequences of the larval species, namely *Pe. caudatum*, *Ph. delphini*, and *Mo. grimaldii*, were located in the *Pelichnibothrium* clade but did not match those of any adults. These three larval species were considered valid species of *Pelichnibothrium*.

The most important taxonomic characteristics of Phyllobothriidea had been believed to be in its scolex and proglottid, but the morphological characteristics did not reflect the phylogeny. Five types of bothridium on the scolex, namely cup, crumple, divided, flat, and loculate margin, were observed in many lineages, but only sac type was uniquely found in *Sc. giganteum*. The hook of scolex was found to be evolved in multiple lineages. Moreover, it was observed not only in Phyllobothriidea but also in Tetracystida and Onchoproteocephalidea. The presence of lacinate and non-lacinate proglottids was also seen in multiple lineages. Further molecular information is essential to clarify the phylogeny of Phyllobothriidea.

Taxonomic studies of Phyllobothriidea have scarcely been done in the coastal seas of Japan, and only 14 species of four genera have been reported thus far. In this study, I have identified 34 species of 14 genera including 14 new species and three new genera from Japanese waters, of which 12 species and five genera are new to Japan. According to the new systematics in this study, the Phyllobothriidea fauna of

the coastal seas of Japan has a total of 47 species of 15 genera.

The life history of Phyllobothriidea has not been studied extensively because identification of its larvae is difficult due to lack of taxonomic characteristics of the larvae. In this study, the identification of phyllobothriidean larvae was attempted by molecular analyses using lsrDNA and mtDNA COI. On investigating 79 species of teleost fishes, mollusks, and decapod crustaceans collected from the coastal seas of Japan, the larvae of Phyllobothriidea or Tetraphyllidea were collected from 25 host species. Molecular analyses of 42 plerocercoids and merocercoids from 25 host species, with the adult and larval species' sequences, revealed plerocercoids of *Pelichnibothrium caudatum* (17 plerocercoids), *Pe. montaukensis* (2), *Pe. cf. montaukensis* (2), *Scyphophyllidium carcharhinus* (1), and *Vertebraeovicestus dobukasube* (1) and merocercoids of *Pe. delphini* (2 merocercoids) and *Pe. grimaldii* (2). Only five plerocercoids were identified to be Phyllobothriidea. The other 10 plerocercoids were found to belong to *Anthobothrium* of Tetraphyllidea. On comparison of the morphology between adults and plerocercoids of the same species, the apical sucker on plerocercoids was shown to be lost in their adult stages. The morphology of bothridium also greatly changed in growth from plerocercoids to adult stages. Molecular identification is a very effective method for the exact identification of phyllobothriidean larvae. In this study, a part of the life cycle of three species was revealed by the exact molecular identification of its larvae.

Comparison of the phyllobothriidean lineage with the host lineage suggested that both host-switch and host-range expansion occurred frequently in the evolution of Phyllobothriidea. These events may play important roles in the diversification of Phyllobothriidea.

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DISCLAIMER: The present study is not issued for the permanent scientific record, and no part of it is to be considered published within the meaning of the International Code of Zoological Nomenclature (see Code, Art. 8b).

General introduction

Cestodes, which infest sharks, skates, and rays (Chondrichthyes: Elasmobranchii), have long attracted parasitologists' attention, probably due to their species diversity and unique morphological characteristics. Therefore, many taxonomic studies have been accumulated to date. However, taxa with unsettled taxonomy have been widely recognized. This is due to species diversity and unique morphological characteristics of cestodes resident in elasmobranchs. For example, the order Tetracanthocephala previously considered as the most diverse, included four families: Onchobothriidae, Phyllobothriidae, Lecanicephalidae, and Ichthyotaeniidae. Subsequently, twenty families were recognized in this order (Table 1). Recent phylogenetic analyses employing morphology (Euzet et al., 1981; Brooks et al., 1991; Hoberg et al., 1997; Caira et al., 1999; 2001) and biomolecular markers (Mariaux, 1998; Olson & Caira, 1999; Kodedová et al., 2000; Olson et al., 2001; Waeschenbach et al., 2007, 2012) revealed that the order Tetracanthocephala was, in fact, polyphyletic.

The order Phyllobothriidae was recently elected from one of the families of the polyphyletic order Tetracanthocephala, based on molecular phylogenetic analysis by Caira et al. (2014). They examined 134 species representing 97 genera across the 15 cestode orders. However, the statistical support for the inclusion of Phyllobothriidae was relatively weak compared with support for that of other orders including Litobothriidae, Lecanicephalidae, Rhinebothriidae, Cathetocephalidae, and Onchoproteocephalidae. Moreover, they also withheld to dismantle the order Tetracanthocephala and remained several uncertain genera in polyphyletic Tetracanthocephala. These uncertainties seem to be consequent to inadequate sampling density (and hence statistical power) across the Tetracanthocephala.

Elasmobranch fauna in the waters around Japan are relatively abundant, comprising 205 species (Nakabo, 2013) which correspond to approximately 1/5 of all elasmobranch species overall. In contrast to the diversity of host elasmobranchs, investigations of parasitic cestode fauna in the coastal seas of Japan have been limited to only 14 species and several unidentified forms from 11 host species, which have been detailed in five reports to date (Yoshida, 1917; Yamaguti, 1934, 1952, 1960; Yamaguchi et al., 2000; Table 2). It is strongly suggested that diverse cestodes, including new species, must infect elasmobranchs in the coastal seas of Japan.

As detailed in Chapter I of the present study, the author conducted several molecular phylogenetic analyses on a wide variety of cestodes using newly collected specimens through intensive and extensive parasitological investigations on

elasmobranchs in the coastal seas of Japan. The sequence data examined by Caira et al. (2014) was augmented to elucidate the precise nature of species diversity and phylogeny of elasmobranch cestodes with special reference to the order Phyllobothriidea. The most accurate, up-to-date statistical analytical techniques were also employed.

Phyllobothriidean cestodes have two or more intermediate hosts for larval development in addition to a final host for adult stages (Euzet, 1952; Caira, 1990). The parasites migrate from an intermediate host to the advanced-stage host through the food chain during their life cycle. Adult worms occupy the spiral intestines of elasmobranchs and chimaeras, and eggs are released into sea water. The eggs hatch and develop into hexacanth embryos, which are usually consumed by copepods or euphausiids, representing the first intermediate host wherein they develop into procercoid larvae. The first intermediate host is consumed by the second intermediate host, which is typically an invertebrate of one of the seven phyla, teleosts, and cetaceans, and procercoids grow to merocercoid or plerocercoid larvae (Caira & Reyda, 2005). When the second intermediate host animals are eaten by elasmobranchs and chimaeras, merocercoids and plerocercoids mature into adult worms. Cestode larvae are frequently found during parasitological surveys of marine fishes, marine mammals, and marine invertebrates. However, these larvae can barely be identified because potentially-distinguishing morphological characteristics are not yet known. This has obstructed detailed characterization of intermediate hosts and life cycle stages for not only phyllobothriidean cestodes but also marine parasites in general.

Molecular methods, i.e., molecular sequence comparisons between adults and larvae should be effective in linking larvae to adults. In Chapter II of the present study, the author employs such molecular techniques in efforts to identify a wide variety of larval cestodes collected from marine fishes, marine mammals, and marine invertebrates, using the sequence data obtained from adults.

Chapter I

Taxonomic revision of the order Phyllobothriidea based on molecular analysis

1. Introduction

Recent taxonomic and phylogenetic studies on cestodes parasitizing in elasmobranchs have been focused on solving polyphyly of the order Tetracophyllidea. Olson & Caira (2001) resurrected the order Litobothriidea and transferred the family Litobothriidae from Tetracophyllidea to Litobothriidea. Caira et al. (2005) resurrected the order Cathetocephalidea and transferred the families Cathetocephalidae and Disculicipitidae from Tetracophyllidea to Cathetocephalidea. Healy et al. (2009) elected a new order Rhinebothriidea and transferred a part of Phyllobothriidae from Tetracophyllidea to Rhinebothriidea. These taxonomical decisions were successful, but polyphyly of Tetracophyllidea was not resolved.

Recently, Ruhnke (2011) re-examined 78 genera historically associated with the family Phyllobothriidae and reported that only *Phyllobothrium* was unambiguously valid, 16 genera were provisionally valid, five genera were *nomina dubia*, one genus was *nomen ad interium*, nine genera were *genera inquirendae*, and 17 genera were *incertae sedis* within the family Phyllobothriidae (Table 3). He transferred eight valid, nine provisionally valid, and one *incertae sedis* genera to the order Rhinebothriidea. In addition, he transferred two valid genera and two genera *inquirenda* to the other tetracophyllidean family Serendipidae (Table 3). He also synonymized the remaining seven genera to some other tetracophyllidean or rhinebothriidean genera. Later, a new genus *Alexandercestus* was elected in Phyllobothriidae by Ruhnke & Workman (2013), and the family Phyllobothriidae included the following 18 genera: *Phyllobothrium*, *Alexandercestus*, *Bibursibothrium*, *Calyptrobothrium*, *Cardiobothrium*, *Clistobothrium*, *Crossobothrium*, *Doliobothrium*, *Flexibothrium*, *Marsupiobothrium*, *Monorygma*, *Nandocestus*, *Orectolobicestus*, *Orygmatobothrium*, *Paraorygmatobothrium*, *Ruhnkecestus*, *Scyphophyllidium*, and *Thysanocephalum* (Ruhnke, 2011; Ruhnke & Workman, 2013).

Recently, Bayesian Inference (BI) analyses of various combinations of complete small subunit ribosomal DNA (ssrDNA) sequences and complete or partial large subunit ribosomal DNA (lsrDNA) sequences that included 134 species representing 97 genera across the 15 eucestode orders, were conducted by Caira et

al. (2014). A monotypic clade consisted of 11 genera, of which 10 genera corresponded to those in the family Phyllobothriidae *sensu* Ruhnke (2011), emerged through an analysis of the complete *ssrDNA* along with the partial *lsrDNA* (D1–D3) sequences of 134 taxa. They raised the clade to the new order Phyllobothriidea elected from a part of Phyllobothriidae and Chimaerocestidae of Tetracanthellidae. Another new order Onchoproteocephalidea was also elected from the order Proteocephalidea and a part of from the family Onchobothriidae.

A monophyletic order Phyllobothriidea was first recognized on a molecular basis by Caira et al. (2014), although the statistical significance was low when comparing with other orders. However, the family Phyllobothriidae, revised by Ruhnke (2011), has not been completely examined on a molecular basis and *incertae sedis* still remains within the family. Moreover, the polyphyly of the order Tetracanthellidae remains unclear because many uncertain genera in polyphyletic Tetracanthellidae remained in the study by Caira et al. (2014).

In Chapter I, 419 specimens of cestodes were obtained and intensive and extensive parasitological investigations were performed on elasmobranchs, teleosts, and mammals in the coastal seas of Japan. The specimens were identified, described, illustrated, and sequenced for molecular analysis. In addition to the original data, data examined by Caira et al. (2014) were also included in the present analysis to precisely elucidate the species diversity and phylogeny of the order Phyllobothriidea and related taxa including the order Tetracanthellidae.

2. Materials and Methods

2.1 Specimens

Elasmobranchs were widely caught in the coastal seas of Japan, including the Ogasawara and Yaeyama Islands (Fig. 1, Table 4). Elasmobranchs and other host animals were collected using scientific research vessels and fishing boats and were also obtained from fishes at fishing ports.

Parasite investigations were performed on a total of 921 individuals obtained from 76 elasmobranch species of 51 genera from 31 families, two teleost species of two genera from two families, and one cetacean species (Table 5). Hosts were preserved on ice or sometimes frozen until examination and dissected to collect cestodes under a stereomicroscope.

2.2 Treatment of specimens

Cestodes were washed and immersed in elasmobranch saline (NaCl, 2 g; KCl, 0.1 g; CaCl₂, 0.02 g; NaHCO₃, 0.002 g; urea, 2.5 g; and distilled water, 100 ml). Specimens were mounted on microscopic slides under coverslips with slight pressure and then fixed in alcohol, formaldehyde, and acetic acid (AFA) fixative (70% ethanol, 40% formaldehyde, and acetic acid glacial at a 20:1:1 ratio) for morphological observations. A part of proglottids of some specimens were separately preserved in 99.5 % ethanol for molecular analyses.

Specimens on slides were stained with Heidenhain's iron hematoxylin or alum carmine. Heidenhain's iron hematoxylin staining was performed as follows: specimens were immersed in 70% ethanol for 1 h, distilled water for 1 h, and 4% ferric ammonium alum for 12h; washed with running water for a half day; stained using Heidenhain's iron hematoxylin from 3–6 h; washed with running water for a half day; immersed in 1 or 2% ammonium ferric ammonium alum until the color was adequate for observation; washed in running water for a half day; dehydrated in graded ethanol series (70%, 90%, 99.5%, and 100%) for one day each; and cleared in creosote for over 3 days. For alum carmine staining, specimens were immersed in 70% ethanol for 1 h and distilled water for 1 h, stained using alum carmine for 3 days, washed with running water for a half day, with 70% ethanol containing 1% HCl until the color was adequate for observations, washed with running water for half a day, dehydrated in graded ethanol series (70%, 90%, 99.5%, and 100%) for one day each, and cleared in xylene for over 3 days. The stained specimens were then mounted in Canada balsam or Entellan new (Merck). Specimens were deposited in the National Museum of Nature and Science (NSMT-PI).

2.3 Morphological observations

Prepared specimens were observed under a light microscope. Illustrations were made using a drawing tube or by tracing the photographs of the specimens. Morphological terms followed Chervy (2002) and Ruhnke (2011) (Fig. 2). Measurements of the following body parts were made directly on the specimens using ocular micrometer or on the drawings using ImageJ 1.49 (Rasband, 1997–2014): total length and maximum width, muscle band widths at anterior and posterior proglottid, scolex length and width, accessory sucker length and width, bothridia length and width, apical sucker length and width, stem length and width,

neck width, anterior region of scolex length and width, testis length and width, cirrus width, cirrus sac length and width, ovary length and width, uterus length and width, vagina width, vitelline follicles length and width, and egg length and width. The number of proglottids and testes were also counted. Specimens were identified as species based on the published keys to the cestodes (Khalil et al., 1994) and every original or revised descriptions.

2.4 DNA extraction, PCR amplification, and DNA sequencing

Total genomic DNA was extracted from a total of 43 species (Table 6) of Phyllobothriidea and its related taxa. In the case that the same species was collected from two or more host species, DNA was obtained from cestode specimen from each host species. DNA extraction was performed using DNeasy Blood & Tissue Kit (Qiagen) following the manufacturer's protocol.

Complete *ssrDNA* and partial *lsrDNA* (D1–D3) were amplified using the primers WormA and WormB, and LSU5 and 1500R, respectively (Waeschenbach et al., 2007; Table 7). PCR amplifications were performed using Premix Taq™ (Ex Taq™ Version 2.0, TaKaRa) or HotStarTaq DNA Polymerase (Qiagen). The total reaction volume was 15 µl containing 7.5 µl of Premix Taq, 0.75 µl of each 0.5 µM primer, and 4.5 µl of Milli-Q water, or 0.075 µl of HotStarTaq DNA Polymerase, 1.5 µl of 10X PCR Buffer, 0.3 µl of 1M dNTP mix, 0.75 µl of each 0.5 µM primer, and 10.125 µl of UPW. Thermal cycling was performed in a Mastercycler® nexus (Eppendorf) or TaKaRa PCR Thermal Cycler Dice® (TaKaRa). PCR reactions were initiated by denaturation at 94°C for 2 min, followed by 40 cycles of the following reactions: denaturation at 94°C for 1 min, annealing at 53°C for 30 s, and extension at 72°C for 2 min, followed by final extension at 72°C for 7 min and completed by preserving the mixture at 10°C. PCR products were loaded on 1.2% agarose gels to confirm amplification and purified by removing excess oligonucleotides using Exo-SAP IT For PCR Product Clean-Up (Affymetrix/USB) following the manufacturer's protocol.

BigDye Terminator version 3.1 Cycle Sequencing Kit (Applied Biosystems) was used for sequencing the purified PCR product. Cycling conditions followed the manufacturer's protocol. PCR and additional sequencing primers (300F, 300R, 930F, 1200F, 1200R, 1270F, 1270R, and 18S-A27 for *ssrDNA* and 300F, ECD2, and 1200R for *lsrDNA*; Table 7) were used. Sequencing was conducted for both strands by Applied Biosystems 3500/3500xL Genetic Analyzer (Applied Biosystems). Contigs were assembled using ATGC5 (GENETYX) or GeneStudio Professional (GeneStudio, Inc.).

2.5 Sequence alignment and tree construction

In the present study, 56 operational taxonomic units (OTUs) from the order Phyllobothriidea and its related taxa, which have not yet been analyzed on a molecular basis, were newly sequenced for the phylogenetic analyses (Table 6). In addition, 55 OTU sequences from GenBank, which were used by Caira et al. (2014), were also analyzed (Table 6). The following two datasets were prepared: Dataset 1 contained a suite of all 111 OTU sequences and Dataset 2 was restricted to 73 OTU sequences, excluding data from the fast-evolving taxa and those from the same species (Table 6, see Results). The sequences of complete *ssrDNA*, partial *lsrDNA* (D1–D3), and their combined sequences were used for analysis.

By the multiple alignments using MAFFT v7.215 (Katoh & Standley, 2013) with the option L-INS-i, 2,322 positions were aligned from 1,770–2,157 bp of *ssrDNA* and 1,549 positions from 1,226–1,390 bp of *lsrDNA* in Dataset 1, whereas 1,905 positions were aligned from 1,816–1,956 bp of *ssrDNA* and 1,396 positions from 1,225–1,331 bp of *lsrDNA* in Dataset 2. To remove unreliably aligned regions, including large gaps, Gblocks v0.91b (Castresana, 2000) was employed with the following parameter settings in Dataset 1: minimum number of sequences for a conserved position = 57, minimum number of sequences for a flank position = 95, maximum number of contiguous non-conserved positions = 8, minimum length of a block = 10, allowed gap positions = all, whereas in dataset 2 the following parameter settings were employed: minimum number of sequences for a conserved position = 37, minimum number of sequences for a flank position = 62, maximum number of contiguous non-conserved positions = 8, minimum length of a block = 10, and allowed gap positions = all. Therefore, 1,859 positions of *ssrDNA* consisting of seven blocks and all 1,386 positions of *lsrDNA* consisting of nine blocks were selected for reliable regions in Dataset 1, whereas all 1,951 positions of *ssrDNA* and all 1,378 positions of *lsrDNA* consisting of two blocks were selected in Database 2.

The phylogenetic trees were constructed using the maximum likelihood (ML) method with IQ-TREE version 1.4.3 (Nguyen et al., 2015) and BI with MrBayes v. 3.2.4 (Ronquist & Huelsenbeck, 2003). The General Time Reversible plus Gamma distributed (GTR + G) model was selected as the best-fitting evolutionary model by IQ-TREE for all analyses. Gaps and missing site characters were treated as unknown characters. In the ML analysis, branch supports with the ultrafast bootstrap (UFBoot) (Minh et al. 2013) were obtained and the Shimodaira-Hasegawa like interpretation of aLRT statistic (SH aLRT) (Guindon et al., 2010) implemented

in the IQ-TREE, UFBoot, and SH aLRT were estimated from 1,000 replicates. SH aLRT was only used to estimate for nodes of orders. In the BI analysis, bayesian posterior probabilities (BPP) were obtained. The Markov-chain Monte Carlo (MCMC) process was run until the average standard deviation of split frequencies became less than 0.01. The MCMC process was run with four chains sampled every 100 generations for 6,500,000 generations for the combined sequence, 1,500,000 generations for *ssrDNA*, and 3,000,000 generations for *lsrDNA* in Dataset 1, whereas 100 generations for 500,000 generations for the combined sequence and *ssrDNA* and 1,500,000 generations for *lsrDNA* were sampled in Dataset 2, and the first 25% trees were discarded as burn-in.

As above mentioned, a total of 12 phylogenetic analyses were conducted and summarized as follows. ML analysis with IQ-TREE in Dataset 1 with a combined sequence of *ssrDNA* and *lsrDNA*, a single *ssrDNA*, and *lsrDNA* (Analyses 1, 2, and 3, respectively) and in Dataset 2 with a combined sequence of *ssrDNA* and *lsrDNA*, a single *ssrDNA*, and a single *lsrDNA* (Analyses 4, 5, and 6, respectively) and BI analysis in Dataset 1 with a combined sequence of *ssrDNA* and *lsrDNA*, a single *ssrDNA*, and a single *lsrDNA* (Analyses 7, 8, and 9, respectively) and in Dataset 2 with a combined sequence of *ssrDNA* and *lsrDNA*, a single *ssrDNA*, and a single *lsrDNA* (Analyses 10, 11, and 12, respectively).

The UFBoot and BPP are shown on each ML and BI tree, respectively. The author interpreted a statistical support as robust when a node was supported by UFBoot ≥ 95 , BPP ≥ 0.98 , or SH aLRT ≥ 80.0 . The members of the order Litobothriidea and Lecanicephalidea were used as outgroups in Analyses 1–3 and 7–9 and those of the order Lecanicephalidea were used as outgroups in Analyses 4–6 and 10–12.

3. Results

3.1 Molecular phylogenetic analysis

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***Calliobothrium shirozame* n. sp.**

Type host: *Mustelus griseus* Pietschmann, 1908 (Carcharhiniiformes: Triakidae).

Type locality: off Chigasaki, Kanagawa Prefecture, Japan.

Site of infection: Spiral intestine.

