

# A new species of *Haliotrema* (Monogenea: Ancyrocephalidae (*sensu lato*) Bychowsky & Nagibina, 1968) from holocentrids off Langkawi Island, Malaysia with notes on the phylogeny of related *Haliotrema* species

Soo O.Y.M.<sup>a,b,\*</sup>

<sup>a</sup> UCSI University KL, No.1, Jalan Menara Gading, Taman Connaught 56000 Cheras, Kuala Lumpur, Malaysia

<sup>b</sup> Institute of Biological Sciences, Faculty of Science, University of Malaya, Kuala Lumpur 50603, Malaysia



## ARTICLE INFO

**Keywords:**  
Monogenea  
Holocentrids  
*Haliotrema*  
28S rDNA  
Malaysia

## ABSTRACT

*Haliotrema susanae* sp. nov. is described from the gills of the pinecone soldierfish, *Myripristis murdjan* off Langkawi Island, Malaysia. This species is differentiated from other *Haliotrema* species especially those from holocentrids in having a male copulatory organ with bract-like extensions at the initial of the copulatory tube, grooved dorsal anchors and ventral anchors with longer shafts. The maximum likelihood (ML) analysis based on partial 28S rDNA sequences of *H. susanae* sp. nov. and 47 closely related monogeneans showed that *H. susanae* sp. nov. is recovered within a monophyletic clade consisting of only species from the genus *Haliotrema*. It is also observed that *H. susanae* sp. nov. forms a clade with *H. cromileptis* and *H. epinepheli* which coincides with a similar grouping by Young based on solely morphological characteristics. The morphological and molecular results validate the identity of *H. susanae* sp. nov. as belonging to the genus *Haliotrema*.

## 1. Introduction

The family Holocentridae (squirrelfish and soldierfish) consists of 84 species globally with 13 species recorded in Malaysia from three genera (*Sargocentron*, *Myripristis* and *Ostichthys*) [1]. To date, species of four genera of monogeneans have been reported from holocentrids; *Benedenia* Diesing, 1858, *Enoplocotyle* Tagliani, 1912, *Pseudohaliotrema* Yamaguti, 1953 (taxonomic status is *incertae sedis*) and *Haliotrema* Johnston & Tiegs, 1922, with the largest number of described species coming from the latter genus (Table 1). During a field trip in 2011, three species of holocentrid fish (*Myripristis murdjan* Forsskål, 1775, *M. hexagona* Lacépède, 1802 and *Sargocentron rubrum* Forsskål, 1775) were examined for monogeneans. A new monogenean species was discovered from *M. murdjan* and its taxonomy is described herein. This new species is not only the first described *Haliotrema* species from Malaysia but is the first study on holocentrids from Malaysian waters. Molecular information from the new species is further analysed with molecular information from related *Haliotrema* species to confirm its validity.

## 2. Materials and methods

The fish hosts, *Myripristis murdjan*, *M. hexagona* and *Sargocentron*

*rubrum* were collected in the coastal waters off the island of Langkawi (6°21'N, 99°46'E) in March 2011. Monogeneans were collected from excised gills of freshly killed or frozen fish and prepared for morphological investigation. Briefly, the monogeneans were collected from the gills, pipetted onto clean glass slides, covered with cover slips and cleared in modified ammonium-picrate-glycerin [2]. These ammonium-picrate-glycerin specimens were later washed and dehydrated through a graded ethanol series and mounted in Canada balsam without staining [2] to study the sclerotized hard parts. To study the soft anatomical structures, some specimens were immediately fixed after collection, in AFA (acetic acid-formalin-alcohol) onto glass slides, then stained in Gomori's triple stain and mounted in Canada balsam after dehydration in increasing ethanol series [3]. Stained and unstained specimens were studied under bright-field and phase contrast microscopes. Images of the hard and soft anatomical structures of the new species were captured using a Leica digital camera and an image analysis software (QWin Plus). The hard and soft parts were illustrated using a digitizing tablet (WACOM) and Adobe Illustrator software (Fig. 1). The specimens were measured as indicated in Fig. 1 and measurements given in micrometres as an average with the ranges in parentheses. Type-specimens of the new species were deposited at the Zoological Reference Collection, Lee Kong Chian Natural History Museum, National University of Singapore, Singapore (ZRC) and Natural History Museum

\* Corresponding author at: UCSI University KL, No.1, Jalan Menara Gading, Taman Connaught 56000 Cheras, Kuala Lumpur, Malaysia.

E-mail address: [michellesoo@ucsiuniversity.edu.my](mailto:michellesoo@ucsiuniversity.edu.my).

<https://doi.org/10.1016/j.parint.2018.09.003>

Received 2 March 2018; Received in revised form 15 August 2018; Accepted 17 September 2018

Available online 20 September 2018

1383-5769/ © 2018 Elsevier B.V. All rights reserved.