Distribution: Pacific Ocean— off Chigasaki, Kanagawa Prefecture, Japan (this study); off Iburi, Kochi Prefecture, Japan (this study); off Kochi, Kochi Prefecture, Japan (this study).

Etymology: The species is named for Japanese name of its type host.

Material examined: NSMT-P1 6037 (Holotype: a whole mounted specimen ex *Mu. griseus*, off Chigasaki, Kanagawa Prefecture, Japan, 31. v. 2009); NSMT-P1 6038–6039 (two whole mounted specimens ex *Mu. griseus*, off Chigasaki, Kanagawa Prefecture, Japan); NSMT-P1 6044, 6048 (two whole mounted specimens ex *Mu. griseus*, off Chigasaki, Kanagawa Prefecture, Japan, 31. v. 2009); NSMT-P1 6040–6041, 6051 (four whole mounted specimens ex *Mu. griseus*, off Chigasaki, Kanagawa Prefecture, Japan, vi. 2009); NSMT-P1 6045 (a whole mounted specimen ex *Mu. griseus*, off Chigasaki, Kanagawa Prefecture, Japan, 24. x. 2009); NSMT-P1 6042 (a whole mounted specimen ex *Mu. griseus* (host voucher NSMT-P 111875) off Iburi, Kochi Prefecture, Japan, 4. xii. 2012); NSMT-P1 6043, 6050 (two whole mounted specimens ex *Mu. griseus*, off Iburi, Kochi Prefecture, Japan, 7. xii. 2012); NSMT-P1 6046–6047 (two whole mounted specimens ex *Mu. griseus*, off Kochi, Kochi Prefecture, Japan, 17. i. 2013).

Description (Fig. 19)

Adults

Based on 17 whole mounted adult specimens.

Worms euapolytic 26.7–108.3 (56.4 mm, N=15) long, with maximum width 596–1,800 (1,062, N=15) at level of mature proglottids, comprised of 240–353 (288, N=15) proglottids (Fig. 19A). Scolex 295–562 (405, N=17) long, bears four bothridia and cephalic peduncle (Fig. 19B). Cephalic peduncle 19–119 (64, N=17) long, gradually widening posteriorly, extends beyond posterior margin of bothridium. Bothridium 257–495 (364, N=16, n=58) long, 129–190 (158, N=14, n=44) wide, each with muscular pad 24–148 (65, N=14, n=29) long, 33–190 (111, N=16, n=34) wide, with central accessory sucker 24–71 (40, N=15, n=41) in diameter and 2 lateral accessory suckers 19–57 (32, N=15, n=44) in diameter, bearing 2 pairs of articulated hooks, divided into 3 loculi by 2 transverse septa. Anterior loculus 105–238 (154, N=16, n=59) long, 81–143 (112, N=13, n=42) wide; middle loculus 33–105 (61, N=16, n=59) long, 62–143 (99, N=13, n=42) wide; posterior loculus 38–100 (61, N=16, n=59) long, 52–129 (80, N=13, n=42) wide.

All hooks hollow. Lateral and medial pair of hooks symmetrical, each tip of prongs extending posteriorly (Fig. 19C). Axial hooks more acutely recurved than abaxial hooks. Lateral hook measurements (Fig. 19D): A, 48–93 (68, N=16, n=53); B,

95–148 (117, N=16, n=53); C, 43–67 (50, N=16, n=53); D, 95–148 (119, N=16, n=53); E, 60–98 (77, N=16, n=53); F, 31–52 (43, N=16, n=53). Medial hook measurements (Fig. 14D): A', 48–93 (68, N=16, n=53); B', 93–145 (116, N=16, n=53); C', 38–62 (49, N=16, n=53); D', 95–157 (118, N=16, n=53); E', 57–107 (77, N=16, n=53); F, 36–52 (43, N=16, n=53).

Proglottids craspedote, laciniate (Fig. 19E). Cephalic peduncle with 2 dorsolateral and 2 ventrolateral pairs of lacinations; immature proglottids with 2 dorsolateral and 2 ventrolateral pairs and 1 central lacination on dorsal and ventral surfaces (Fig. 19F), middle lacinations gradually disappearing toward mature proglottids; mature proglottids with 2 dorsolateral and 2 ventrolateral pairs of lacinations. Mature proglottids 576–3,323 (1,975, N=14, n=39) long, 506–1,573 (977, N=14, n=39) wide, becoming longer than wide with maturity. Testes 77–116 (93, N=13, n=36) in total number, 41–143 (85, N=13, n=37) long, 57–157 (100, N=13, n=37) wide, densely arranged, extend from near anterior margin of proglottid to anterior margin of ovary; postvaginal testes 26–43 (33, N=13, n=36) in number. Cirrus-sac spherical, 62–233 (160, N=13, n=35) long, 57–176 (116, N=13, n=35) wide, contains coiled cirrus, covered with small microtriches. Vas deferens medial, coiled lateral and anterior to cirrus-sac, enters the proximal end of cirrus-sac. Ovary posterior, 144–915 (438, N=12, n=39) long, 326–1,223 (708, N=12, n=39) wide, H-shaped in dorsoventral view; ovarian margins digitiform. Genital pores lateral, irregularly alternating, 62–81 (74, N=14, n=39)% from anterior margin of each proglottid. Vagina sinuous, arises from oötype, extends anteriorly along mid-line of proglottid, then laterally along the anterior margin of cirrus-sac, and opens in common genital atrium next to cirrus. Vitellarium follicular, follicles arranged in two lateral fields consisting of multiple columns, interrupted by vagina and cirrus-sac on the poral side. Uterus median, occupies region from oötype to approximately anterior 1/5 of proglottid, ventral to the vagina.

Remarks

Calliobothrium shirozame was distinguished from the congeners by the number and form of lacinations. *Calliobothrium shirozame* has four posterior triangular lacinations in cephalic peduncle, six in immature proglottids and four in mature proglottids whereas *Ca. australis* and *Ca. verticillatum* have four posterior triangular lacinations in cephalic peduncle, six in early immature proglottids, eight in late immature proglottids, and four in mature proglottids (Euzet, 1959; Ivanov &

Brooks, 2002) while *Ca. creeveyae* and *Ca. tylotocephalum* have two posterior triangular lacinations in cephalic peduncle and all proglottids (Alexander, 1963; Butler, 1987).

All *Calliobothrium* species are oioxenous species. Two triakid sharks of the genus *Mustelus*, *Mu. griseus* and *Mu. manazo*, are sympatrically distributed in Japanese waters (Ebert et al., 2013), but *Calliobothrium* species except this new species have been reported only from *Mu. manazo* (Yoshida, 1917; Yamaguti, 1952; Yamaguchi et al., 2000). In this investigation, *Calliobothrium shirozame* was only found in *Mu. griseus*. Yamaguchi et al. (2000) also found no specimen of this new species from *Mu. manazo* in their surveys on parasites of *Mustelus* species in the coastal seas of Japan and Taiwan.

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4. Discussion

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Chapter II

Identification of phyllobothriidean larvae and their development

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General discussion

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Conclusion

The primary objective of this study was to revise the taxonomy of Phyllobothriidea genera based on their molecular phylogeny. According to the molecular analyses, Phyllobothriidea is composed of 17 genera, three of which are newly described. Considerable morphological variation was observed in each genus. The form of the bothridium has traditionally been used as the primary characteristic for the classification of phyllobothriidean genera, but this did not reflect the molecular phylogeny.

Based on these new systematics, the Phyllobothriidea fauna of the coastal seas of Japan were summarized. Phyllobothriidea from Japan are composed of 41 species belonging to 15 genera, which included 14 new species, and 12 species and five genera new to Japan. This suggests that the phyllobothriidean fauna of the coastal seas of Japan is rich in species diversity.

The second objective of this study was to identify plerocercoid larvae using molecular techniques for better understanding of the life cycle of Phyllobothriidea. Seven species of phyllobothriidean plerocercoids were found in teleosts and cephalopods. Four of these were identified as *Pelichnibothrium montaukensis*, *Pe. cf. montaukensis*, *Scyphophyllidium carcharhinus*, and *Vertebraeovicestus dobukasube* based on lsrDNA and mtDNA COI, and their intermediate host species were found to be *Alepisaurus ferox*, *Todarodes pacificus* and *Berryteuthis magister*, *Coryphaena hippurus*, and *Lycodes nakamurae*, respectively. The remaining three species were not identified. These Phyllobothriidea larvae were categorized into three morphological types based on the present observations and information from previous reports.

The evolution of Phyllobothriidea has resulted not only from co-speciation, but more importantly also from host switching events. Such events can readily occur between closely related hosts and also results in high host specificity at the genus level. Furthermore, host switching events may have caused the observed morphological diversification, resulting in phyllobothriidean morphology not reflecting their phylogeny.

Acknowledgements

I am deeply grateful to Dr. Toshihiko Fujita (National Museum of Nature and Science, Tokyo) for his support and guidance throughout my study. I would like to thank Dr. Fumio Tajima and Dr. Rei Ueshima (the University of Tokyo), Dr. Toshiaki Kuramochi and Dr. Tsuyoshi Hosoya (National Museum of Nature and Science, Tokyo), for reviewing the manuscript. I greatly thank to Dr. Hitomi Hirose, Dr. Toru Taniuchi, Dr. Kiyoshi Yoshihara (Nihon University), Dr. Kazuo Ogawa (Meguro Parasitological Museum) for their supporting throughout my study; to Dr. Nobuhiro Mano, Toshiya Shimizu, Akiko Tsuchiya (Kitamura) (Nihon University) for their insightful comments and suggestions.

Many people helped me to collect host specimens: Dr. Tadasu Yamada, Dr. Gento Shinohara, Dr. Kazunori Hasegawa, Dr. Yuko Tajima, Dr. Taiju Kitayama, Dr. Goro Tanifuji, Dr. Satoru Chiba, Dr. Eri Katayama (National Museum of Nature and Science, Tokyo), Dr. Fumihito Tashiro (Kyoto University), Dr. Takahito Kojima, Dr. Noriyuki Takai, Hideaki Takashiro, Genki Obara, Naoki Takamoto, Takashi Tamaru and Keisuke Tanabe, Naoki Tokisaki, Ayaka Hasaka (Nihon University), Ohkawa Toshiyuki, Taniguchi Masao and Saida Hisaki (Kochi Prefecture Fisheries Experiment Station), Dr. Hiromitsu Endo, Naohide Nakayama, Ryo Asaoka, Ryo Misawa (Kochi University), Hiroyuki Irino (Osaka Aquarium KAIYUKAN), Dr. Yuji Ueda (Japan Sea National Fisheries Research Institute), Dr. Kei Sakuma (National Research Institute of Far Seas Fisheries), Dr. Minoru Toda, Dr. Kei Miyamoto (Okinawa Churashima Foundation), Dr. Katsunori Tachihara, Shohei Koizumi (University of the Ryukyus), Dr. Sho Tanaka, Dr. Taku Horie and Asako Mumaguchi (Tokai University), Dr. Masahito Higuchi (Niigata Prefecture Fisheries Experiment Station), and Nobuaki Akao (Tokyo Medical and Dental University).

The materials for this study were collected by R/V Soyo-Maru of the Fisheries Research Agency, F/S Chogoro-Maru of Kanaya Fishing Port, F/S Marukin-Maru of Nagai Fishing Port, Tokuami Inc., Akazawa Gyogyo Inc., T/S Suzaki II of the Nihon University, F/S Shojin-Maru of Heda Fishing Port, T/S Tanshu-Maru of the Hyogo Prefectural Kasumi Fishery High School, Kochi Prefectural Fisheries Experimental Station, Mimase Fishing Port, Tosasaga Fishing Port, Iburi Triangular Set Nets Cooperative, Tosashimizu Fisheries Cooperative, Tosashimizu Fisheries Cooperative, Yomitan Fisheries Cooperative, Yaeyama Fisheries Cooperative, Fishing boat Aippara II.

I would like to thank the following scientists for their comments to my study

and help in collecting samples: Dr. Keiichi Matsuura, Dr. Masanori Nakae, Dr. Takuya Kiyoshi, Dr. Hiromori Komatsu, Dr. Takashi Sato, Dr. Shinpei Ohhashi, Dr. Takuma Haga (National Museum of Nature and Science, Tokyo), Dr. Masaaki Machida, Dr. Jun Araki and Dr. Takashi Iwaki (Meguro Parasitological Museum), Dr. Sho Shirakashi (Kindai University), Dr. Tetsuya Yanagida (Yamaguchi University), Dr. Yuzo Ota (Lake Biwa Museum), Dr. Daisuke Ueno (University of Florida), Dr. Kenta Suda (Okabe Co. Ltd.). Thanks are also due to my colleagues: Dr. Masato Hirose (the University of Tokyo), Dr. Masanori Okanishi (Ibaraki University), Junko Inoue, Mayumi Masuda, Mikihiro Arai, Ikumi Tahara, Takato Izumi, Akito Ogawa (the University of Tokyo), Woo Sau Pinn (Hokkaido University), Suguru Ujino (Fukuoka Prefectural Office), Dr. Noko Kuze (National Museum of Nature and Science, Tokyo), Akito Ogawa (Toho University), Dr. Yoshiaki Ishida, Dr. Seiichi Irimura, Hiromi Takeuchi, Tomoko Iida (National Museum of Nature and Science, Tokyo), Rie Tajiri (Waseda University). The authors would like to thank Enago (www.enago.jp) for the English language review.

This study is partly supported by the project studies conducted by the National Museum of Nature and Science, “Integrated research on biodiversity of interspecies relationships”, “Research on the Earth’s surface processes and biota in and near the Sea of Japan” and “Biodiversity hot spots in Japan”.

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Table 1 Families of Tetrephyllidea in major taxonomical works before Caira et al. (2014). Circles denote that the family was included in Tetrephyllidea in each paper. Serendipidae was most recently a family of Tetrephyllidea, no comprehensive taxonomical works of tetrephyllidean families exist since the establishment of this family.

Family	Braun (1900)	Meggitt (1924)	Southwell (1925)	Fuhrmann (1931)	Joyeux & Baer (1936)	Hyman (1951)	Wardle & McLeod (1952)	Euzet (1959)	Yamaguti (1959)	Joyeux & Baer (1961)	Schmidt (1986)	Euzet (1994)
Ichthyotaeniidae Ariola, 1899	x	x			x							
Onchobothriidae Braun, 1900	x	x	x	x	x	x	x	x	x	x	x	x
Phyllobothriidae Braun, 1900	x	x	x	x	x	x	x	x	x	x	x	x
Lecanicephalidae Braun, 1900	x	x		x				x		x		
Monticelliidae La Rue, 1911				x								
Proteocephalidae La Rue, 1911				x								
Polypocephalidae Miggitt, 1924		x										
Cephalobothriidae Pintner, 1928				x	x			x		x		
Discocephalidae Pintner, 1928				x								
Disculicipitidae Joyeux & Baer, 1935					x			x		x		x
Gastrolecithidae Euzet, 1955								x		x		
Phoreiobothriidae Baer & Euzet 1955								x		x		
Prosobothriidae Baer & Euzet, 1955								x		x		x
Maccallumiellidae Yamaguti, 1959									x			
Triloculariidae Yamaguti, 1959									x		x	
Dioecotaeniidae Schmidt, 1969												x
Litobothriidae Dailey, 1969												x
Cathetocephalidae Dailey & Overstreet, 1973											x	x
Chimaerocestidae Williams & Bray, 1984												x
Serendipidae Brooks & Barriga, 1995												

Table 2 Phyllobothriidean species reported in coastal seas of Japan.

Species	Host species	Location	Reference
<i>Phyllobothrium</i>			
<i>Phyllobothrium lactuca</i>	<i>Mustelus manazo</i>	Hamajima (Mie Pref.)	Yoshida (1917)
		Sea of Japan	Yamaguti (1952)
<i>Phyllobothrium serratum</i>	<i>Triakis scyllium</i>	Hamajima (Mie Pref.)	Yamaguti (1952)
<i>Phyllobothrium</i> sp.	<i>Isurus glaucus</i>	Pacific Ocean	Yamaguti (1934)
<i>Phyllobothrium</i> sp.	<i>Mustelus manazo</i>	Coatal seas of Japan	Yamaguchi et al. (2000)
<i>incertae sedis</i>			
<i>Phyllobothrium biacetabulatum</i>	<i>Rhinobatus schlegeli</i>	Inland sea (Okayama Pref.)	Yamaguti (1960)
<i>Phyllobothrium dasybati</i>	<i>Dasrbatus akajei</i>	Pacific coast	Yamaguti (1934)
<i>Phyllobothrium marginatum</i>	<i>Squatina japonica</i>	Toyama Bay	Yamaguti (1934)
<i>Phyllobothrium loculatum</i>	<i>Heterodontus zebra</i>	East China Sea	Yamaguti (1952)
<i>Phyllobothrium squali</i>	<i>Squalus sucklii</i>	Onahama (Fukushima Pref.)	Yamaguti (1952)
<i>Orygmatobothrium</i>			
<i>Orygmatobothrium musteli</i>	<i>Mustelus manazo</i>	Hamajima (Mie Pref.)	Yamaguti (1952)
		Pacific	Yamaguti (1952)
		Sea of Japan	Yamaguti (1952)
<i>species inquirenda</i>			
<i>Orygmatobothrium velaentum</i>	<i>Mustelus manazo</i>	Hiroshima	Yoshida (1917)
<i>Orygmatobothrium</i> sp.	<i>Mustelus manazo</i>	Coatal seas of Japan	Yamaguchi et al. (2000)
<i>Paraorygmatobothrium</i>			
<i>Paraorygmatobothrium angustum</i>	<i>Triakis scyllium</i>	Misaki (Kanagawa Pref.)	Yoshida (1917)
<i>Paraorygmatobothrium filiforme</i>	<i>Alopias vulpinus</i>	Pacific Ocean	Yamaguti (1934)
<i>Paraorygmatobothrium tiacis</i>	<i>Triakis scyllium</i>	Pacific Ocean	Yamaguti (1952)
<i>Paraorygmatobothrium prionacis</i>	<i>Prionace glauca</i>	Pacific Ocean	Yamaguti (1934)
<i>Marsupiobothrium</i>			
<i>Marsupiobothrium alopias</i>	<i>Alopias vulpinus</i>	Pacific Ocean	Yamaguti (1934)

Table 3 Genera reported in Phyllobothriidae so far and their status (Ruhnke, 2011; Ruhnke & Workman, 2013). Genera included in Phyllobothriidea in Caira et al. (2014) are shown in bold face.

Order	Family	Genus
Tetraphyllidea		
	Phyllobothriidae	<i>Phyllobothrium</i> van Beneden, 1849
		<i>Alexandercestus</i> Ruhnke & Workman, 2013
		<i>Bibursibothrium</i> McKenzie & Caira, 1998
		<i>Calyptribothrium</i> Monticelli, 1893
		<i>Cardiobothrium</i> McKenzie & Caira, 1998
		<i>Clistobothrium</i> Dailey & Vogelbein, 1990
		<i>Crossobothrium</i> Linton, 1889
		<i>Marsupiobothrium</i> Yamaguti, 1952
		<i>Doliobothrium</i> Malek, Caira & Ruhnke, 2010
		<i>Flexibothrium</i> McKenzie & Caira, 1998
		<i>Monorygma</i> Diesing, 1863
		<i>Nandocestus</i> Reyda, 2008
		<i>Orectolobicestus</i> Ruhnke, Caira & Carpenter, 2006
		<i>Orygmatobothrium</i> Diesing, 1863
		<i>Paraorygmatobothrium</i> Ruhnke, 1994
		<i>Ruhnkecestus</i> Caira & Durkin, 2006
		<i>Scyphophyllidium</i> Woodl&, 1927
		<i>Thysanocephalum</i> Linton, 1889
		<i>nomina dubia</i>
		<i>Aacobothrium</i> Mola, 1907
		<i>Dittocephalus</i> Parona, 1887
		<i>Hoaleshwaria</i> Shinde & Chincholikar, 1975
		<i>Phanobothrium</i> Mola, 1907
		<i>Shindeobothrium</i> Shinde & Chincholikar, 1975
		<i>nomen ad interim</i>
		<i>Phyllobothrideum</i> Olsson, 1866
		<i>inquirenda</i>
		<i>Bilocularia</i> Obersteiner, 1914
		<i>Biporophyllaeus</i> Subramaniam, 1939
		<i>Cyatocotyle</i> Mola, 1908
		<i>Kowsalyabothrium</i> Muralidhar, Shinde & Jadhav, 1987
		<i>Maccallumiella</i> Yamaguti, 1959
		<i>Mastacembellophyllaeus</i> Shinde & Chincholikar, 1977
		<i>Pillersium</i> Southwell, 1927
		<i>Pithophorus</i> Southwell, 1925
		<i>Polipobothrium</i> Mola, 1908
	Serendipidae	<i>Duplicibothrium</i> Williams & Campbell, 1978
		<i>Glyphobothrium</i> Williams & Campbell, 1977
		<i>inquirenda</i>
		<i>Myliobatibothrium</i> Shinde & Mohekar, 1983
		<i>Tiarabothrium</i> Shipley & Hornell, 1906

Table 3 continued.

Order	Family	Genus
	<i>incertae sedis</i>	<i>Anindobothrium</i> Marques, Brooks & Lasso, 2001 <i>Anthobothrium</i> van Beneden, 1850 <i>Carpobothrium</i> Shipley & Hornell, 1906 <i>Caulobothrium</i> Baer, 1948 <i>Caulopatera</i> Cutmore, Bennett & Cribb, 2010 <i>Ceratobothrium</i> Monticelli, 1892 <i>Chimaerocestos</i> Williams & Bray, 1984 <i>Dinobothrium</i> van Beneden, 1889 syn. <i>Diplobothrium</i> van Beneden, 1889 syn. <i>Reesium</i> Euzet, 1955 <i>Gastrolecithus</i> Yamaguti, 1952 <i>Guidus</i> Ivanov, 2006 <i>Mixophyllobothrium</i> Shinde & Chincholikar, 1980 <i>Myzocephalus</i> Shipley & Hornell, 1906 <i>Myzophyllobothrium</i> Shipley & Hornell, 1906 <i>Pelichnibothrium</i> Monticelli, 1889 syn. <i>Prionacestus</i> Mete & Euzet, 1996 <i>Rhoptrobothrium</i> Shipley & Hornell, 1906 <i>Trilocularia</i> Olsson, 1867 syn. <i>Urogonoporus</i> Lühe, 1902 <i>Zyxibothrium</i> Hayden & Campbell, 1981
	Rhinebothriidea	<i>Anthocephalum</i> Linton, 1890 <i>Biotobothrium</i> Tan, Zhou & Yang, 2009 <i>Clydonobothrium</i> Euzet, 1959 <i>Echeneibothrium</i> van Beneden, 1850 <i>Escherbothrium</i> Berman & Brooks, 1994 <i>Notomegarhynchus</i> Ivanov & Campbell, 2002 <i>Pararhinebothroides</i> Zamparo, Brooks & Barriga, 1999 <i>Pentaloculum</i> Alexander, 1963 <i>Phormobothrium</i> Alexander, 1963 <i>Pseudanthobothrium</i> Baer, 1956 <i>Rhabdotobothrium</i> Euzet, 1953 <i>Rhinebothrium</i> Linton, 1890 <i>Rhinebothroides</i> Mayes, Brooks & Thorson, 1981 <i>Rhodobothrium</i> Linton, 1889 syn. <i>Inermiphyllidium</i> Riser, 1955 syn. <i>Sphaerobothrium</i> Euzet, 1959 syn. <i>Proboscidosaccus</i> Gallien, 1949 <i>Scalithrium</i> Ball, Neifar, & Euzet, 2003 <i>Spongiobothrium</i> Linton, 1889 <i>Tritaphros</i> Lönnberg, 1889 <i>inquirenda</i> <i>Shindeiobothrium</i> Jadhav, Shinde & Deshmukh, 1981

Table 4 Sampling data. See Fig. 1 for positions of the stations.

Station	Locality	Depth	Fishing gear	Remarks
1	off Erimo, Hokkaido Prefecture	-	set net	Bought from Onsenichiba Inc.
2	off Ooma, Aomori Prefecture	ca. 200 m	long line	Bought from Tamukaisyoten Inc.
3	off Sado Island, Niigata Prefecture	-	set net	Collected by Kamo-Suisan Set Nets Cooperative
4	off Ibaraki, Ibaraki Prefecture	ca. 200 m	crab cage	Collected using R/V Soyo-Maru of the Fisheries Research
5	Ooarai, Ibaraki Prefecture	-	-	A whale stranded on the beach
6	off Toyama, Toyama Prefecture	-	set net	Tapeworms specimens were provided by Tokyo Medical and Dental University.
7	off Kanaya, Chiba Prefecture	ca. 160–500 m	gill net	Collected using F/S Tyogoro-Maru of Kanaya Fishing Port
8	off Miura, Kanagawa Prefecture	ca. 160–500 m	gill net	Collected using F/S Marukin-Maru of Nagai Fishing Port
9	off Chigasaki, Kanagawa Prefecture	ca. 1–10 m	dragnet	Collected by Tokunami Inc.
10	off Izu-Ooshima Island, Tokyo Metropolitan	ca. 200 m	long-line	
11	off Akazawa, Shizuoka Prefecture	ca. 25 m	set net	Collected by Akazawa Gyogyo Inc.
12	off Shimoda, Shizuoka Prefecture	-	angling	
13	off Shimoda, Shizuoka Prefecture	ca. 30 m	long line	Collected using T/S Suzuki II of the Nihon University
14	off Toshima Island, Tokyo Metropolitan	ca. 30 m	long line	Collected using T/S Suzuki II of the Nihon University
15	off Yui, Shizuoka Prefecture	-	set net	
16	off Heda, Shizuoka Prefecture	ca. 140–690 m	bottom trawl	Collected using F/S Shoujin-Maru of Heda Fishing Port
17	off Irouzaki, Shizuoka Prefecture	ca. 30 m	long line	Collected using T/S Suzuki II of the Nihon University
18	off Obama, Fukui Prefecture	-	set net	
19	off Maizuru, Kyoto Prefecture	-	-	Bought from Mizushima-Suisan Inc.
20	off Oku Islands, Shimane Prefecture	ca. 200–400 m	otter trawl	Collected using T/S Tansyu-Maru of the Hyogo Prefectural
21	off Kochi, Kochi Prefecture	ca. 55–70 m	beam trawl	Kasumi Fishery High School
22	off Mimase, Kochi Prefecture	-	trawl	Collected by Kochi
23	off Tosasaga, Kochi Prefecture	-	trawl	Prefectural Fisheries Experimental Station
24	off Iburi and Kubotsu, Kochi Prefecture	ca. 25 m	set net	Landed at Mimase Fishing Port
25	off Kubotsu, Kochi Prefecture	ca. 25 m	set net	Landed at Tosasaga Fishing Port
				Collected by Iburi Triangular Set Nets Cooperative
				Collected by Kubotsu Triangular Set Nets Cooperative

Table 4 continued.

Station	Locality	Depth	Fishing gear	Remarks
26	off Tosazhimizu, Kochi Prefecture	-	long line	Collected by Tosashimizu Fisheries Cooperative
27	Minamisanaka, Kagoshima Prefecture	-	-	A whale stranded on the beach
28	off Yomotan, Okinawa Prefecture	ca. 25 m	set net	Collected by Yomitan Fisheries Cooperative
29	off Yaeyama Islands, Okinawa, Prefecture	-	long line	Collected by Yaeyama Fisheries Cooperative
30	off Ogasawara Islands, Tokyo Metropolitan	-	hook and line	Using by F/B Aippara II or from land
31	off Ogasawara Islands, Tokyo Metropolitan	-	long line	Collected by R/V Koyo of the Tokyo Metropolitan Islands Area Research and Development Center of Agriculture

Table 5 List of the examined host specimens of elasmobranchs, teleosts and cetaceans. N means the number of host specimens and the number of measured specimens is shown in parentheses if all specimens were not measured. Length was measured by total length except some batoids of Dasyatidae, Gymnuridae, and Myliobatidae by disk length (marked by asterisks). Weight is shown by gram for small specimens (indicated by g).

Host species	N	Length (mm) Range	Weight (kg) Range
Elasmobranchii			
Galeomorphii			
Heterodontidae			
<i>Heterodontus japonicus</i>	19	326 – 1031	0.3 – 5.3
Orectolobidae			
<i>Orectolobus japonicus</i>	1	1071	10.2
Ginglymostomatidae			
<i>Nebrius ferrugineus</i>	1	2960	141.0
Mitsukurinidae			
<i>Mitsukurina owstoni</i>	8	1050 – 3600	1.7 – 167.0
Odontaspidae			
<i>Odontaspis ferox</i>	1	no data	no data
Megachasmidae			
<i>Megachasma pelagios</i>	1	4460	677.0
Lamnidae			
<i>Carcharodon carcharias</i>	1	4070	no data
<i>Isurus oxyrinchus</i>	3 (1)	1500	26.1
Scyliorhinidae			
<i>Apristurus macrorhynchys</i>	17	360 – 672	97 – 786 g
<i>Cephaloscyllium umbratile</i>	60	213 – 1022	0.03 – 8.6
<i>Galeus eastmani</i>	6	342 – 699	0.09 – 1.3
<i>Galeus nipponensis</i>	18	375 – 699	0.1 – 1.5
<i>Parmaturus pilosus</i>	1	622	860 g
<i>Scyliorhinus torazame</i>	17	390 – 489	194 – 490 g
Proscylliidae			
<i>Proscyllium habereri</i>	3	572 – 704	0.4 – 1.0
Triakidae			
<i>Mustelus manazo</i>	69	300 – 935	0.03 – 2.8
<i>Mustelus griseus</i>	20	460 – 880	0.3 – 2.9
<i>Triakis scyllium</i>	12	245 – 992	0.2 – 3.6
<i>Hemitriakis japonica</i>	6	750 – 954	1.3 – 4.1

Table 5 continued.

Table 3. Continued.

Host species	N	Length (mm)	Weight (kg)
		Range	Range
Carcharhinidae			
<i>Carcharhinus albimarginatus</i>	4	1380 – 1560	18.0 – 25.0
<i>Carcharhinus brevipinna</i>	23	700 – 1020	1.3 – 5.5
<i>Carcharhinus dussumieri</i>	4	556 – 730	1.0 – 2.1
<i>Carcharhinus galapagensis</i>	9	790 – 1910	2.7 – 36.1
<i>Carcharhinus leucas</i>	1	2650	147.0
<i>Carcharhinus longimanus</i>	1	1730	32.0
<i>Carcharhinus limbatus</i>	2	1430 – 2090	1.1 – 1.5
<i>Carcharhinus plumbeus</i>	4	1120 – 1420	8.0 – 15.0
<i>Galeocerdo cuvier</i>	6	1130 – 2990	5.4 – 158.0
<i>Loxodon macrorhinus</i>	2	710 – 950	1.0 – 3.0
<i>Prionace glauca</i>	3	2080 – 2690	8.3 – 29.0
<i>Triaenodon obesus</i>	2	1265 – 1320	10.7 – 17.8
Sphyrnidae			
<i>Sphyrna zygaena</i>	5	933 – 1775	4.0 – 24.5
Squalomorphi			
Chlamydoselachidae			
<i>Chlamydoselachus anguineus</i>	18	1005 – 1485	1.7 – 6.7
Hexanchidae			
<i>Heptranchias perlo</i>	30	682 – 1150	0.9 – 5.9
<i>Hexanchus griseus</i>	1	4020	505.5
Echinorhinidae			
<i>Echinorhinus cookei</i>	1	no data	no data
Dalatiidae			
<i>Dalatias licha</i>	52	281 – 1542	0.14 – 29.1
Etmopteridae			
<i>Etmopterus brachyurus</i>	9	241 – 467	39 – 400 g
<i>Etmopterus lucifer</i>	19	284 – 430	70 – 274 g
<i>Etmopterus molleri</i>	17	202 – 385	20 – 179 g
<i>Etmopterus pusillus</i>	16	168 – 288	12 – 82 g
<i>Etmopterus unicolor</i>	1	520	606 g
Centrophoridae			
<i>Deania calcea</i>	46	315 – 1170	0.09 – 8.0
<i>Deania hystricosa</i>	34	324 – 1230	0.12 – 10.4
<i>Centrophorus squamosus</i>	7	461 – 1270	0.42 – 13.0
<i>Centrophorus tessellatus</i>	1	618	1.0
<i>Centrophorus</i> sp. 1	1	1220	12.7
<i>Centrophorus</i> sp. 2	1	895	3.7
Squalidae			
<i>Cirrhigaleus barbifer</i>	1	796	4.3
<i>Squalus acanthias</i>	10	745 – 840	1.6 – 2.2
<i>Squalus brevirostris</i>	69	510 – 880	0.6 – 3.6
<i>Squalus japonicus</i>	18	323 – 690	0.1 – 1.4
<i>Squalus mitsukurii</i>	72	348 – 1090	0.2 – 8.4

Table 5 continued.

Host species	N	Length (mm)	Weight (kg)
		Range	Range
Squatinae			
<i>Squatina japonica</i>	5	588 – 840	1.7 – 6.0
<i>Squatina nebulosa</i>	3	484 – 1490	1.0 – 23.1
Pristiophoridae			
<i>Pristiophorus japonicus</i>	1	1010	1.3
Batoidea			
Rhynchobatidae			
<i>Rhynchobatus djiddensis</i>	1	2410	8.4
Rhinobatidae			
<i>Rhinobatos schlegelii</i>	16	753 – 955	1.1 – 3.4
<i>Rhinobatos hynnicephalus</i>	1	660	1.1
Platyrrhinidae			
<i>Platyrrhina tangi</i>	3	145 – 537	0.02 – 1.1
Narcinidae			
<i>Narke japonica</i>	18	104 – 351	20 – 750 g
Torpedinidae			
<i>Torpedo tokionis</i>	2	530 – 629	2.3 – 4.0
Arhynchobatidae			
<i>Bathyrhaja smirnovi</i>	6	517 – 1030	1.0 – 9.8
Rajiidae			
<i>Amblyrhaja badia</i>	1	922	6.5
<i>Dipterus macrocauda</i>	1	1075	7.6
<i>Okamejei kenojei</i>	2	360 – 373	200 – 460 g
<i>Okamejei schmidtii</i>	1	473	550 g
<i>Raja pulchra</i>	17	648 – 885	0.5 – 6.2
Urolophidae			
<i>Urolophus aurantiacus</i>	3	275 – 362	220 – 534 g
Dasyatidae			
<i>Dasyatis akajei</i>	33	135 – 791*	0.07 – 23.3
<i>Dasyatis izuensis</i>	5	162 – 368*	0.2 – 3.0
<i>Dasyatis matsubarae</i>	2	376 – 380*	1.5 – 1.6
<i>Taeniura meyeni</i>	5 (3)	1180 – 1560	24.8 – 77.0
<i>Pleroplatytrygon violacea</i>	1	585*	6.0
Gymnuridae			
<i>Gymnura japonica</i>	7	324 – 1045*	0.3 – 10.0
Myliobatidae			
<i>Myliobatis tobijei</i>	25	220 – 405*	0.2 – 1.4
Neopterygii			
Alepisauridae			
<i>Alepisaurus ferox</i>	4	905 – 1050	840 – 1360
Salmonidae			
<i>Oncorhynchus keta</i>	1	630	2630
Mammalia			
Delphinidae			
<i>Stenella coeruleoalba</i>	3 (1)	2340	133

Table 6 List of the specimens of two Dataset for molecular phylogenetic analyses. Systematic positions and scientific names of the specimens from GeneBank follow Caira et al. (2014). Type species are shown in bold face. Specimens marked by single asterisk are undescribed species reported by Caira et al. (2014), and specimens marked by double asterisks are unidentified species. Double quotation denotes species of undescribed genus reported by previous papers. Abbreviations of country names in locality: AUS; Australia, CAN; Canada, CHL; Chile, DNK; Denmark, GBR; United Kingdom, JAP; Japan, NZL; New Zealand, MEX; Mexico, MYS; Malaysia, PRY; Paraguay, PEL; Peru, ROU; Romania, SEN; Senegal, USA; U.S.A.

Specimens	Genotype	Host species	Locality	Accession Number		Dataset	
				ssrDNA	lstrDNA	1	2
Phyllobothriidea							
<i>Phyllobothrium</i> cf. <i>lactuca</i>		<i>Mustelus mento</i>	Puerto Montt, CHL	KF685770	KF685845	x	x
<i>Phyllobothrium biacetabulatum</i>		<i>Rhinobatos schlegelii</i>	off Chigasaki, JAP	This study	This study	x	
<i>Phyllobothrium delphini</i>		<i>Stenella coeruleoalba</i>	Minami-satsuma, JAP	This study	This study	x	x
<i>Phyllobothrium hoshizame</i> n. sp.		<i>Mustelus manazo</i>	off Kanaya, JAP	This study	This study	x	x
<i>Phyllobothrium serratum</i>		<i>Triakis scyllium</i>	off Shimoda, JAP	This study	This study	x	x
<i>Phyllobothrium squali</i>	1	<i>Squalus japonicus</i>	off Kanaya, JAP	This study	This study	x	
	2	<i>Squalus mitsukurii</i>	off Kanaya, JAP	This study	This study	x	x
	3	<i>Squalus japonicus</i>	off Mimase, JAP	This study	This study	x	
	4	<i>Squalus mitsukurii</i>	off Kanaya, JAP	This study	This study	x	
	5	<i>Squalus acanthias</i>	Rhode Island, USA	KF685897	KF685846	x	
<i>Calypotrobothrium</i> sp. *		<i>Torpedo nobiliana</i>	Rhode Island, USA	KF685754	KF685848	x	x
<i>Calypotrobothrium</i> sp. 1**		<i>Torpedo tokionis</i>	off Heda, JAP	This study	This study	x	x
<i>Chimaerocesost</i> sp. 1*		<i>Rhinochimaera pacifica</i>	Chatham Rise, NZL	KF685758	KF685827	x	x
<i>Chimaerocesost</i> sp. 2*		<i>Rhinochimaera pacifica</i>	Chatham Rise, NZL	KF685882	KF685850	x	x
<i>Marsupiobothrium gobelinus</i>		<i>Mitsukurina owstoni</i>	off Kanaya, JAP	This study	This study	x	x
<i>Marsupiobothrium</i> sp.**		<i>Alopias pelagicus</i>	Santa Rosalia, MEX	KF685771	KF685821	x	x
<i>Nandocesostus guariticus</i>		<i>Paratrygon aiereba</i>	Mouth of Rio Shilve, PER	KF685888	KF685817	x	x
<i>Orectolobicesstus tyleri</i>		<i>Chiloscyllium punctatum</i>	Mukah, Borneo, MYS	KF685890	KF685819	x	x
<i>Orectolobicesstus randyi</i>		<i>Chiloscyllium hasseltii</i>	Mukah, Borneo, MYS	KF685767	KF685820	x	x
<i>Orygmatobothrium musteli</i>	1	<i>Mustelus manazo</i>	off Heda, JAP	This study	This study	x	x
	2	<i>Mustelus griseus</i>	off Chigasaki, JAP	This study	This study	x	
<i>Orygmatobothrium</i> cf. <i>musteli</i> 1		<i>Mustelus mustelus</i>	Soumbédioune, SEN	KF685768	KF685815	x	
<i>Orygmatobothrium</i> cf. <i>musteli</i> 2		<i>Mustelus mustelus</i>	Ouakam, SEN	KF685891	KF685816	x	x

Table 6 continued.

Specimens	Genotype	Host species	Locality	Accession Number		Dataset	
				ssrDNA	lstrDNA	1	2
<i>Paraorygmatobothrium prionacis</i>		<i>Prionace glauca</i>	Montauk, New York, USA	KF685892	KF685818	x	x
<i>Paraorygmatobothrium exiguum</i>		<i>Prionace glauca</i>	off Tosashimizu, JAP	This study	This study	x	
<i>Paraorygmatobothrium paulum</i>		<i>Alopias vulpinus</i>	Montauk, New York, USA	KF685769	KF685822	x	x
<i>Paraorygmatobothrium triacis</i>		<i>Galeocerdo cuvier</i>	off Iburi, JAP	This study	This study	x	x
<i>Paraorygmatobothrium</i> sp. 1		<i>Triakis scyllium</i>	off Shimoda, JAP	This study	This study	x	x
		<i>Carcharhinus plumbeus</i>	off Yaeyama, JAP	This study	This study	x	x
		<i>Carcharhinus galapagensis</i>	off Ogasawara Is., JAP	This study	This study	x	
<i>Paraorygmatobothrium</i> sp. 2		<i>Hemitriakis japonica</i>	off Shimoda, JAP	This study	This study	x	x
<i>Paraorygmatobothrium</i> sp. 3		<i>Trienodon obesus</i>	off Ogasawara Is., JAP	This study	This study	x	x
		<i>Carcharhinus galapagensis</i>	off Ogasawara Is., JAP	This study	This study	x	
<i>Paraorygmatobothrium</i> sp. 4		<i>Carcharhinus brevipinna</i>	off Iburi, JAP	This study	This study	x	x
<i>Paraorygmatobothrium</i> sp. 5		<i>Carcharhinus limbatus</i>	off Yaeyama, JAP	This study	This study	x	x
<i>Paraorygmatobothrium</i> sp. 6		<i>Trienodon obesus</i>	off Ogasawara Is., JAP	This study	This study	x	x
<i>Paraorygmatobothrium</i> sp. 7		<i>Carcharhinus longimanus</i>	off Yaeyama, JAP	This study	This study	x	x
<i>Paraorygmatobothrium</i> sp. 8		<i>Carcharhinus limbatus</i>	off Yaeyama, JAP	This study	This study	x	x
		<i>Carcharhinus leucas</i>	off Yaeyama, JAP	This study	This study	x	
		<i>Carcharhinus galapagensis</i>	off Ogasawara Is., JAP	This study	This study	x	
<i>Paraorygmatobothrium</i> sp. 9		<i>Carcharhinus dussumieri</i>	off Ogasawara Is., JAP	This study	This study	x	
<i>Paraorygmatobothrium</i> sp. 10		<i>Carcharhinus brevipinna</i>	off Tosasaga, JAP	This study	This study	x	x
		<i>Carcharhinus galapagensis</i>	off Iburi, JAP	This study	This study	x	
<i>Ruhnkecestus latipi</i>		<i>Carcharhinus galapagensis</i>	off Ogasawara Is., JAP	This study	This study	x	
<i>Scyphophyllidium</i> cf. <i>giganteum</i>		<i>Scoliodon macrorhynchus</i>	Mukah, Borneo, MYS	KF685900	KF685853	x	x
<i>Thysanocephalum thysanocephalum</i>		<i>Galeorhinus galeus</i>	Chatham Rise, NZL	KF685901	KF685854	x	x
<i>Thysanocephalum crispum</i>		<i>Galeocerdo cuvier</i>	off Yaeyama, JAP	This study	This study	x	x
<i>Vertebraeovicescus dobukasube</i> n. sp.		<i>Galeocerdo cuvier</i>	off Iburi, JAP	KF685902	KF685823	x	
"Phyllbothriidea gen. sp."*		<i>Bathyraja smirnovi</i>	off Oki Is., JAP	This study	This study	x	x
		<i>Sphyrna lewini</i>	Straits of Florida, USA	KF685889	KF685814	x	x

Table 6 continued.

Specimens	Genotype	Host species	Locality	Accession Number		Dataset	
				ssrDNA	1stDNA	1	2
Tetraphyllidea							
<i>Anthobothrium caseyi</i>		<i>Prionace glauca</i>	Montauk, New York, USA	KF685879	KF685805	x	
		<i>Prionace glauca</i>	off Tosashimizu, JAP	This study	This study		x
<i>Anthobothrium</i> sp. 1 *		<i>Carcharhinus tilstoni</i>	Wessel Is., AUS	KF685752	KF685806	x	
<i>Anthobothrium</i> sp. 2		<i>Carcharhinus brevipinna</i>	off Iburi, JAP	This study	This study	x	
<i>Anthobothrium</i> sp. 3		<i>Carcharhinus plumbeus</i>	off Yaeyama, JAP	This study	This study	x	
<i>Anthobothrium</i> sp. 4		<i>Carcharhinus brevipinna</i>	off Iburi, JAP	This study	This study	x	
<i>Anthobothrium</i> sp. 5		<i>Carcharhinus dussumieri</i>	off Tosasaga, JAP	This study	This study	x	
<i>Alexandercestus oomejirozame</i> n. sp.		<i>Carcharhinus leucas</i>	off Yaeyama, JAP	This study	This study	x	x
<i>Balanobothrium</i> sp.*		<i>Stegostoma fasciatum</i>	Mukah, Borneo, MYS	KF685880	KF685802	x	x
<i>Bilocularia hyperapolytica</i>		<i>Dalatias licha</i>	off Kanaya, JAP	This study	This study	x	x
<i>Calliobothrium</i> cf. <i>verticillatum</i>		<i>Mustelus canis</i>	Old Lyme, USA	KF685753	KF685812	x	x
<i>Calliobothrium shirozame</i> n. sp.		<i>Mustelus griseus</i>	off Shimoda, JAP	This study	This study	x	x
<i>Ceratobothrium xanthocephalum</i>		<i>Isurus oxyrinchus</i>	Montauk, New York, USA	KF685756	KF685849	x	x
<i>Ceratobothrium</i> sp. 1		<i>Carcharodon carcharias</i>	off Iburi, JAP	This study	This study	x	x
<i>Clistobothrium carcharodoni</i>		<i>Carcharodon carcharias</i>	off Iburi, JAP	This study	This study	x	x
<i>Clistobothrium montaukensis</i>	1	<i>Isurus oxyrinchus</i>	Montauk, New York, USA	EF095259	AF286996	x	
	2	<i>Isurus oxyrinchus</i>	off Toshima I., JAP	This study	This study	x	x
<i>Clistobothrium tumidum</i>		<i>Carcharodon carcharias</i>	off Iburi, JAP	This study	This study	x	x
<i>Crossobothrium laciniatum</i>		<i>Hexanchus griseus</i>	Puerto Montt, CHL	KF685883	KF685824	x	x
<i>Crossobothrium campanulatum</i>		<i>Hexanchus griseus</i>	off Sado I., JAP	This study	This study	x	x
<i>Crossobothrium dohrnii</i>		<i>Heptranchias perlo</i>	off Kanaya, JAP	This study	This study	x	x
<i>Crossobothrium</i> cf. <i>dohrnii</i>		<i>Heptranchias perlo</i>	Gulf of Mexico	KF685759	KF685847	x	
<i>Megalonchos shawae</i>		<i>Hemipristis elongata</i>	Wessel Is., AUS	KF685764	KF685807	x	
<i>Monorygma chamydoselachii</i>		<i>Chlamydoselachus anguineus</i>	off Kanaya, JAP	This study	This study	x	x
<i>Monorygma grimaldii</i>		<i>Stenella coeruleoalba</i>	Minami-satsuma, JAP	This study	This study	x	x

Table 6 continued.

Specimens	Genotype	Host species	Locality	Accession Number		Dataset	
				ssrDNA	lstrDNA	1	2
<i>Monorygma megacotyle</i>	1	<i>Cephaloscyllium umbratile</i>	off Heda, JAP	This study	This study	x	
	2	<i>Cephaloscyllium umbratile</i>	off Heda, JAP	This study	This study	x	
	3	<i>Galeus nipponensis</i>	off Kanaya, JAP	This study	This study	x	
	4	<i>Cephaloscyllium umbratile</i>	off Kanaya, JAP	This study	This study	x	
<i>Pachybothrium hutsoni</i>		<i>Nebrius ferrugineus</i>	Weipa, AUS	EF095260	EF095246	x	x
<i>Pedibothrium mounseyi</i>		<i>Nebrius ferrugineus</i>	Weipa, AUS	KF685893	KF685803	x	x
<i>Pelichnibothrium speciosum</i>		<i>Prionace glauca</i>	off Iburi, JAP	This study	This study	x	
		<i>Alepisaurus ferox</i>	off Toshima I., JAP	This study	This study	x	x
<i>Pelichnibothrium caudatum</i>		<i>Oncorhynchus keta</i>	off Erimo, JAP	This study	This study	x	x
<i>Symcallio violae</i>		<i>Mustelus canis</i>	Old Lyme, USA	KF685881	KF685813	x	x
<i>Trilocularia gracilis</i>		<i>Squalus acanthias</i>	Rhode I., USA	KF685776	KF685855	x	x
<i>Yorkeria hilli</i>		<i>Chiloscyllium punctatum</i>	Cairns, AUS	KF685903	KF685798	x	
<i>Rosebothrium ootenjikuzame n. sp.</i>		<i>Nebrius ferrugineus</i>	off Yaeyama, JAP	This study	This study	x	x
Onchoproteocephalidea							
<i>Acanthobothrium parviuncinatum</i>		<i>Urobatis maculatus</i>	Bahia de Los Angeles, MEX	EF095264	EF095250	x	x
<i>Acanthobothrium santarosaliense</i>		<i>Heterodontus mexicanus</i>	Santa Rosalia, MEX	KF685751	KF685834	x	x
<i>Pelidocotyle rugosa</i>		<i>Pseudoplattystoma fasciatum</i>	Rio Paraguay, PRY	AF286937	AF286989	x	x
<i>Phoreiobothrium lewinense</i>		<i>Sphyrna lewini</i>	Straits of Florida, USA	KF685896	KF685830	x	x
<i>Platybothrium auriculatum</i>		<i>Prionace glauca</i>	Montauk, New York, USA	KF685898	KF685837	x	x
<i>Platybothrium jondoeorum</i>		<i>Negaprion acutidens</i>	Darwin, AUS	KF685772	KF685829	x	x
<i>Potamotrygonocetus cf. fitzgeraldae</i>		<i>Potamotrygon castexi</i>	Madre de Dios River, PEL	KF685773	KF685832	x	x
<i>Prosobothrium armigerum</i>		<i>Prionace glauca</i>	Montauk, New York, USA	KF685899	KF685828	x	x
<i>Proteocephalus macrocephalus</i>		<i>Anguilla anguilla</i>	River Thames, GBR	EF095261	EF095247	x	x
<i>Proteocephalus perplexus</i>		<i>Amia calva</i>	Hay Bay, CAN	KF685873	KF685833	x	x
<i>Trilocularium anderssonorum</i>		<i>Negaprion acutidens</i>	Weipa, AUS	KF685895	KF685831	x	x
<i>Uncibilocularis okei</i>		<i>Pastinachus atrus</i>	Dundee Beach, AUS	KF685777	KF685835	x	x
<i>Onchoproteocephalidea gen. sp.*</i>		<i>Pristis clavata</i>	Darwin, AUS	KF685765	KF685836	x	x

Table 6 continued.

Specimens	Genotype	Host species	Locality	Accession Number		Dataset	
				ssrDNA	lsrDNA	1	2
Cathetocephalidea							
<i>Cathetocephalus thatcheri</i>		<i>Carcharhinus leucas</i>	Gulf of Mexico, USA	KF685884	KF685838	x	x
Rhinebothriidea							
<i>Anthocephalum</i> cf. <i>centrurum</i>		<i>Dasyatis centroura</i>	Mbour, SEN	FJ177099	FJ177059		x
<i>Rhinebothrium megacanthophallus</i>		<i>Himantura polylepis</i>	Kinabatangan River, MSY	AF286962	FJ177080		x
<i>Rhodobothrium paucitesticulare</i>		<i>Rhinoptera bonasus</i>	Davis, USA	FJ177100	FJ177060		x
Nippoteaniidea							
<i>Nippotaenia chaenogobii</i>		<i>Chaenogobius urotaenia</i>	Lake Taupo, NZL	AF286933	AF286987		x
Tetrabothriidea							
<i>Tetrabothrius erostris</i>		<i>Larus argentatus</i>	Danube Delta, ROU	AF286950	AJ287581		x
Cyclophyllidea							
<i>Hymenolepis diminuta</i>		<i>Rattus norvegicus</i>	Univ. Copenhagen, DNK	AY157181	AF286983		x
Lecanicephalidea							
<i>Adelobothrium</i> cf. <i>aetiobatidis</i>		<i>Aetobatus ocellatus</i>	Darwin, AUS	EF095257	EF095249	x	x
<i>Paraberrapex manifestus</i>		<i>Squatina californica</i>	Santa Rosalia, MEX	KF685868	KF685781	x	x
Litobothriidea							
<i>Litobothrium amplifica</i>		<i>Alopias pelagicus</i>	Santa Rosalia, MEX	KF685906	KF685843		x
<i>Litobothrium nickoli</i>		<i>Alopias pelagicus</i>	Santa Rosalia, MEX	KF685907	KF685844		x

Table 7 PCR and sequence primers of complete *ssrDNA* and partial *lsrDNA*. F means forward primer and R means reverse primer.

Primer names	Direction	Sequence	Reference
<i>ssrDNA</i>			
<i>PCR primers</i>			
WormA	F	GCG AAT GGC TCA TTA AAT CAG	Waeschenbach et al. (2007)
WormB	R	CTT GTT ACG ACT TTT ACT TCC	Waeschenbach et al. (2007)
<i>Sequence primers</i>			
300F	F	AGG GTT CGA TTC CGG AG	Waeschenbach et al. (2007)
300R	R	TCA GGC TCC CTC TCC GGA	Waeschenbach et al. (2007)
930F	F	GCA TGG AAT AAT GGA ATA GG	Waeschenbach et al. (2007)
1200F	F	CAG GTC TGT GAT GCC C	Waeschenbach et al. (2007)
1200R	R	GGG CAT CAC AGA CCT G	Waeschenbach et al. (2007)
1270F	F	ACT TAA AGG AAT TGA CGG	Waeschenbach et al. (2007)
1270R	R	CCG TCA ATT CCT TTA AGT	Waeschenbach et al. (2007)
18S-A27	R	CCA TAC AAA CGT CCC CGC CTG	Olson & Caira (1999)
<i>lsrDNA</i>			
<i>PCR primers</i>			
LSU5	F	TAG GTC GAC CCG CTG AAY TTA AGC A	Waeschenbach et al. (2007)
1500R	R	GCT ATC CTG AGG GAA ACT TCG	Waeschenbach et al. (2007)
<i>Sequence primers</i>			
300F	F	CAA GTA CCG TGA GGG AAA GTT G	Waeschenbach et al. (2007)
ECD2	R	CTT GGT CCG TGT TTC AAG ACG GG	Waeschenbach et al. (2007)
1200R	R	GCA TAG TTC ACC ATC TTT CGG	Waeschenbach et al. (2007)

Table 8 Sequence data of *ssrDNA* and *lsrDNA* of *Pelichnibothrium*, *Clistobothrium* and its related species. Only variable sites are shown, and numbers indicates the positions. Question marks mean missing sites. Completely identical sequences are indicated by gray markers.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 9 Sequence data of *ssrDNA* and *lsrDNA* of *Crossobothrium* and its related species. Only variable sites are shown, and numbers indicates the positions. Hyphen mean gaps.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 10 Sequence data of ssrDNA and lsrDNA of *Oryzomatobothrium musteli* and its related species. Only variable sites are shown, and numbers indicates the positions.

<p>本表については、5年以内に雑誌等で刊行予定のため、非公開。</p>

Table 11 Sequence data of ssrDNA and 1srDNA of *Phyllobothrium squali*, *Monorygma megacotyla* and its related species. Only variable sites are shown, and numbers indicates the positions. Question marks mean missing sites. Completely identical sequences are indicated by gray markers.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 12 Sequence data of *ssrDNA* and *lsrDNA* of *Paraoryzmatobothrium* and its related species. Only variable sites are shown, and numbers indicates the positions. Question marks mean missing sites, hyphen showed gaps. Completely identical sequences are indicated by gray markers.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 12 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 12 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 12 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 12 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 12 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 13 Comparison of statistical inference at each order. Order names were used revised orders in this study. UFBoot, ultrafast bootstrap; BPP, bayesian posterior probabilities; SH aLRT, SH-like interpretation of aLRT statistic. UFBoot ≥ 95 , BPP ≥ 0.98 , SH aLRT ≥ 80.0 are shown in bold face. Hyphens indicate branches which were not observed in each analysis.

本表については、5年以内に雑誌等で刊行予定のため、非公開.

Table 14 Morphological differences in the scolex among the revised genera of Phyllobothriidae. Degree means the degree of crumpling for the crumple type of bothridium. Asterisks mean the egg morphology was observed only for a part of the species of each genus. Old genera were also shown for *Pelichnibothrium* and *Scyphophyllidium* which included other synonymous old genera.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 14 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 15 New species and new combinations in this study. Abbreviations of the orders: Phy, Phyllobothriiidea; Tet, Tetraphyllidea; Rhi, Rhinebothriiidea.

本表については，5年以内に雑誌等で刊行予定のため，非公開.

Table 16 Sequence data of ssrDNA and lsrDNA of *Phyllobothrium* and its related species. Only variable sites are shown. Gray markers show the same sequence.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 17 Difference in measurements of *Pe. speciosum* between two different host species. Length and width were shown μm unit, large measurements were shown mm unit. L; length, MW; maxium width, W; width.

本表については，5年以内に雑誌等で刊行予定のため，非公開.

Table 18 Interspecific variations of *Sc. carcharinus*. L; length, MW; maxium width, W; width.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 19 Interspecific variations of *Sc. insulaeum*. L; length, MW; maxium width, W; width.

本表については，5年以内に雑誌等で刊行予定のため，非公開.

Table 20 Interspecific variations of *Sc.maritimum*. L; length, MW; maxium width, W; width.

本表については，5年以内に雑誌等で刊行予定のため，非公開.

Table 21 Interspecific variations of *Sc.ogasawaraensis*. L; length, MW; maxium width, W; width.

本表については，5年以内に雑誌等で刊行予定のため，非公開.

Table 22 Interspecific variations of *Ya. megacotyla*. L; length, MW; maxium width, W; width

本表については，5年以内に雑誌等で刊行予定のため，非公開.

Table 22 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開.

Table 23 Phyllobothriidean fauna in the coastal seas of Japan. Single asterisks mean species of new to Japan, double asterisks mean new host records.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 23 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 23 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 23 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 23 continued.

本表については、5年以内に雑誌等で刊行予定のため、非公開。

Table 24 List of examined host specimens for investigation of larvae of Phyllobothriidea and its related orders. N means the number of examined host specimens and the number of measured host specimens is shown in parentheses if all specimens were not measured, and n means the number of larvae used in molecular analyses/ the total number of collected larvae.

Host species	Station	N	Total Length (mm)		Weight (g)		n	Specimen ID
			Min	Max	Min	Max		
Mollusca								
Cephalopoda								
Enoploteuthidae								
<i>Watasenia scintillans</i>	St. 6	1	no data		no data		1/1	Larva 14
Gonatidae								
<i>Berryteuthis magister</i>	St. 20	6	235 – 361		133 – 315		4/97	Larva 11–13, Larva 27
Octopodidae								
<i>Octopus vulgaris</i>	St. 20	1	374		315			
Ommastrephidae								
<i>Todarodes pacificus</i>	St. 20	6 (4)	331 – 375		40 – 271		1/7	Larva 26
Sepiolidae								
<i>Rossia pacifica</i>	St. 20	6	163 – 252		35 – 110			
Gastropoda								
Buccininae								
<i>Buccinum striatissimum</i>	St. 20	3	138 – 166		156 – 221			
Arthropoda								
Malacostraca								
Crangonidae								
<i>Argis dentata</i>	St. 20	6	134 – 160		20 – 30			
Hippolytidae								
<i>Lebbeus groenlandicus</i>	St. 20	6	118 – 154		25 – 44			
Chordata								
Elasmobranchii								
Arhynchobatidae								
<i>Bathyrhaja smirnovi</i>	St. 20	Examined by investigation of Chapter I					2/2	Larva 15, 16

Table 24 continued.

Host species	Station	N	Total Length (mm)		Weight (g)		n	Specimen ID
			Min	Max	Min	Max		
Actinopterygii								
Acanthuridae								
<i>Zebrasoma flavescens</i>	St. 30	1	80		13			
Alepisauridae								
<i>Alepisaurus ferox</i>	St. 14	Examined by investigation of Chapter I					2/258	Larva 24, 25
Argentinidae								
<i>Glossanodon semifasciatus</i>	St. 20	5	202 – 229		40 – 66			
Balistidae								
<i>Sufflamen fraenatum</i>	St. 30	2(1)	340		840			
<i>Xanthichthys mento</i>	St. 30	1	no data		no data			
Bramidae								
<i>Taractichthys steindachneri</i>	St. 31	1	712		no data		2/3	Larva 30, 31
Carangidae								
<i>Carangoides orthogrammus</i>	St. 30	3	375 – 580		700 – 3060			
<i>Caranx lugubris</i>	St. 30	2	530 – 550		1990 – 3030			
<i>Seriola rivoliana</i>	St. 30	1	640 – 1000		2700 – 10800			
<i>Trachurus japonicus</i>	St. 19	2	280		230 – 250		1/16	Larva 5
Coryphaenidae								
<i>Coryphaena hippurus</i>	St. 19	1	880		4000		2/32	Larva 4, 34
Chaetodontidae								
<i>Chaetodon daedalma</i>	St. 30	1	175		160			
Cyclopteridae								
<i>Aptocyclus ventricosus</i>	St. 20	2	no data		184 – 246		2/38	Larva 20, 21
<i>Eumicrotremus asperimus</i>	St. 20	4 (3)	77 – 102		67 – 81		1/31	Larva 22
Gadidae								
<i>Gadus macrocephalus</i>	St. 20	2	324 – 614		162 – 2000		1/10	Larva 17
Holocentridae								
<i>Neoniphon sammara</i>	St. 30	1	150		46			
<i>Sargocentron ittodai</i>	St. 30	2	130 – 140		40 – 51			

Table 24 continued.

Host species	Station	N	Total Length (mm)		Weight (g)		n	Specimen ID
			Min	Max	Min	Max		
Kyphosidae								
<i>Girella leonina</i>	St. 30	1	590		3470			
<i>Kyphosus pacificus</i>	St. 30	1	430		1580			
Labridae								
<i>Bodianus perditio</i>	St. 30	2	190 – 380		130 – 750			
<i>Coris aygula</i>	St. 30	1	500		1990		1/1	Larva 10
<i>Thalassoma lutescens</i>	St. 30	4	170 – 230		53 – 150		1/1	Larva 9
Liparidae								
<i>Careproctus trachysoma</i>	St. 20	6	141 – 320		27 – 410		1/1	Larva 19
Lutjanidae								
<i>Gnathodentex aureolineatus</i>	St. 30	2	215 – 220		160 – 200			
<i>Gymnocranius euanus</i>	St. 30	11	270 – 475		400 – 2175			
<i>Gymnocranius griseus</i>	St. 30	2	325 – 345		620 – 710			
<i>Lethrinus rubrioperculatus</i>	St. 30	12 (11)	300 – 430		390 – 1000			
<i>Lutjanus kasmira</i>	St. 30	3	220 – 300		170 – 430			
<i>Lutjanus stellatus</i>	St. 30	1	585		3330			
<i>Paracaesio xanthura</i>	St. 30	3	220 – 390		160 – 560			
<i>Pristipomoides argyrogrammicus</i>	St. 30	1	350		620			
<i>Pristipomoides filamentosus</i>	St. 30	1	535		1930			
<i>Pristipomoides zonatus</i>	St. 30	3	365 – 420		770 – 1110			
Monacanthidae								
<i>Aluterus scriptus</i>	St. 30	1	475		550			
<i>Sufflamen bursa</i>	St. 30	1	210		230			
Mullidae								
<i>Mulloidichthys vanicolensis</i>	St. 30	2	240 – 260		160 – 210			
<i>Parupeneus multifasciatus</i>	St. 30	3	225 – 265		180 – 220			
<i>Parupeneus pleurostigma</i>	St. 30	5	210 – 305		120 – 310			

Table 24 continued.

Host species	Station	N	Total Length (mm)		Weight (g)		n	Specimen ID
			Min	Max	Min	Max		
Muraenesocidae								
<i>Muraenesox cinereus</i>	St. 19	1	514		no data			
Nemipteridae								
<i>Pentapodus nagasakiensis</i>	St. 30	1	160		60			
Pleuronectidae								
<i>Glyptocephalus stelleri</i>	St. 20	2	no data		no data			
<i>Hippoglossoides dubius</i>	St. 20	3	340 – 414		316 – 570		1/1	Larva 23
<i>Hippoglossoides pinetorum</i>	St. 20	2	281 – 431		159 – 625			
Pomacentridae								
<i>Abudefduf sexfasciatus</i>	St. 30	2	150	160	72	82		
Psychrolutidae								
<i>Dasycottus setiger</i>	St. 20	5 (4)	124 – 158		32 – 68		1/2	Larva 33
<i>Malacocottus gibber</i>	St. 20	3	208 – 272		207 – 530			
Salmonidae								
<i>Oncorhynchus keta</i>	St. 1	Examined by investigation of Chapter I					4/63	Larva 35–38
Sebastidae								
<i>Sebastes owstoni</i>	St. 20	4	172 – 283		52 – 190			
Serranidae								
<i>Cephalopholis miniata</i>	St. 30	2	305 – 355		400 – 670			
<i>Cephalopholis urodeta</i>	St. 30	1	240		220			
<i>Epinephelus fasciatus</i>	St. 30	8 (5)	245 – 315		190 – 410		3/24	Larva 6–8
<i>Epinephelus merra</i>	St. 30	2	148 – 170		35 – 67			
<i>Epinephelus retouti</i>	St. 30	4	335 – 375		620 – 850			
<i>Epinephelus tauvina</i>	St. 30	1	265		285			
<i>Variola louti</i>	St. 30	4(3)	320 – 750		320 – 5700		1/1	Larva 2
Scaridae								
<i>Scarus ghobban</i>	St. 30	1	205		160			

Table 24 continued.

Host species	Station	N	Total Length (mm)		Weight (g)		n	Specimen ID
			Min	Max	Min	Max		
Scombridae								
<i>Scomber australasicus</i>	St. 19	2	315	320	345	363	1/1	Larva 1
Siganidae								
<i>Siganus fuscescens</i>	St. 30	1	350		620			
Sparidae								
<i>Pagrus major</i>	St. 19	3	90	95	11	16		
Sphyraenidae								
<i>Sphyraena pinguis</i>	St. 19	2	135	150	9	15		
Tetraodontidae								
<i>Lagocephalus lagocephalus</i>	St. 30	2	360	365	500	570		
<i>Lagocephalus spadiceus</i>	St. 19	2	105	115	18	27		
Trichiuridae								
<i>Trichiurus japonicus</i>	St. 25	1	no data		no data		1/2	Larva 3
Trichodontidae								
<i>Arctoscopus japonicus</i>	St. 20	3	211	272	50	163	1/6	Larva 18
Xiphiidae								
<i>Xiphias gladius</i>	St. 31	1	1735		no data		2/104	Larva 28, 29
Zoarcidae								
<i>Bothrocara hollandi</i>	St. 20	6	244	294	53	89		
<i>Lycodes nakamurai</i>	St. 20	5 (3)	302	314	69	97	1/22	Larva 32
<i>Lycodes tanakae</i>	St. 20	3 (2)	475	784	1350	2250		
<i>Lycodes toyamensis</i>	St. 20	3	318	380	147	235		
Mammalia								
Delphinidae								
<i>Stenella coeruleoalba</i>	St. 25		Examined by investigation of Chapter I				4/7	Larva 39-42

Table 25 List of the specimens for molecular phylogenetic analyses of larvae. Species names revised in this study are used. Larvae specimens which collected in this study were given the number to distinguish the individual, Larva 1-42 (see Table 24). Alphabet specific names of *Scyphophyllidium* are undescribed species reported by Jensen & Bullard (2010). Abbreviations of country names in the locality: AUS; Australia, GBR; United Kingdom, JAP; Japan, NZL; New Zealand, MYS; Malaysia, SEN; Senegal, USA; U.S.A.

Specimens	Host species	Locality	Accession Number		Dataset		
			1stDNA	COI	3	4	5
Adult							
Phyllobothriidea							
<i>Phyllobothrium</i> cf. <i>lactuca</i>	<i>Mustelus mento</i>	Puerto Montt, CHL	KF685845	-		x	
<i>Phyllobothrium hoshizame</i>	<i>Mustelus manazo</i>	off Kanaya, JAP	This study	-		x	
<i>Phyllobothrium serratum</i>	<i>Triakis scyllium</i>	off Shimoda, JAP	This study	-		x	
<i>Alexandercestus oomejirozame</i>	<i>Carcharhinus leucas</i>	off Yaeyama, JAP	This study	-		x	
<i>Alexandercestus gibsoni</i>	<i>Negaprion acutidens</i>	Timor Sea, AUS	KC505623	-		x	
<i>Bilocularia hyperapolytica</i>	<i>Dalatias licha</i>	off Kanaya, JAP	This study	-		x	
<i>Calliobothrium</i> cf. <i>verticillatum</i>	<i>Mustelus canis</i>	Old Lyme, USA	KF685812	-		x	
<i>Calliobothrium shirozame</i>	<i>Mustelus griseus</i>	off Shimoda, JAP	This study	-		x	
<i>Calyptrobthrium</i> sp.	<i>Torpedo nobiliana</i>	Rhode Island, USA	KF685848	-		x	
<i>Calyptrobthrium</i> sp. 1	<i>Torpedo tokionis</i>	off Heda, JAP	This study	-		x	
<i>Chimaerocestos</i> sp. 1	<i>Rhinochimaera pacifica</i>	Chatham Rise, NZL	KF685827	-		x	
<i>Chimaerocestos</i> sp. 2	<i>Rhinochimaera pacifica</i>	Chatham Rise, NZL	KF685850	-		x	
<i>Crossobothrium laciniatum</i>	<i>Hexanchus griseus</i>	Puerto Montt, CHL	KF685824	-		x	
<i>Crossobothrium campanulatum</i>	<i>Hexanchus griseus</i>	off Sado I., JAP	This study	-		x	
<i>Crossobothrium dohrnii</i>	<i>Heptranchias perlo</i>	off Kanaya, JAP	This study	-		x	
<i>Crossobothrium</i> cf. <i>dohrnii</i>	<i>Heptranchias perlo</i>	Gulf of Mexico	KF685847	-		x	
<i>Mitsukurincestus gobelinus</i>	<i>Mitsukurina owstoni</i>	off Kanaya, JAP	This study	-		x	
<i>Monorygma chamydoselachi</i>	<i>Chlamydoselachus</i>	off Kanaya, JAP	This study	This study		x	x
<i>Orygmatobothrium musteli</i>	<i>Mustelus manazo</i>	off Heda, JAP	This study	-		x	
	<i>Mustelus griseus</i>	off Chigasaki, JAP	This study	-		x	
<i>Orygmatobothrium</i> cf. <i>musteli</i> 1	<i>Mustelus mustelus</i>	Soumbédioune, SEN	KF685815	-		x	
<i>Orygmatobothrium</i> cf. <i>musteli</i> 2	<i>Mustelus mustelus</i>	Ouakam, SEN	KF685816	-		x	

Table 25 continued.

Specimens	Host species	Locality	Accession Number		Dataset				
			IsrDNA	COI	3	4	5		
<i>Pelichnibothrium speciosum</i>	<i>Prionace glauca</i>	off Iburi, JAP	This study	This study	x	x	x		
	<i>Prionace glauca</i>	off Iburi, JAP	This study	This study			x		
	<i>Alepisaurus ferox</i>	off Toshima I., JAP	This study	This study	x	x	x		
	<i>Alepisaurus ferox</i>	off Toshima I., JAP	This study	This study			x		
<i>Pelichnibothrium carcharodoni</i>	<i>Carcharodon carcharias</i>	Kaipara Harbour, NZL	HM856633	-	x	x			
	<i>Carcharodon carcharias</i>	off Iburi, JAP	This study	This study	x	x	x		
	<i>Carcharodon carcharias</i>	off Iburi, JAP	This study	This study			x		
<i>Pelichnibothrium montaukensis</i>	<i>Isurus oxyrinchus</i>	Montauk, New York, USA	AF286996	JQ268541	x	x	x		
	<i>Isurus oxyrinchus</i>	off Toshima I., JAP	This study	This study	x	x	x		
	<i>Isurus oxyrinchus</i>	off Toshima I., JAP	This study	This study			x		
	<i>Isurus oxyrinchus</i>	off Ogasawara Is., JAP	This study	This study			x		
	<i>Isurus oxyrinchus</i>	off Obama, JAP	This study	This study			x		
<i>Pelichnibothrium tumidum</i>	<i>Carcharodon carcharias</i>	off Iburi, JAP	This study	This study	x	x	x		
	<i>Carcharodon carcharias</i>	off Iburi, JAP	This study	This study			x		
<i>Pelichnibothrium cf. montaukensis</i>	<i>Lamna nasus</i>	Falkland Is., GBR	JF436969	-	x	x	x		
<i>Scyphophyllodium bai</i>	<i>Mustelus mustelus</i>	Ouakam Senegal, SEN	KC505625	-	x				
<i>Scyphophyllidium carcharhinus</i>	<i>Carcharhinus limbatus</i>	off Yaeyama, JAP	This study	This study	x				
	<i>Carcharhinus leucas</i>	off Yaeyama, JAP	This study	This study	x				
	<i>Carcharhinus galapagensis</i>	off Ogasawara Is., JAP	This study	This study	x				
<i>Scyphophyllidium eirakubuka</i>	<i>Hemirhamphys japonica</i>	off Shimoda, JAP	This study	This study	x				
<i>Scyphophyllodium exiguum</i>	<i>Alopias vulpinus</i>	Montauk, New York, USA	KF685769	KF685822	x				
<i>Scyphophyllodium guariticus</i>	<i>Paratrygon aiereba</i>	Mouth of Rio Shilve, PER	KF685888	KF685817	x				
<i>Scyphophyllidium hanazame</i>	<i>Carcharhinus brevipinna</i>	off Iburi, JAP	This study	This study	x				
<i>Scyphophyllidium insulatum</i>	<i>Carcharhinus plumbeus</i>	off Yaeyama, JAP	This study	This study	x				
	<i>Carcharhinus galapagensis</i>	off Ogasawara Is., JAP	This study	This study	x				
<i>Scyphophyllodium janineae</i>	<i>Hemigaleus australiensis</i>	Moreton Bay, AUS	HQ680627	-	x				
<i>Scyphophyllodium kirstenae</i>	<i>Hemigaleus microstoma</i>	Mukah, Borneo, MYS	KC505626	-	x				
<i>Scyphophyllidium latipi</i>	<i>Scoliodon macrorhynchus</i>	Mukah, Borneo, MYS	KF685853	-	x				
<i>Scyphophyllidium longiproglottius</i>	<i>Carcharhinus limbatus</i>	off Yaeyama, JAP	This study	-	x				

Table 25 continued.

Specimens	Host species	Locality	Accession Number		Dataset				
			IsrDNA	COI	3	4	5		
<i>Scyphophyllidium maritimum</i>	<i>Carcharhinus brevipinna</i>	off Iburi, JAP	This study	-				x	
	<i>Carcharhinus galapagensis</i>	off Ogasawara Is., JAP	This study	-				x	
<i>Scyphophyllidium nenuribuka</i>	<i>Trienodon obesus</i>	off Ogasawara Is., JAP	This study	-				x	
<i>Scyphophyllidium ogasawaraensis</i>	<i>Trienodon obesus</i>	off Ogasawara Is., JAP	This study	-				x	
	<i>Carcharhinus galapagensis</i>	off Ogasawara Is., JAP	This study	-				x	
<i>Scyphophyllidium paulum</i>	<i>Galeocerdo cuvier</i>	off Iburi, JAP	This study	-				x	
	<i>Galeocerdo cuvier</i>	Moreton Bay, AUS	HQ680630	-				x	
<i>Scyphophyllidium prionacis</i>	<i>Prionace glauca</i>	Montauk, New York, USA	KF685818	-				x	
	<i>Prionace glauca</i>	off Tosashimizu, JAP	This study	-				x	
<i>Scyphophyllidium randyi</i>	<i>Chiloscyllium hasseltii</i>	Mukah, Borneo, MYS	KF685820	-				x	
<i>Scyphophyllidium sumitukizame</i>	<i>Carcharhinus dussumieri</i>	off Tosasaga, JAP	This study	-				x	
<i>Scyphophyllidium triacis</i>	<i>Triakis scyllium</i>	off Shimoda, JAP	This study	-				x	
<i>Scyphophyllidium taylori</i>	<i>Hemigaleus australiensis</i>	Moreton Bay, AUS	HQ680633	-				x	
<i>Scyphophyllidium taylori</i>	<i>Chiloscyllium punctatum</i>	Mukah, Borneo, MYS	KF685819	-				x	
<i>Scyphophyllidium yogore</i>	<i>Carcharhinus longimanus</i>	off Yaeyama, JAP	This study	-				x	
<i>Scyphophyllidium</i> sp. A	<i>Carcharhinus brevipinna</i>	Gulf of Mexico, USA	GQ470001	-				x	
<i>Scyphophyllidium</i> sp. B	<i>Carcharhinus brevipinna</i>	Gulf of Mexico, USA	GQ470031	-				x	
	<i>Carcharhinus isodon</i>	Gulf of Mexico, USA	GQ470037	-				x	
	<i>Carcharhinus limbatus</i>	Gulf of Mexico, USA	GQ470034	-				x	
	<i>Rhizoprionodon terraenovae</i>	Gulf of Mexico, USA	GQ470033	-				x	
<i>Scyphophyllidium</i> sp. C	<i>Rhizoprionodon terraenovae</i>	Gulf of Mexico, USA	GQ470020	-				x	
<i>Scyphophyllidium</i> sp. D	<i>Carcharhinus brevipinna</i>	Gulf of Mexico, USA	GQ470011	-				x	
	<i>Carcharhinus limbatus</i>	Gulf of Mexico, USA	GQ470005	-				x	
	<i>Rhizoprionodon terraenovae</i>	Gulf of Mexico, USA	GQ470006	-				x	
<i>Scyphophyllidium</i> cf. <i>giganteum</i>	<i>Galeorhinus galeus</i>	Chatham Rise, NZL	KF685854	-				x	
<i>Scyphophyllidium</i> sp.	<i>Alopias pelagicus</i>	Santa Rosalia, MEX	KF685821	-				x	
<i>Symcallio violae</i>	<i>Mustelus canis</i>	Old Lyme, USA	KF685813	-				x	
<i>Thysanocephalum thysanocephalum</i>	<i>Galeocerdo cuvier</i>	off Yaeyama, JAP	This study	-				x	
	<i>Galeocerdo cuvier</i>	off Iburi, JAP	KF685823	-				x	

Table 25 continued.

Specimens	Host species	Locality	Accession Number		Dataset		
			IsrDNA	COI	3	4	5
<i>Trilocularia gracilis</i>	<i>Squalus acanthias</i>	Rhode I., USA	KF685855	-		x	
<i>Yamaguticestus squali</i>	<i>Cephaloscyllium umbratile</i>	off Heda, JAP	This study	-		x	
	<i>Cephaloscyllium umbratile</i>	off Heda, JAP	This study	-		x	
	<i>Cephaloscyllium umbratile</i>	off Kanaya, JAP	This study	-		x	
	<i>Galeus nipponensis</i>	off Kanaya, JAP	This study	-		x	
	<i>Squalus acanthias</i>	Rhode Island, USA	KF685846	-		x	
	<i>Squalus japonicus</i>	off Kanaya, JAP	This study	-		x	
	<i>Squalus japonicus</i>	off Mimase, JAP	This study	-		x	
	<i>Squalus mitsukurii</i>	off Kanaya, JAP	This study	-		x	
	<i>Squalus mitsukurii</i>	off Kanaya, JAP	This study	-		x	
<i>Vertebraeovicescus dobukasube</i>	<i>Bathyraja smirnovi</i>	off Oki Is., JAP	This study	-		x	
"Phyllorhina gen. sp."	<i>Sphyrna lewini</i>	Straits of Florida, USA	KF685814	-		x	
Tetrapteryllidae							
<i>Anthobothrium caseyi</i>	<i>Prionace glauca</i>	Montauk, New York, USA	KF685805	-		x	
	<i>Prionace glauca</i>	off Tosashimizu, JAP	This study	-		x	
<i>Anthobothrium</i> sp. 1	<i>Carcharhinus tilstoni</i>	Wessel Is., AUS	KF685806	-		x	
<i>Anthobothrium</i> sp. 2	<i>Carcharhinus brevipinna</i>	off Iburu, JAP	This study	-		x	
<i>Anthobothrium</i> sp. 3	<i>Carcharhinus plumbeus</i>	off Yaeyama, JAP	This study	-		x	
<i>Anthobothrium</i> sp. 4	<i>Carcharhinus brevipinna</i>	off Iburu, JAP	This study	-		x	
<i>Anthobothrium</i> sp. 5	<i>Carcharhinus dussumieri</i>	off Tosasaga, JAP	This study	-		x	
<i>Balanobothrium</i> sp.	<i>Stegostoma fasciatum</i>	Mukah, Borneo, MYS	KF685802	-		x	
<i>Ceratobothrium xanthocephalum</i>	<i>Isurus oxyrinchus</i>	Montauk, New York, USA	KF685849	-		x	
<i>Ceratobothrium</i> sp. 1	<i>Carcharodon carcharias</i>	off Iburu, JAP	This study	-		x	
<i>Megalonchos shawae</i>	<i>Hemipristis elongata</i>	Wessel Is., AUS	KF685807	-		x	
<i>Pachybothrium hutsoni</i>	<i>Nebrius ferrugineus</i>	Weipa, AUS	EF095246	-		x	
<i>Pedibothrium mounseyi</i>	<i>Nebrius ferrugineus</i>	Weipa, AUS	KF685803	-		x	
<i>Yorkeria hilli</i>	<i>Chiloscyllium punctatum</i>	Cairns, AUS	KF685798	-		x	
<i>Rosebothrium ootenjikuzame</i>	<i>Nebrius ferrugineus</i>	off Yaeyama, JAP	This study	-		x	

Table 25 continued.

Specimens	Host species	Locality	Accession Number		Dataset		
			IsrDNA	COI	3	4	5
Onchoproteocephalidea							
<i>Acanthobothrium parvuncinatum</i>	<i>Urobatis maculatus</i>	Bahia de Los Angeles, MEX	EF095250	-		x	
<i>Acanthobothrium santarosaliense</i>	<i>Heterodontus mexicanus</i>	Santa Rosalia, MEX	KF685834	-		x	
<i>Pelidocotyle rugosa</i>	<i>Pseudoplattystoma fasciatum</i>	Rio Paraguay, PRY	AF286989	-		x	
<i>Phoreiobothrium lewinense</i>	<i>Sphyrna lewini</i>	Straits of Florida, USA	KF685830	-		x	
<i>Platybothrium auriculatum</i>	<i>Prionace glauca</i>	Montauk, New York, USA	KF685837	-		x	
<i>Platybothrium jondoeorum</i>	<i>Negaprion acutidens</i>	Darwin, AUS	KF685829	-		x	
<i>Potamotrygonocestus</i> cf. <i>fitzgeraldae</i>	<i>Potamotrygon castexi</i>	Madre de Dios River, PEL	KF685832	-		x	
<i>Prosobothrium armigerum</i>	<i>Prionace glauca</i>	Montauk, New York, USA	KF685828	-		x	
<i>Proteocephalus macrocephalus</i>	<i>Anguilla anguilla</i>	River Thames, GBR	EF095247	-		x	
<i>Proteocephalus perplexus</i>	<i>Amia calva</i>	Hay Bay, CAN	KF685833	-		x	
<i>Triloculatum andersonorum</i>	<i>Negaprion acutidens</i>	Weipa, AUS	KF685831	-		x	
<i>Uncibilocularis okei</i>	<i>Pastinachus atrus</i>	Dundee Beach, AUS	KF685835	-		x	
Onchoproteocephalidea gen. sp.	<i>Pristis clavata</i>	Darwin, AUS	KF685836	-		x	
Cathetocephalidea							
<i>Cathetocephalus thatcheri</i>	<i>Carcharhinus leucas</i>	Gulf of Mexico, USA	KF685838	-		x	
Rhinebothriidea							
<i>Anthocephalum biacetabulatum</i>	<i>Rhinobatos schlegelii</i>	off Chigasaki, JAP	This study	-		x	
<i>Anthocephalum</i> cf. <i>centrurum</i>	<i>Dasyatis centroura</i>	Mbour, SEN	FJ177059	-		x	
<i>Rhinebothrium megacanthophallus</i>	<i>Himantura polylepis</i>	Kinabatangan River, MSY	FJ177080	-		x	
<i>Rhodobothrium paucitesticulare</i>	<i>Rhinoptera bonasus</i>	Davis, USA	FJ177060	-		x	
Lecanicephalidea							
<i>Adelobothrium</i> cf. <i>aetiobatidis</i>	<i>Aetobatus ocellatus</i>	Darwin, AUS	EF095249	-		x	
<i>Paraberrapex manifestus</i>	<i>Squatina californica</i>	Santa Rosalia, MEX	KF685781	-		x	
Litobothriidea							
<i>Litobothrium amplifica</i>	<i>Alopias pelagicus</i>	Santa Rosalia, MEX	KF685843	-		x	
<i>Litobothrium nickoli</i>	<i>Alopias pelagicus</i>	Santa Rosalia, MEX	KF685844	-		x	

Table 25 continued.

Specimens	Host species	Locality	Accession Number		Dataset				
			IsrDNA	COI	3	4	5		
Plerocercoid									
<i>Pelichnibothrium carcarodoni</i>	<i>Grampus griseus</i>	Spain	GQ839587	-		x	x		
	<i>Stenella coeruleoalba</i>	Spain	GQ839588	-		x	x		
<i>Pelichnibothrium caudatum</i>	-Larva 35 <i>Oncorhynchus keta</i>	off Erimo, JAP	This study	This study		x	x		x
	-Larva 36 <i>Oncorhynchus keta</i>	off Erimo, JAP	This study	This study		x	x		x
	-Larva 37 <i>Oncorhynchus keta</i>	off Erimo, JAP	This study	This study		x	x		x
	-Larva 38 <i>Oncorhynchus keta</i>	off Erimo, JAP	This study	This study		x	x		x
<i>Pelichnibothrium cf. montaukensis</i>	<i>Doryteuthis gahi</i>	Falkland Is., GBR	AF382072	-		x	x		
	<i>Regalecus glesne</i>	Santa Catalina I., USA	KM272992	-		x	x		
<i>Scyphophylodinium</i> sp. C	<i>Trichiurus lepturus</i>	Gulf of Mexico, USA	GQ470022	-		x			
<i>Scyphophylodinium</i> sp. D	<i>Lobotes surinamensis</i>	Gulf of Mexico, USA	GQ470014	-		x			
	<i>Paralichthys lethostigma</i>	Gulf of Mexico, USA	GQ470008	-		x			
Plerocercoid	<i>Scomber australasicus</i>	off Maizuru, JAP	This study	-		x			
-Larva 1		off Ogasawara Is., JAP	This study	-		x			
-Larva 2	<i>Variola louti</i>	off Kubotsu, JAP	This study	-		x			
-Larva 3	<i>Trichiurus japonicus</i>	off Maizuru, JAP	This study	-		x			
-Larva 4	<i>Trachurus japonicus</i>	off Maizuru, JAP	This study	-		x			
-Larva 5	<i>Coryphaena hippurus</i>	off Maizuru, JAP	This study	-		x			
-Larva 6	<i>Epinephelus fasciatus</i>	off Ogasawara Is., JAP	This study	-		x			
-Larva 7	<i>Epinephelus fasciatus</i>	off Ogasawara Is., JAP	This study	-		x			
-Larva 8	<i>Epinephelus fasciatus</i>	off Ogasawara Is., JAP	This study	-		x			
-Larva 9	<i>Thalassoma lutescens</i>	off Ogasawara Is., JAP	This study	-		x			
-Larva 10	<i>Coris aygula</i>	off Ogasawara Is., JAP	This study	-		x			
-Larva 11	<i>Berryteuthis magister</i>	off Oki Is., JAP	This study	This study		x	x		x
-Larva 12	<i>Berryteuthis magister</i>	off Oki Is., JAP	This study	This study		x	x		x
-Larva 13	<i>Berryteuthis magister</i>	off Oki Is., JAP	This study	This study		x	x		x
-Larva 14	<i>Watasenia scintillans</i>	off Toyama, JAP	This study	This study		x	x		x
-Larva 15	<i>Bathyraja smirnovi</i>	off Oki Is., JAP	This study	This study		x	x		x
-Larva 16	<i>Bathyraja smirnovi</i>	off Oki Is., JAP	This study	This study		x	x		x

Table 25 continued.

Specimens	Host species	Locality	Accession Number		Dataset		
			IsrDNA	COI	3	4	5
Plerocercoid -Larva 17	<i>Gadus macrocephalus</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 18	<i>Arctoscopus japonicus</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 19	<i>Careproctus trachysomai</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 20	<i>Aptocyclus ventricosus</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 21	<i>Aptocyclus ventricosus</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 22	<i>Eumicrotremus asperrimus</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 23	<i>Hippoglossoides dubius</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 24	<i>Alepisaurus ferox</i>	off Toshima I., JAP	This study	This study	x	x	x
-Larva 25	<i>Alepisaurus ferox</i>	off Toshima I., JAP	This study	This study	x	x	x
-Larva 26	<i>Todarodes pacificus</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 27	<i>Berryteuthis magister</i>	off Oki Is., JAP	This study	This study	x	x	x
-Larva 28	<i>Xiphias gladius</i>	off Ogasawara Is., JAP	This study	-	x		
-Larva 29	<i>Xiphias gladius</i>	off Ogasawara Is., JAP	This study	-	x		
-Larva 30	<i>Taractichthys steindachneri</i>	off Ogasawara Is., JAP	This study	-	x		
-Larva 31	<i>Taractichthys steindachneri</i>	off Ogasawara Is., JAP	This study	-	x		
-Larva 32	<i>Lycodes nakamurae</i>	off Oki Is., JAP	This study	-	x		
-Larva 33	<i>Dasycottus setiger</i>	off Oki Is., JAP	This study	-	x		
-Larva 34	<i>Coryphaena hippurus</i>	off Maizuru, JAP	This study	-	x		
Plerocercoid	<i>Ariopsis felis</i>	Gulf of Mexico, USA	GQ470051	-	x		
	<i>Cynoscion nebulosus</i>	Gulf of Mexico, USA	GQ470049	-	x		
	<i>Opsanus beta</i>	Gulf of Mexico, USA	GQ470030	-	x		
	<i>Stenella coeruleoalba</i>	Spain	DQ839568	-	x	x	
	<i>Grampus griseus</i>	Spain	DQ839573	-	x	x	
	<i>Tursiops truncatus</i>	Spain	DQ839580	-	x	x	

Table 25 continued.

Specimens	Host species	Locality	Accession Number		Dataset				
			IsrDNA	COI	3	4	5		
Meroceroid									
<i>Pelichnibothrium delphini</i> - Larva 39 - Larva 40	<i>Stenella coeruleoalba</i>	Minami-satsuma, JAP	This study	This study		x	x		x
	<i>Stenella coeruleoalba</i>	Minami-satsuma, JAP	This study	This study			x	x	x
	<i>Stenella coeruleoalba</i>	Spain	AY741599	-		x	x		
	<i>Grampus griseus</i>	Spain	DQ839593	-		x	x		
	<i>Tursiops truncatus</i>	Spain	DQ839590	-			x	x	
<i>Pelichnibothrium grimaldii</i> - Larva 41 - Larva 42	<i>Stenella coeruleoalba</i>	Minami-satsuma, JAP	This study	This study		x	x		x
	<i>Stenella coeruleoalba</i>	Minami-satsuma, JAP	This study	This study			x	x	x
	<i>Stenella coeruleoalba</i>	Spain	AY741591	-			x	x	
	<i>Grampus griseus</i>	Spain	DQ839585	-			x	x	
	<i>Tursiops truncatus</i>	Spain	DQ839582	-			x	x	

Table 26 Sequence data of 1srDNA of plerocercoid, merocercoid, and adult *Pelichnibothrium*. Only variable sites are shown, and numbers indicate the positions. Question marks mean missing sites.

本表については、5年以内に雑誌等で刊行予定のため、非公開.

Table 27 List of phyllobothriidean species having multiple final host species. Abbreviations of host order names in order: Car; Carcharinidae, Etm; Etomopteridae, Hex; Hexanchidae, Odo; Odontaspidae, Tri; Triakidae, Scy; Scyliorhinidae, Squ; Squalidae.

本表については，5年以内に雑誌等で刊行予定のため，非公開.

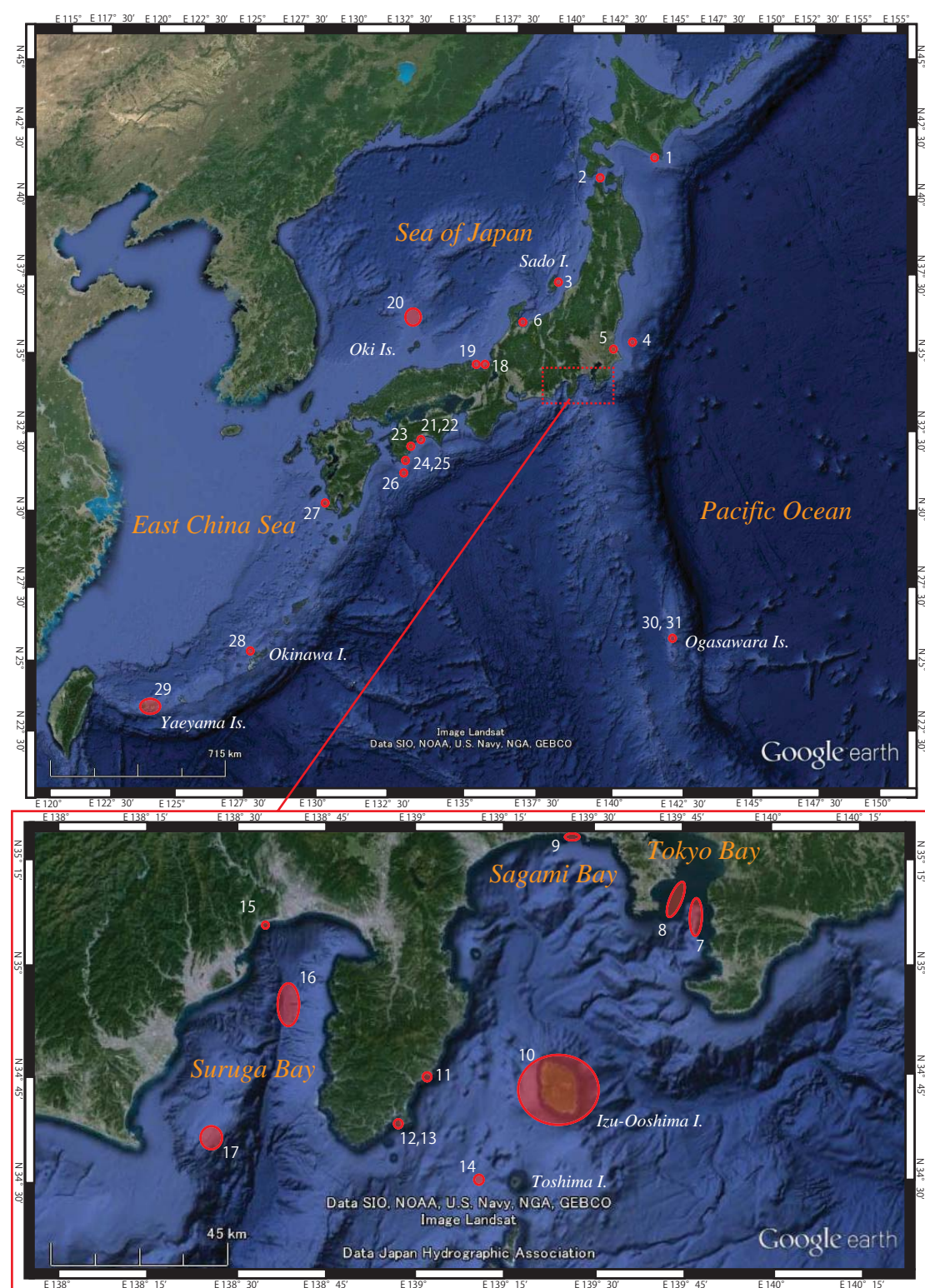


Fig. 1 Sampling locality. Circles show sampling locality, and numbers denote the station number in Table 4. See Table 4 for the detailed information on each station.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 2 Morphological terminology of Phyllobothriidea. **A**, Whole worm (*Scyphophyllidium brevipinnacis*), **B**, Scolex (*Pelichnibothrium speciosum*), **C**, Scolex (*Orygmatobothrium musteli*), **D**, Scolex (*Mitsukuricestus gobelinus*), **E**, Mature proglottid (*Sc. brevipinnacis*), **F**, Free proglottid (*Pe. speciosum*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 3 Maximum likelihood tree by Analysis 1 (Dataset 1, ssrDNA and lsrDNA). UFBootstrap and BPP are shown at each branch, and hyphens indicate branches which were not observed in the bayesian tree. Red arrowhead shows the node of the clade of revised order supported by SH aLRT ≥ 80 (see Table 8). Current scientific names are used and some of them were revised in this study (see Table 15). Sequences obtained in this study are shown in black color, and those collected from GenBank are in gray color. Open and solid circles near species names indicate the current order sensu Caira et al. (2014), Phyllobothriidea and Tetraphyllidea, respectively. Open boxes indicate the revised orders in this study (see Section 3.3). Polyphyletic genera are shown by dotted lines: A, *Phyllobothrium*; B, *Clistobothrium*; C, *Monorygma*; D, *Marsupiobothrium*; E, *Paraorygmatobothrium*.

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 4 Maximum likelihood tree by Analysis 2 (Dataset 1, ssrDNA). See Fig. 3 for the explanations of symbols in the tree. Phyllobothriidea, Tetraphyllidea and Onchoproteocephalidea were polyphyletic in this analysis (connected by dotted lines).

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 5 Maximum likelihood tree by Analysis 3 (Dataset 1, lsrDNA). See Fig. 3 for the explanations of symbols in the tree. Phyllobothriidea were polyphyletic in this analysis (connected by dotted lines).

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 6 Maximum likelihood tree by Analysis 4 (Dataset 2, ssrDNA and lsrDNA). See Fig. 3 for the explanations of symbols in the tree.

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 7 Maximum likelihood tree by Analysis 5 (Dataset 2, ssrDNA). See Fig. 3 for the explanations of symbols in the tree. Phyllobothriidea were polyphyletic in this analysis (connected by dotted lines).

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 8 Maximum likelihood tree by Analysis 6 (Dataset 2, lsrDNA). See Fig. 3 for the explanations of symbols in the tree. Phyllobothriidea were polyphyletic in this analysis (connected by dotted lines).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 9 Phylogenetic consideration of the hook in Phyllobothriidea and related orders. Based on the tree of Analysis 1 (Fig. 3), robust nodes (UFBoot ≥ 95 or BPP ≥ 0.98) are shown by solid circles, and the other nodes are shown by open circles. Hooked taxa are shown in red color, and non-hooked taxa are shown in black. Open boxes show the revised orders (see Section 3.3).

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 10 Phylogenetic consideration of the bothridium types in the revised Phyllobothriidea. Based on the tree of Analysis 1 (Fig. 3), robust nodes (UFBoot ≥ 95 or BPP ≥ 0.99) are shown by solid circles, and the other nodes are shown by open circles. Colors show bothridium types: black, flat; blue, crumple; pink, cup; green, divided; light green, loculate margin; purple, sac.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 11 Phylogenetic consideration of the types of bothridium base in the revised Phyllobothriidea. See Fig. 10 for the explanations of the tree. Stalked taxa are shown in red color, and sessile taxa are shown in black.

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 12 Phylogenetic consideration of the laciniation in Phyllobothriidea. See Fig. 10 for the explanations of the tree. Laciniated taxa were shown by red color, and non-laciniated taxa shown by black.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 13 Phylogenetic consideration of the additional suckers and the reduced suckers in Phyllobothriidea. See Fig. 10 for the explanations of the tree. Colors show additional suckers: black, general; blue, apical sucker; yellow, central sucker; green, no sucker.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 14 Revision of the genera in the revised Phyllobothriidea. The phylogenetic tree was based on Analysis 1 (Fig. 3). Bars indicate the revised genera. Type species of the current genera are shown in bold face. Robust nodes (UFBoot ≥ 95 or BPP ≥ 0.98) are shown by solid circles and thick lines, and the other nodes are shown by open circles and thin lines.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 15 *Phyllobothrium hoshizame* n. sp. **A**, Scolex, **B**, Mature proglottid, No. *Mustelus manazo* 27 (Host: *Mustelus manazo*).

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 16 *Phyllobothrium serratum*. **A**, Scolex, No. *Triakis scyllium* 32 (Host: *Triakis scyllium*). **B**, Mature proglottid, **C**, Gravid proglottid, **D**, Eggs, MPM 22715, SY7159 (Host: *Triakis scyllium*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 17 *Alexandercestus oomejirozame* n. sp. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, No. *Carcharhinus leucas* 10 (Host: *Carcharhinus leucas*). Overlapping vitelline follicles are not drawn on the left side of a broken line.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 18 *Bilocularia hyperapolytica*. **A**, Scolex, No. *Dalatias licha* 15 (Host: *Dalatias licha*). **B**, Free proglottid, **C**, Egg, No. *Dalatias licha* 16 (Host: *Dalatias licha*).

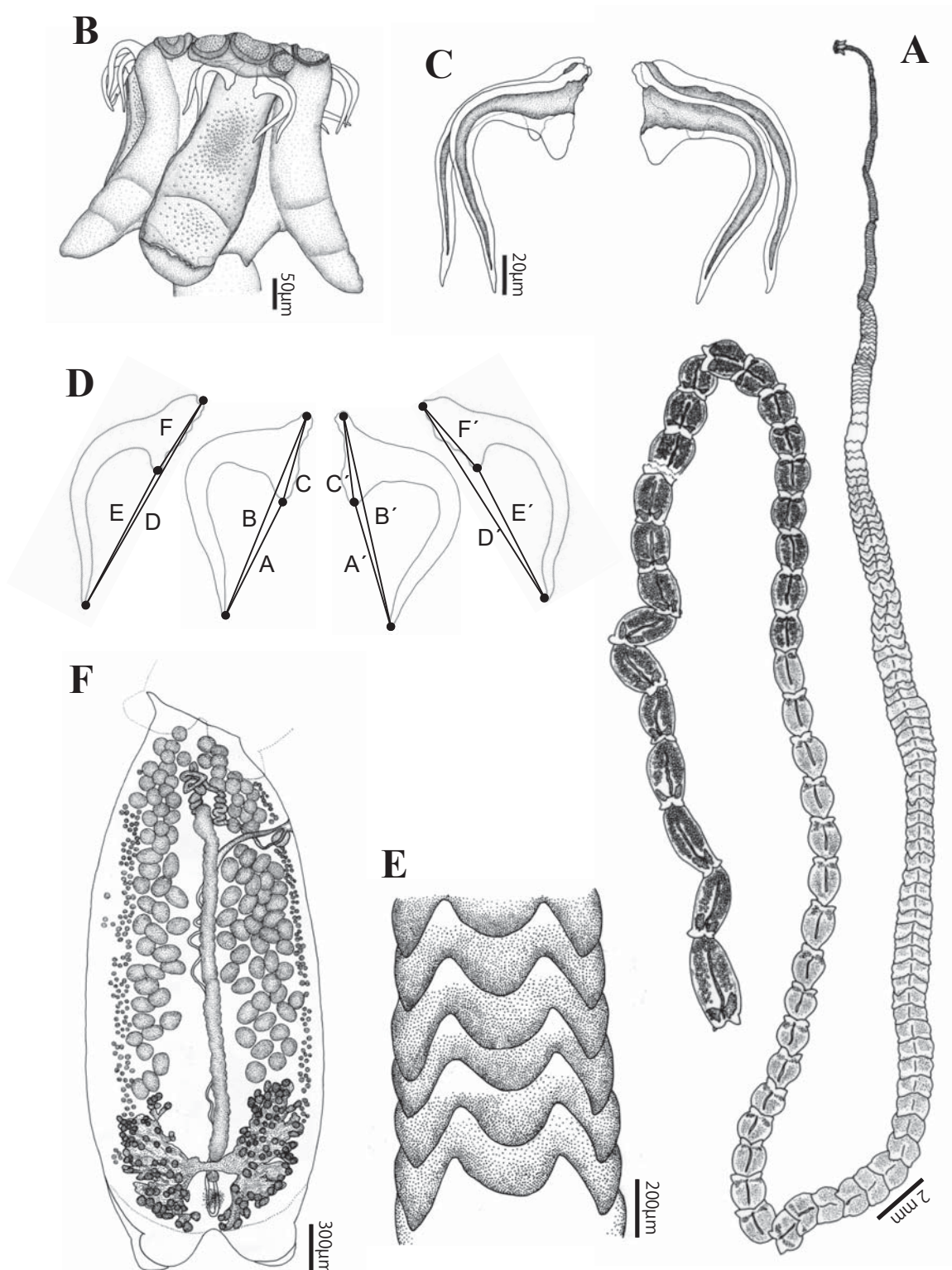


Fig. 19 *Calliobothrium shirozame* n. sp. **A**, Whole worm, **B**, Scolex, **C**, Hook, **D**, Hook measurements, letters reflect those used in the description, **E**, Mature proglottid, **F**, Lacination, NSMT-PI 6037 (Host: *Mustelus griseus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 20 *Calyptrorhynchium* sp. Scolex, No. *Torpedo tokionis* 1 (Host: *Torpedo tokionis*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 21 *Crossobothrium campanulatum*. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, **D**, Egg, No. *Hexanchus griseus* 1 (Host: *Hexanchus griseus*). Overlapping vitelline follicles are not drawn on the right side of a broken line.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 22 *Crossobothrium dohrni*. **A**, Whole worm, No. *Heptranchias perlo* 23 (Host: *Heptranchias perlo*). **B**, Scolex, **B-1**, Scolex with opened bothridia, No. *Heptranchias perlo* 23 (Host: *Heptranchias perlo*). **B-2**, Scolex with closed bothridia, **C**, Mature proglottid, No. *Heptranchias perlo* 20 (Host: *Heptranchias perlo*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 23 *Mitsukuricestus gobelinus* n. comb. **A**, Whole worm, No. *Miysukurina owstoni* 4 (Host: *Miysukurina owstoni*). **B**, Scolex, No. *Miysukurina owstoni* 1 (Host: *Miysukurina owstoni*). **C**, Mature proglottid, No. *Miysukurina owstoni* 4 (Host: *Miysukurina owstoni*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 24 *Monorygma chlamydoselachi*. **A**, Whole worm, No. *Chlamydoselachus anguineus* 1 (Host: *Chlamydoselachus anguineus*). **B**, Scolex, No. *Chlamydoselachus anguineus* 9 (Host: *Chlamydoselachus anguineus*). **C**, Mature proglottid, No. *Chlamydoselachus anguineus* 10 (Host: *Chlamydoselachus anguineus*). **D**, Egg, No. *Chlamydoselachus anguineus* 1 (Host: *Chlamydoselachus anguineus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 25 *Orygmatobothrium musteli*. **A**, Whole worm, No. *Mustelus manazo* 8 (Host: *Mustelus manazo*). **B**, Scolex, **C**, Mature proglottid, No. *Mustelus griseus* 20 (Host: *Mustelus griseus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 26 *Pelichinibothrium speciosum*. **A**, Whole worm, No. *Prionace glauca* 29 (Host: *Prionace glauca*). **B**, Mature proglottid, MPM 23879, SY3102 (Host: *Prionace glauca*). **C**, Free proglottid, MPM 23879, SY5624 (Host: *Prionace glauca*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 27 *Pelichinibothrium speciosum*. **A**, Whole worm, **B**, Mature proglottid, No. others 365 (Host: *Alepisaurus ferox*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 28 *Pelichinibothrium carcharodoni* n. comb. **A**, Whole worm, No. *Carcharodon carcharias* 19 (Host: *Carcharodon carcharias*). **B**, Mature proglottid, No. *Carcharodon carcharias* 15 (Host: *Carcharodon carcharias*).

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 29 *Pelichnibothrium caudatum*. **A**, Plerocercoid, No. *others* 512 (Host: *Onchorhynchus keta*). **B**, Plerocercoid, **C**, Plerocercoid, No. *others* 61 (Host: *Berryteuthis magister*). **D**, Plerocercoid, No. *others* 101 (Host: *Gadus macrocephalus*), **E**, Plerocercoid, arrow showed site which caused initial proglottization, **F**, Proglottids, No. *others* 68 (Host: *Aptocyclus ventricosus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 30 *Pelichinibothrium delphni* n. comb. Merocercoid, No. *others* 629 (Host: *Stenella coeruleoalba*).

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 31 *Pelichinibothrium grimaldii* n. comb. **A**, Merocercoid, **B**, Scolex, No. *others* 633 (Host: *Stenella coeruleoalba*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 32 *Pelichinibothrium montaukensis* n. comb. **A**, Whole worm, No. *Isurus oxyrinchus* 18 (Host: *Isurus oxyrinchus*). **B**, Scolex, No. *Isurus oxyrinchus* 1 (Host: *Isurus oxyrinchus*). **C**, Mature proglotiid, No. *Isurus oxyrinchus* 18 (Host: *Isurus oxyrinchus*). Overlapping vitelline follicles are not drawn on the right side of a broken line. **D**, Juvenil worm, No. *Isurus oxyrinchus* 25 (Host: *Isurus oxyrinchus*). **E**, Plerocercoid, No. *others* 343 (Host: *Alepisaurus ferox*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 33 *Pelichinibothrium tumidum* n. comb. **A**, Whole worm, No. *Carcharodon carcharias* 12 (Host: *Carcharodon carcharias*). **B**, Scolex, No. *Carcharodon carcharias* 7 (Host: *Carcharodon carcharias*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 34 *Scyphophyllidium carcharhinus* n. sp. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, No. *Carcharhinus limbatus* 14 (Host: *Carcharhinus limbatus*). **D**, Plerocercoid, No. *othres* 426 (Host: *Coryphaena hippurus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 35 *Scyphophyllidium eirakubuka* n. sp. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, No. *Hemitriakis japonica* 64 (Host: *Hemitriakis japonica*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 36 *Scyphophyllidium hanazame* n. sp. **A**, Whole worm, *Carcharhinus brevipinna* 11 (Host: *Carcharhinus brevipinna*). **B**, Scolex, No. *Carcharhinus brevipinna* 74 (Host: *Carcharhinus brevipinna*). **C**, Mature proglottid, No. *Carcharhinus brevipinna* 11 (Host: *Carcharhinus brevipinna*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 37 *Scyphophyllidium insulaeum* n. sp. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, No. *Carcharias galapagensis* 3 (Host: *Carcharias galapagensis*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 38 *Scyphophyllidium longiproglottidum* n. sp. **A**, Whole worm, **B**, Mature proglottid, No. *Carcharhinus limbatus* 6 (Host: *Carcharhinus limbatus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 39 *Scyphophyllidium maritimum* n. sp. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, No. *Carcharhinus brevipinna* 78 (Host: *Carcharhinus brevipinna*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 40 *Scyphophyllidium nemuribuka* n. sp. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, No. *Triaenodon obesus* 26 (Host: *Triaenodon obesus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 41 *Scyphophyllidium ogasawaraensis* n. sp. **A**, Whole worm, No. *Triaenodon obesus* 27 (Host: *Triaenodon obesus*). **B**, Scolex, **C**, Mature proglottid, No. *Triaenodon obesus* 6 (Host: *Triaenodon obesus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 42 *Scyphophyllidium paulum* n. com. **A**, Whole worm, No. *Galeocerdo cuvier* 7 (Host: *Galeocerdo cuvier*). **B**, Scolex, **C**, Mature proglottid, No. *Galeocerdo cuvier* 5 (Host: *Galeocerdo cuvier*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 43 *Scyphophyllidium prionacis* n. comb. **A**, Whole worm, No. *Prionace glauca* 24 (Host: *Prionace glauca*). **B**, Scolex, **C**, Mature proglottid, No. *Prionace glauca* 23 (Host: *Prionace glauca*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 44 *Scyphophyllidium sumitsukizame* n. sp. **A**, Whole worm, No. *Carcharhinus dussumieri* 14 (Host: *Carcharhinus dussumieri*). **B**, Scolex, No. *Carcharhinus dussumieri* 11 (Host: *Carcharhinus dussumieri*). **C**, Mature proglottid, No. *Carcharhinus dussumieri* 14 (Host: *Carcharhinus dussumieri*).

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 45 *Scyphophyllidium triacis* n. comb. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, No. *Triakis scyllium* 27 (Host: *Triakis scyllium*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 46 *Scyphophyllidium yogore* n. sp. **A**, Scolex, **B**, Mature proglottid, No. *Carcharhinus longimanus* 9 (Host: *Carcharhinus longimanus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 47 *Thysanocephalum thysanocephalum*. **A**, Whole worm, No. *Galeocerdo cuvier* 8 (Host: *Galeocerdo cuvier*). **B**, Scolex, No. *Galeocerdo cuvier* 1 (Host: *Galeocerdo cuvier*). **C**, Mature proglottid, No. *Galeocerdo cuvier* 8 (Host: *Galeocerdo cuvier*). Overlapping vitelline follicles are not drawn on the right side of a broken line. **D**, Egg, No. *Galeocerdo cuvier* 24 (Host: *Galeocerdo cuvier*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 48 *Vertebraevicestus dobukasube* n. sp. **A**, Whole worm, No. *Bathyraja smirnovi* 11 (Host: *Bathyraja smirnovi*). **B**, Scolex, No. *Bathyraja smirnovi* 13 (Host: *Bathyraja smirnovi*). **C**, Mature proglottid, No. *Bathyraja smirnovi* 11 (Host: *Bathyraja smirnovi*). Overlapping vitelline follicles are not drawn on the left side of a broken line. **D**, Eggs, No. *Bathyraja smirnovi* 13 (Host: *Bathyraja smirnovi*). **E**, Plerocercoid, No. *others* 34 (Host: *Lycodes nakamurae*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 49 *Yamaguticestus squali* n. comb. **A**, Whole worm, No. *Squalus japonicus* 11 (Host: *Squalus japonicus*). **B**, Scolex, No. *Squalus japonicus* 62 (Host: *Squalus mitsukurii*). **C**, Mature proglottid, **D**, Free proglottid, No. *Squalus japonicus* 11 (Host: *Squalus japonicus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 50 *Yamaguticestus squali* n. comb. **A**, Whole worm, No. *Cephaloscyllium umbratile* 56 (Host: *Cephaloscyllium umbratile*). **B**, Scolex, No. *Cephaloscyllium umbratile* 19 (Host: *Cephaloscyllium umbratile*). **C**, Mature proglottid, No. *Cephaloscyllium umbratile* 11 (Host: *Cephaloscyllium umbratile*). **D**, Eggs, No. *Cephaloscyllium umbratile* 5 (Host: *Cephaloscyllium umbratile*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 51 *Rosebothrium ootenjikuzame* n. sp. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, **D**, Microtriches in cirrus, No. *Nebrius ferrugineus* 2 (Host: *Nebrius ferrugineus*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 52 *Anthocephalum biacetabulatum* n. comb. **A**, Whole worm, **B**, Scolex, **C**, Mature proglottid, No. *Rhinobatos schlegelii* 31 (Host: *Rhinobatos schlegelii*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 53 Morphological terminology of larvae. Phyllobothriidean larvae, **A**, Plerocercoid of *Vertebraeovicestus dobukasube*; **B**, Plerocercoid of *Pelichnibothrium* cf. *montaukensis*; **C**, Merocercoid of *Pelichnibothrium delphini*. Tetraphyllidean larva, **D**, Plerocercoid of *Anthobothrium* sp. L3.

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 54 Maximum likelihood tree using Dataset 3 (lsrDNA). Bootstrap values and BPP are shown at each branch, and hyphens indicate branches which were not observed in the Bayesian Inference tree. Larval sequences obtained in this study are shown in red color, adult sequences obtained in this study are shown in black color, and those collected from GenBank are in gray color. Open boxes indicate the revised orders. Solid bars indicate species which identified by this study. *Pelichnibothrium* clade is shown by dotted lines.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 55 Maximum likelihood tree of *Pelichnibothrium* clade using Dataset 4 (lsrDNA). See Fig. 54 for the explanations of trees and open box. Solid bars indicate clades of each species clades.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 56 Maximum likelihood tree of *Pelichnibothrium* clade using Dataset 5 (mtDNA COI). See Fig. 54 for the explanations of trees and open box. Solid bars indicate clades of each species clades.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 57 Plerocercoids. **A.** *Pelichnibothrium* cf. *montaukensis*, No. *others* 52 (Host: *Todarodes pacificus*). **B.** Phyllobothriidae gen. sp. L1, No. *others* 577 (Host: *Xiphias gladius*). **C.** Phyllobothriidae gen. sp. L2, No. *others* 99 (Host: *Dasycottus setiger*). **D.** *Anthobothrium* sp. L3, No. *others* 390 (Host: *Trichiurus japonicus*), **E.** *Anthobothrium* sp. L4, No. *others* 430 (Host: *Coryphaena hippurus*). **F.** *Anthobothrium* sp. L5, No. *others* 132 (Host: *Epinephelus fasciatus*). **G.** *Anthobothrium* sp. L5, No. *others* 553 (Host: *Coris aygula*).

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 58 Morphological variation of the scolex of plerocercoids of *Pelichnibothrium caudatum*. **A**, flatted bothridium, No. *others* 48 (Host: *Hippoglossoides dubius*). **B**, flatted bothridium, No. *others* 29 (Host: *Careproctus trachysoma*). **C**, slightly crumpled bothridium, No. *others* 22 (Host: *Aptocyclus ventricosus*).

本図については、5年以内に雑誌等で刊行予定のため、非公開。

Fig. 59 Invagination of scolex of *Pelichnibothrium caudatum*, No. others 527 (Host: *Oncchorhynchus keta*). Pictures were arranged in time series from left to right. Drawings were given to show outlines of the body and organs for pictures of 0 sec and 30 sec.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 60 Life cycle of *Pelichnibothrium montaukensis*. Arrows show the flow of life cycle, black arrows show the free-living flows, red arrows show host transportation by food web. Squares show host, life stages in the square are a parasitic stage.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 61 Life cycle of *Pelichnibothrium* cf. *montaukensis*. The information was followed as Fig. 60.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 62 Life cycle of *Pelichnibothrium carcharodoni*. The information was followed as Fig. 60.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 63 Life cycle of *Pelichnibothrium speciosum*. The other information was followed as Fig. 60.

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 64 Life cycle of *Scyphophyllidium carcharhinus*. The other information was followed as Fig. 60.

本図については、5年以内に雑誌等で刊行予定のため、非公開.

Fig. 65 Life cycle of *Vertebraeovicestus dobukasube*. The other information was followed as Fig. 60.

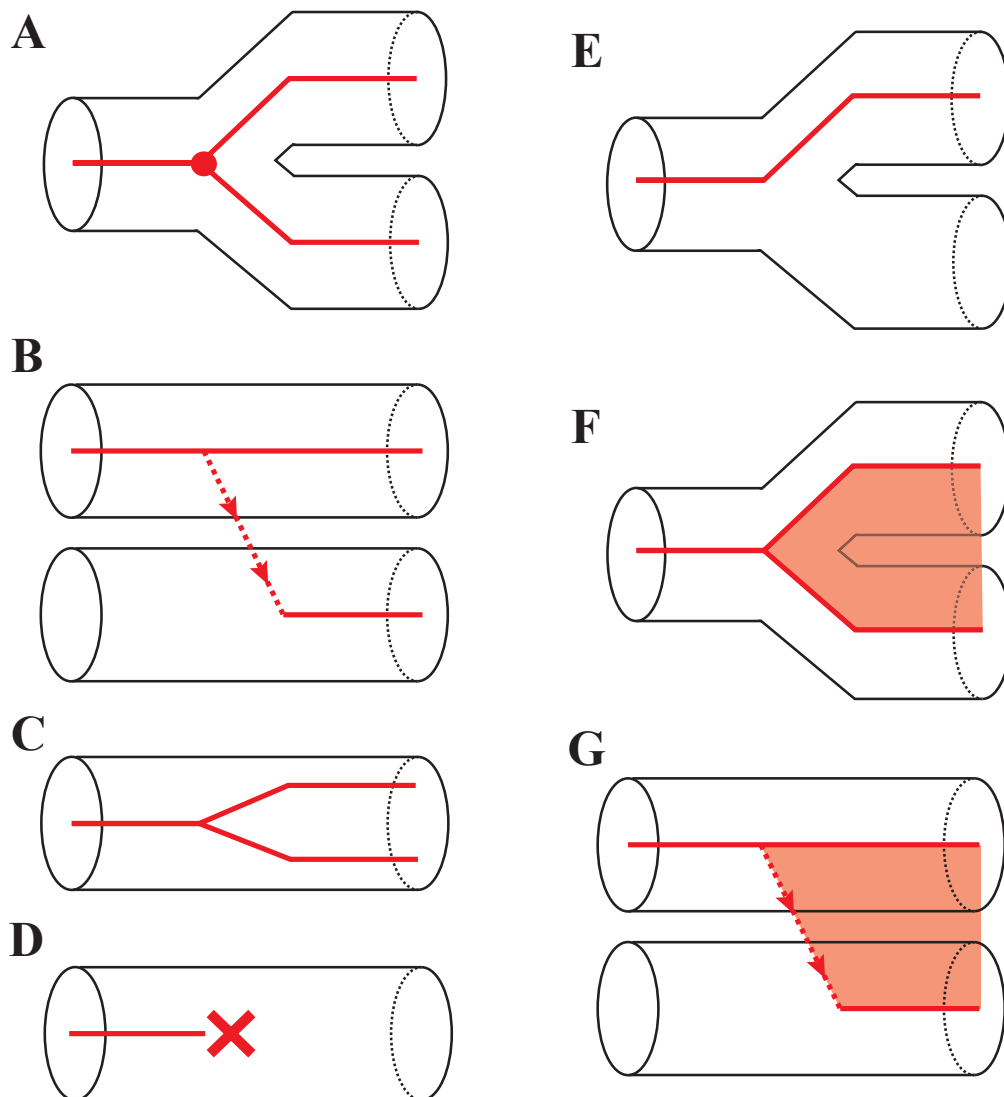


Fig. 66 Evolutionally processes. **A.** co-speciation, **B.** host switch, **C.** independent speciation, **D.** extinction, **E.** missing the boat, **F.** failure to speciation, **G.** host range expansion. Black tubes represent host lineages, red lines represent parasite lineages. A circle shows co-speciation, arrows and dotted lines show host switching, and a cross shows extinction. This figures were modified from Page (2003) and Clayton et al. (2003).

本図については、5年以内に雑誌等で刊行予定のため、非公開。

Fig. 67 Relationship between phyllobothriidean lineages and their host elasmobranch lineages. The phylogenetic tree of phyllobothriideans was based on Fig. 14. The phylogenetic tree of elasmobranch families was based on Naylor et al. (2012). Single asterisks mean polyphyletic families, and double asterisks mean paraphyletic families. Lines show parasite-host associations, and their color shows subclasses of the host families: black, Holocephali; blue, Galeomorphi; green, Batoidea; red, Squalomorphi.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 68 Relationship between *Pelichnibothrium* lineages and their host's lineages. Phylogenetic tree of *Pelichnibothrium* was based on Fig. 55. Phylogenetic tree of their host elasmobranchs was based on Naylor et al. (2012). Single asterisks mean polyphyletic families. Lines show parasite-host relationship.

本図については，5年以内に雑誌等で刊行予定のため，非公開.

Fig. 69 Estimation of co-evolutionary history between *Pelichnibothrium* and their host shark species. See Fig. 66 for the explanations of symbols and lines in the tree. Species in red show parasite, and species in black color show host.

本図については、5年以内に雑誌等で刊行予定のため、非公開。

Fig. 70 Relationship between *Alexanderecestus*, *Scyphophyllidium* and *Thysanoccephalum* lineages and their host elasmobranch lineages. The phylogenetic tree of the three genera was based on Fig. 14. The phylogenetic tree of elasmobranchs was based on Naylor et al. (2012). See Fig. 67 for the explanations of symbols in the tree. Red color Phyllobothriidean species in red color indicates species parasitic in multiple hosts. Lines show the relationship between phyllobothriidean and carcharhinid sharks.

Appendix History of systematics of the order Phyllobothriidea and Tetrephyllidea. The genera included in Phyllobothriidea in this study are shown in bold face.

Braum (1900)

Order Tetrephyllidea

Family Onchobothriidae

Onchobothrius, *Acanthobothrium*, ***Calliobothrium***, *Ceratobothrium*,
Cylindrophorus, *Phoreiobothrium*, *Platybothrium*, *Prosthecobothrium*,
Thysanocephalum

Family Phyllobothriidae

Phyllobothrium, *Anthobothrium*, ***Calyptrbothrium***, ***Crossobothrium***,
Dinobothrium, *Diplobothrium*, *Echeneibothrium*, ***Monorygma***,
Orygmatobothrium, *Spongiobothrium*, ***Trilocularia***, *Tritaphros*

Family Lecanicephalidae

Discocephalum, *Lecanicephalum*, *Tylocephalum*

Family Ichthyoteaniidae

Ichthyotaenia, *Corallobothrium*, *Crepidobothrium*

Meggitt (1924)

Order Tetrephyllidea

Family Ichthyoteaniidae

Ichthyoteania, *Corallobothrim*, *Crepidobothrium*, *Gangesia*

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, *Balanobothrium*, ***Calliobothrium***,
Ceratobothrium, *Cylindrophorus*, *Pedibothrium*, *Phyllobothroides*,
Platybothrium, *Prosthecobothrium*, ***Thysanocephalum***

Family Phyllobothriidae

Phyllobothrium, *Anthobothrium*, *Aocobothrium*, ***Bilocularia***,
Calyptrbothrium, *Carpobothrium*, *Dinobothrium*, *Diplobothrium*,
Echeneibothrium, *Eniochobothrium*, *Hornellobothrium*, ***Monorygma***,
Myzocephalus, *Myzophyllobothrium*, *Oriana*, ***Orygmatobothrium***,
Pelichnibothrium, *Peltidocotyle*, *Prosobothrium*, *Rhinebothrium*,
Rhoptrobobothrium, *Spongiobothrium*, *Tiarabothrium*, ***Trilocularia***,
Tritaphros, *Zygobothrium*

Family Lecanicephalidae

Anthemobothrium, *Calycobothrium*, *Polypocephalus*

Family Polypocephalidae

Polypocephalus, *Anthemobothrium*, *Calycobothrium*

Southwell (1925)

Order Tetraphyllidea

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, ***Calliobothrium***, *Ceratobothrium*,
Cylindrophorus, *Uncibilocularis*, *Platybothrium*, *Pedibothrium*,
Tetracampos, ***Thysanocephalum***

Family Phyllobothriidae

Phyllobothrium, *Anthobothrium*, *Aocobothrium*, *Carpobothrium*,
Echeneibothrium, *Myzophyllobothrium*, ***Orygmatobothrium***, *Pithophorus*

Fuhrmann (1931)

Order Tetraphyllidea

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, *Balanobothrium*, ***Calliobothrium***,
Cylindrophorus, *Pedibothrium*, *Platybothrium*, *Spiniloculus*,
Thysanocephalum *Uncibilocularis*, *Yorkeria*

Family Phyllobothriidae

Phyllobothrium, *Anthobothrium*, ***Bilocularia***, ***Calyptrbothrium***,
Carpobothrium, *Ceratobothrium*, *Dinobothrium*, *Echeneibothrium*,
Myzophyllobothrium, ***Orygmatobothrium***, ***Scyphophyllidium***, ***Trilocularia***,
Tritaphros

Family Lecanicephalidae

Lecanicephalum, *Adelobothrium*, *Anthemobothrium*, *Polypocephalus*

Family Monticelliidae

Monticellia, *Goezeella*, *Ephedrocephalus*, *Marsypocephalus*, *Peltidocotyle*

Family Proteocephalidae

Proteocephalus, *Acanthoteania*, *Corallobotilrium*, *Crepidobothrium*,
Ophioteania

Family Cephalobothriidae

Cephalobothrium, *Tylocephalum*, *Discobothrium*

Family Discocephalidae

Discocephalum

Joyeux & Baer (1936)

Order Tetracystida

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, ***Calliobothrium***, *Cylindrophorus*

Family Phyllobothriidae

Phyllobothrium, *Anthobothrium*, *Ceratobothrium*, *Dinobothrium*,
Echeneibothrium, ***Orymatobothrium***, *Prosobothrium*, ***Scyphophyllidium***

Family Ichthyotaeniidae

Ichthyotaenia, *Ophiotaenia*

Family Cephalobothriidae

Discobothrium

Family Disculicipitidae

Disculiceps

Hyman (1951)

Order Tetracystida

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, ***Calliobothrium***, *Thysanocephalum*

Family Phyllobothriidae

Phyllobothrium, *Anthobothrium*, *Carpobothrium*, *Dinobothrium*,
Echeneibothrium, *Myzophyllobothrium*, ***Orymatobothrium***, ***Trilocularia***

Wardle & McLeod (1952)

Order Tetracystida

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, ***Calliobothrium***, *Thysanocephalum*

Family Phyllobothriidae

Phyllobothrium, *Anthobothrium*, *Carpobothrium*, *Dinobothrium*,
Echeneibothrium, *Mysophorus*, *Myzophyllobothrium*, ***Orymatobothrium***,
Pithophorus, ***Pelichnibothrium***, ***Scyphophyllidium***

Euzet (1959)

Order Tetrephyllidea

Superfamily Phyllobothrioidea

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, ***Calliobothrium***

Family Phyllobothriidae

Subfamily Phyllobothriinae

Phyllobothrium, *Anthobothrium*, ***Calyptrbothrium***, *Ceratobothrium*,
Crossobothrium, *Dinobothrium*, ***Monorygma***, ***Orygmatobothrium***,
Scyphophyllidium, *Sphaerobothrium*

Subfamily Echeneibothriinae

Echeneibothrium, *Clydonobothrium*, *Tritaphros*

Subfamily Rhinebothriinae

Rhinebothrium, *Caulobothrium*, *Rhabdotobothrium*

Subfamily Thysanocephalinae

Thysanocephalum

Superfamily Prosobothrioidea

Family Prosobothriidae

Subfamily Prosobothriinae

Prosobothrium

Subfamily Platybothriinae

Platybothrium

Family Phoreiobothriidae

Subfamily Phoreiobothriinae

Phoreiobothrium

Subfamily Reesiinae

Reesium, ***Trilocularia***

Family Gastrolecithidae

Gastrolecithus

Superfamily Lecanicephaloidea

Family Lecanicephalidae

Lecanicephalum, *Tetragonocephalum*

Family Disculicipitidae

Disculiceps

Family Cephalobothriidae

Cephalobothrium

Yamaguti (1959)

Order Tetraphyllidea

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, ***Calliobothrium***, *Ceratobothrium*,
Cylindrophorus, *Phoreiobothrium*, *Pinguicollum*, *Platybothrium*,
Spiniloculus, ***Thysanocephalum***, *Uncibilocularis*, *Yorkeria*

Family Phyllobothriidae

Phyllobothrium, *Anthobothrium*, *Aocobothrium*, *Carpobothrium*,
Cyatocotyle, *Dinobothrium*, *Echeneibothrium*, *Gastrolecithus*,
Marsupiobothrium, ***Monorygma***, ***Orygmatobothrium***, ***Pelichnibothrium***,
Pithophorus, *Pseudanthobothrium*, *Reesium*, *Rhodobothrium*,
Scyphophyllidium

Family Maccallumiellidae

Maccallumiella

Family Triloculariidae

Trilocularia

Joyeux & Baer (1961)

Order Tetraphyllidea

Superfamily Phyllobothrioidea

Family Onchobothriidae

Onchobothrium, *Acanthobothrium*, *Balanobothrium*, ***Calliobothrium***,
Pediobothrium, *Pinguicollum*, *Spiniloculus*, *Uncibilocularis*, *Yorkeria*

Family Phyllobothriidae

Subfamily Phyllobothriinae

Phyllobothrium, *Anthobothrium*, ***Calyptrbothrium***, *Carpobothrium*,
Ceratobothrium, ***Crossobothrium***, *Dinobothrium*, *Inermiphyllidium*,
Marsupiobothrium, ***Monorygma***, ***Orygmatobothrium***, *Pillersium*,
Pithophorus, *Polypobothrium*, ***Scyphophyllidium***, *Sphaerobothrium*,
Staurobothrium

Subfamily Echeneibothriinae

Clydonobothrium, *Echeneibothrium*, *Pseudanthobothrium*, *Tritaphros*

Subfamily Rhinebothriinae

Caulobothrium, *Rhabdotobothrium*, *Rhinebothrium*, *Tiarabothrium*

Subfamily Thysanocephalinae

Myzocephalus, *Myzophyllobothrium*, *Rhoptrobothrium*, ***Thysanocephalum***

Superfamily Lecanicephaloidea

Family Lecanicephalidae

Anthemobothrium, Calycobothrium, Lecanocephalum, Polypocephalus

Family Cephalobothriidae

Adelobothrium, Cephalobothrium, Hexacanalus, Tetragonocephalum

Superfamily Prosobothrioidea

Family Disculicepitidae

Disculiceps

Family Gastrolecithidae

Gastrolecithus

Family Phoreiobothriidae

Subfamily Phoreiobothiinae

Phoreiobothrium

Subfamily Reesiinae

*Reesium, **Trilocularia***

Family Prosobothiidae

Subfamily Platybothriinae

Dicranobothrium, Platybothrium

Schmidt (1986)

Order Tetraphyllidea

Family Onchobothriidae

*Onchobothrium, Acanthobothroides, Acanthobothrium, **Calliobothrium**,
Ceratobothrium, Cyliodrophorus, Pachybothrium, Pedibothrium,
Phoreiobothrium, Pinguicollum, Platybothrium, Pomatotrygonocetus,
Spiniloculus, **Thysanocephalum**, Uncibilocularis, Yorkeria*

Family Phyllobothriidae

***Phyllobothrium**, Anthobothrium, Aocobothrium, Carpobothrium,
Caulobothrium, Clydonobothrium, Cyatocotyle, Dinobothrium,
Duplicibothrium, Echeneibothrium, Gastrolecithus, Glyphobothrium,
Marsupiobothrium, Mixophyllobothrium, **Monorygma**, Myzophyllobothrium,
Orygmatobothrium, **Pelichnibothrium**, Phormobothrium, Pithophorus,
Pseudanthobothrium, Rhabdotobothrium, Rhinebothrium, Rhinebothroides,
Rhodobothrium, Reesium, **Scyphophyllidium***

Family Triloculariidae

***Trilocularia**, Pentaloculum, Zyxibothrium*

Family Cathetocephalidae

Cathetocephalus

Euzet (1994)

Order Tetraphyllidea

Family Onchobothriidae

Onchobothrium, *Acanthobothroides*, *Acanthobothrium*, *Balanobothrium*,
Calliobothrium, *Dicranobothrium*, *Megalonchos*, *Pachybothrium*,
Pedibothrium, *Phoreiobothrium*, *Platybothrium*, *Pomatotrygonocetus*,
Spiniloculus, *Uncibilocularis*, *Yorkeria*

Family Phyllobothriidae

Subfamily Thysanocephalinae

Thysanocephalum, *Myzocephalus*, *Myzophyllobothrium*

Subfamily Echeneibothriinae

Pseudoanthobothrium, *Clydonobothrium*, *Phormbothrium*, *Tritaphros*,
Echeneibothrium

Subfamily Phyllobothriinae

Phyllobothrium, *Anthobothrium*, *Calyptrbothrium*, *Carpobothrium*,
Ceratobothrium, *Clistobothrium*, *Crossobothrium*, *Dinobothrium*,
Gastrolecithus, *Marsupiobothrium*, ***Monorygma***, ***Orygmatobothrium***,
Pithophorus, *Rhodobothrium*, ***Scyphophyllidium***

Subfamily Triloculariinae

Trilocularia, *Pentaloculum*, *Zyxibothrium*

Subfamily Rhinebothriinae

Rhinebothrium, *Caulobothrium*, *Duplicibothrium*, *Glyphobothrium*,
Rhabdotobothrium, *Rhinebothroides*

Family Disculidpitidae

Disculiceps

Family Prosobothriidae

Prosohothrium

Family Dioecotaeniidae

Dioecotaenia

Family Litobothriidae

Litobothrium

Family Cathetocephalidae

Cathetocephalus

Family Chimaerocestidae

Chimaerocestos

Caira et al. (2014)

Order Phyllobothriidea

Family Phyllobothriidae

Phyllobothrium, *Calyptrbothrium*, *Chimaerocestos*, *Marsupiobothrium*,
Nandocestus, *Orectolobicestus*, *Orygmatobothrium*, *Paraorygmatobothrium*,
Ruhnkecestus, *Scyphophyllidium*, *Thysanocephalum*

Order Tetracophyllidea

incertae sedis

Alexandercestus, *Anthobothrium*, *Balanobothrium*, *Bibursibothrium*,
Biloculuncus, *Calliobothrium*, *Caulobothrium*, *Carpobothrium*,
Cardiobothrium, *Ceratobothrium*, *Clistobothrium*, *Crossobothrium*,
Dinobothrium, *Dioecotaenia*, *Duplicibothrium*, *Erudituncus*, *Flexibothrium*,
Megalonchos, *Myzocephalus*, *Myzophyllobothrium*, *Pachybothrium*,
Pedibothrium, *Pelichnibothrium*, *Rhoptrobothrium*, *Spiniloculus*,
Trilocularia, *Yorkeria*

This study

付録については、5年以内に雑誌等で刊行予定のため、非公開。

Appendix Table 1 Revised nominal species of Phyllobothriidea. Taxonomic status excepted species using in this study were given from Ruhnke (2011) and followed descriptions.

付録については，5年以内に雑誌等で刊行予定のため，非公開.

Appendix Table 1 continued.

付録については，5年以内に雑誌等で刊行予定のため，非公開.

Appendix Table 1 continued.

付録については，5年以内に雑誌等で刊行予定のため，非公開.

Appendix Table 1 continued.

付録については，5年以内に雑誌等で刊行予定のため，非公開.

付録については、5年以内に雑誌等で刊行予定のため、非公開.

Appendix Fig. 1 Bayesian Inference tree of Analysis 7 (Dataset 1, ssrDNA+lssrDNA). BPP were showed upper orders clade. Current scientific names are used and some of them were revised in this study (see Table 15). Sequences obtained in this study are shown in black color, and those collected from GenBank are in gray color. Open boxes indicate the revised orders in this study (see Section 3.3 of Chapter I).

付録については，5年以内に雑誌等で刊行予定のため，非公開.

Appendix Fig. 2 Bayesian Inference tree of Analysis 8 (Dataset 1, ssrDNA). See Appendix Fig. 1 for the explanations of symbols in the tree. Phyllobothriidea, Tetraphyllidea and Onchoproteocephalidea were polyphyletic in this analysis (connected by dotted lines).

付録については、5年以内に雑誌等で刊行予定のため、非公開.

Appendix Fig. 3 Bayesian Inference tree of Analysis 9 (Dataset 1, lsrDNA). See Appendix Fig. 1 for the explanations of symbols in the tree. Phyllobothriidea were polyphyletic in this analysis (connected by dotted lines).

付録については，5年以内に雑誌等で刊行予定のため，非公開.

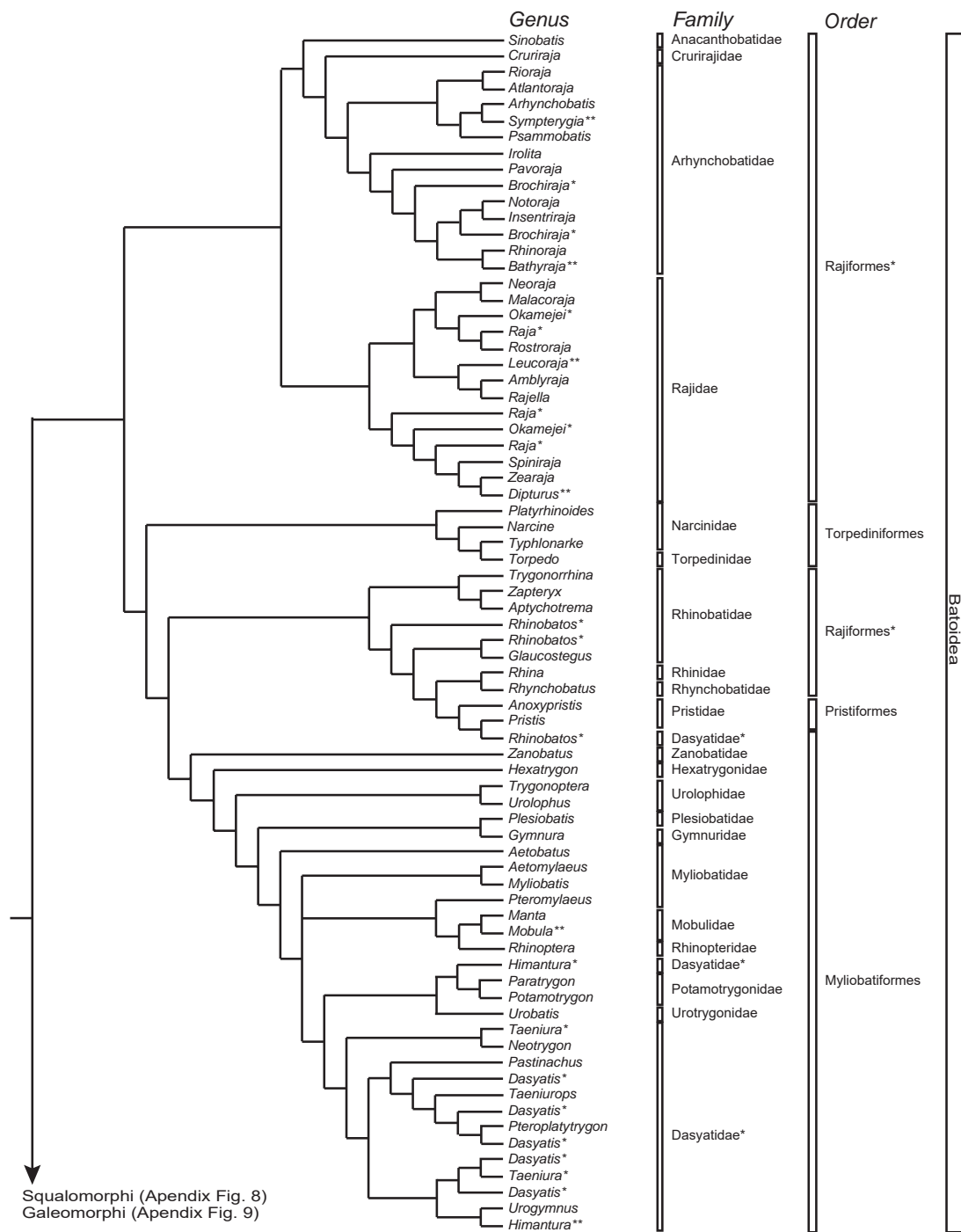
Appendix Fig. 4 Bayesian Inference tree of Analysis 10 (Dataset 2, ssrDNA and lsrDNA). See Appendix Fig. 1 for the explanations of symbols in the tree.

付録については，5年以内に雑誌等で刊行予定のため，非公開.

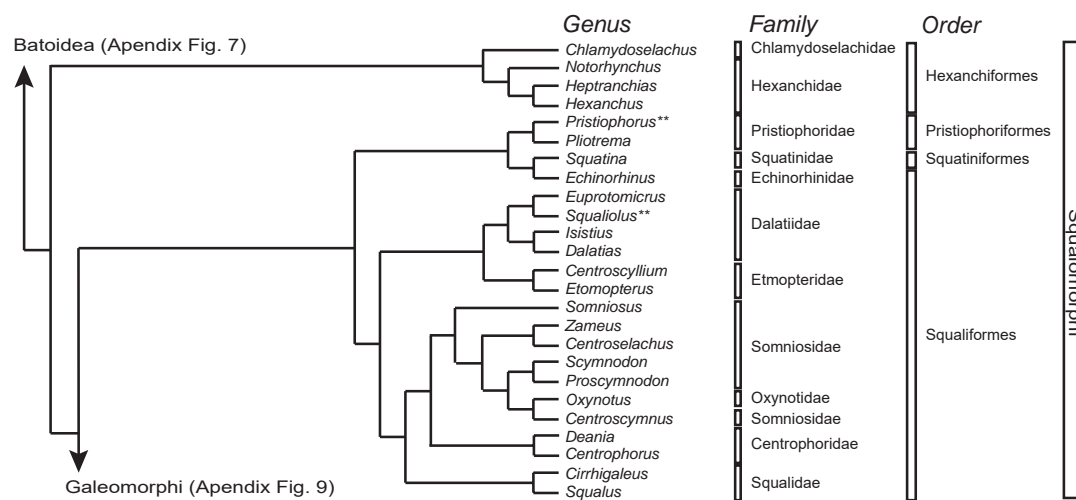
Appendix Fig. 5 Bayesian Inference tree of Analysis 11 (Dataset 2, ssrDNA). See Appendix Fig. 1 for the explanations of symbols in the tree. Phyllobothriidea were polyphyletic in this analysis (connected by dotted lines).

付録については，5年以内に雑誌等で刊行予定のため，非公開.

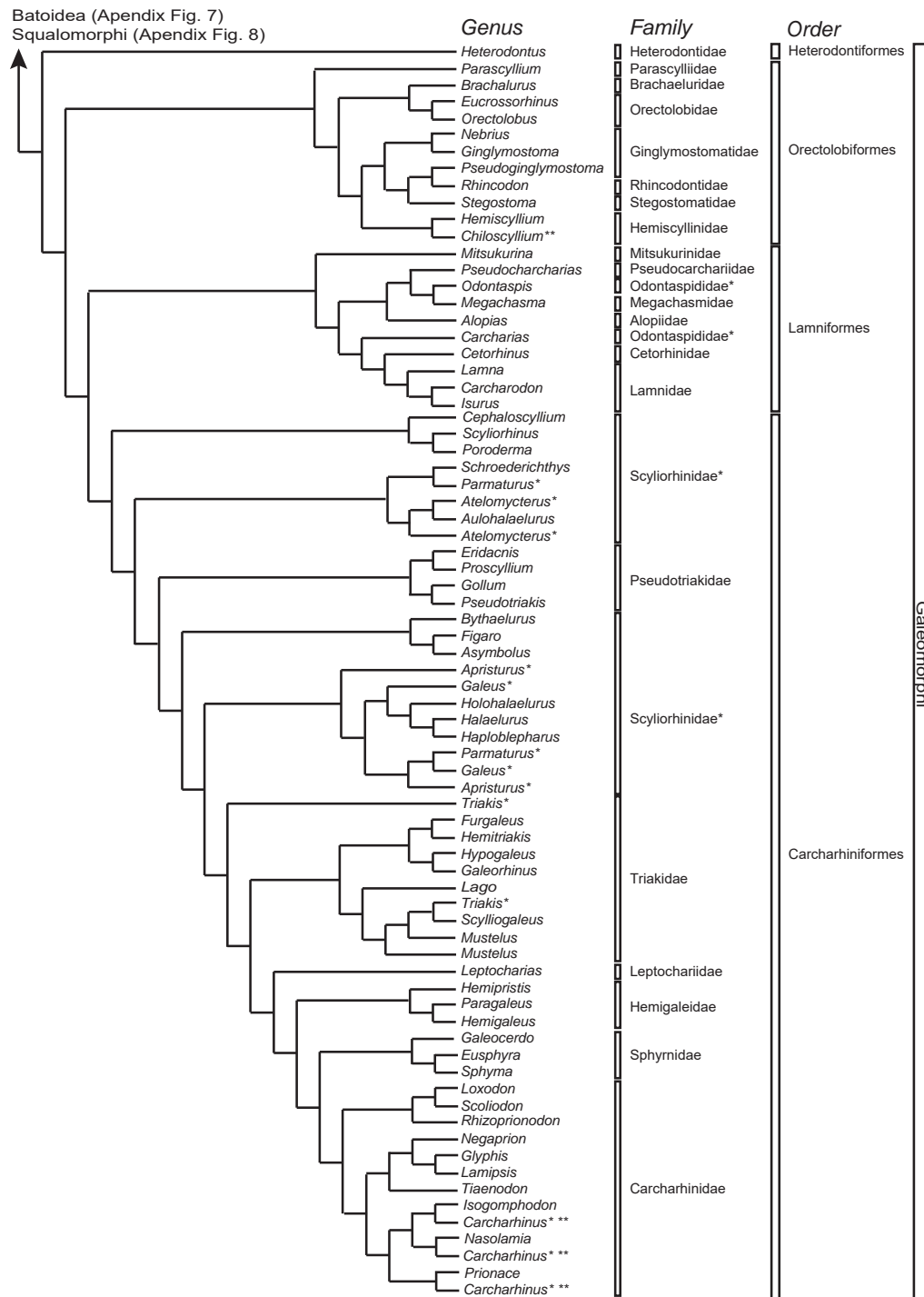
Appendix Fig. 6 Bayesian Inference tree of Analysis 12 (Dataset 2, lsrDNA). See Appendix Fig. 1 for the explanations of symbols in the tree. Phyllobothriidea were polyphyletic in this analysis (connected by dotted lines).



Appendix Fig. 7 Phylogenetic tree of Batoidea of Elasmobranchii (Naylor et al., 2012). Single asterisks mean polyphyletic taxa, and double asterisks mean paraphyletic taxa.



Appendix Fig. 8 Phylogenetic tree of Squalomorphi of Elasmobranchii (Naylor et al., 2012). Single asterisks mean polyphyletic taxa, and double asterisks mean paraphyletic taxa.



Appendix Fig. 9 Phylogenetic tree of Galeomorphi of Elasmobranchii (Naylor et al., 2012). Single asterisks mean polyphyletic taxa, and double asterisks mean paraphyletic taxa.