**Table 1**  
Monogeneans from fish of the Family Holocentridae.

Monogenean	Host	Localities	Present status	Authority
<i>Haliotrema auribaculum</i> Zhukov, 1980	<i>Holocentrus adscensionis</i> Osbeck (formerly known as <i>Holocentrus ascensionis</i> )	Gulf of Mexico	Valid	Zhukov, 1980
<i>Haliotrema chelicirrus</i> Yamaguti, 1968	<i>Neoniphon aureolineatus</i> Liénard (formerly known as <i>Holocentrus scythrops</i> ), <i>N. samara</i> Forsskål, <i>Sargocentron punctatissimum</i> Cuvier (formerly known as <i>Holocentrus lacteoguttatus</i> )	Hawaii	Valid	Yamaguti, 1968
<i>Haliotrema curvicirrus</i> Zhukov, 1980 (questionable)	<i>Holocentrus adscensionis</i> Osbeck (formerly known as <i>Holocentrus ascensionis</i> )	Gulf of Mexico	Valid	Zhukov, 1980
<i>Haliotrema holocentri</i> Young, 1968	<i>Sargocentron rubrum</i> Forsskål	Heron Island, Queensland, Australia	Valid	Young, 1968
<i>Haliotrema longirectocirrus</i> Zhukov, 1980	<i>Holocentrus adscensionis</i> Osbeck (formerly known as <i>Holocentrus ascensionis</i> )	Gulf of Mexico	Valid	Zhukov, 1980
<i>Haliotrema myrpiristisi</i> Zhukov, 1980	<i>Myrpiristis jacobus</i> Cuvier	Gulf of Mexico	Valid	Zhukov, 1980
<i>Haliotrema papillibaculum</i> Zhukov, 1980	<i>Myrpiristis jacobus</i> Cuvier	Gulf of Mexico	Valid	Zhukov, 1980
<i>Haliotrema tenuhamus</i> Zhukov, 1980	<i>Myrpiristis jacobus</i> Cuvier	Gulf of Mexico	Valid	Zhukov, 1980
<i>Benedenia hawaiiensis</i> Yamaguti, 1968	<i>Neoniphon aureolineatus</i> Liénard (formerly known as <i>Holocentrus scythrops</i> )	Hawaii	Valid	Yamaguti, 1968
<i>Enoplocoryle hawaiiensis</i> Yamaguti, 1968	<i>Neoniphon aureolineatus</i> Liénard (formerly known as <i>Holocentrus scythrops</i> )	Hawaii	Valid	Yamaguti, 1968
<i>Pseudohaliotrema</i> <i>incertae sedis</i> <i>falcatus</i> Yamaguti, 1968	<i>Sargocentron spiniferum</i> Forsskål	Hawaii	<i>Incertae sedis</i>	Yamaguti, 1968

London (NHMUK). The following material was examined: paratypes, *Haliotrema banana* Lim & Justine, 2007 (MZUM 578–580). The author was not able to obtain Zhukov's type specimens (from holocentrid hosts) because the slides cannot be taken out of the museum (Gerasev, pers. comm., 2017). Monogeneans collected from *S. rubrum* were found to be *Haliotrema holocentri* Young, 1968 but a redescription was not carried out as there weren't enough specimens for a revision.

For extraction of molecular information, the monogenean worms were removed from the gills of the hosts, identified and preserved in 75% ethanol. Genomic DNA was then extracted using DNEasy extraction kit from QIAGEN. The extracted DNA (5 µl) was used as a template in the PCR reaction to amplify the partial D1-D2 domain of the 28S rDNA, using forward primers C1 (5'-ACCCGCTGAATTTAAGCAT-3') [4] and reverse primer D2 (5'-TGGTCCGTGTTTCAAGAC-3') [5]. The PCR reaction (50 µl) was performed in 1.5 mM MgCl<sub>2</sub>, PCR buffer (Fermentas), 200 µM of each deoxyribonucleotide triphosphate, 1.0 µM of each PCR primer and 1 U of *Taq* polymerase (Fermentas) in a thermocycler (Eppendorf Mastercycler) using the following conditions: an initial denaturation at 95 °C for 4 min, followed by 35 cycles of 95 °C for 1 min, 50 °C for 1 min and 72 °C for 1 min, followed by a final extension at 72 °C for 10 min. An aliquot (10 µl) from the amplicon was electrophoresed in a 1.3% agarose gel, stained with ethidium bromide and viewed under a UV illuminator. The remaining 40 µl of each amplicon was purified using a DNA purification kit (QIAGEN) and subjected to automated DNA sequencing (ABI 3730 DNA Sequencer, First Base Laboratories, KL) using the same primers used for PCR amplification. Partial 28S rDNA sequence data of the new species and selected monogeneans from the GenBank (Table 2) were analysed with *Calydiscoides indianus* Oliver, 1987 (from GenBank) as the outgroup species. It should be noted that the names of the species used for building the phylogenetic tree follow those in the World Register of Marine Species (WoRMS) [6] and not Genbank, where the species names have not been updated. These DNA sequences were edited and aligned with MAFFT v7.164 [7] and verified/edited visually using BioEdit v7.0.5.3 [8]. Maximum likelihood (ML) analysis was performed with MEGA7 [9]. Bootstrap procedure with 1000 replications was performed to assess the robustness of the inferred relationships.

### 3. Results

#### 3.1. Description of new species

##### **Monogenea (Van Beneden, 1858) Bychowsky, 1937**

##### **Dactylogyridea Bychowsky, 1937**

##### **Ancyrocephalidae (sensu lato) Bychowsky & Nagibina, 1968**

##### ***Haliotrema susanae* sp. nov. (Fig. 1).**

Body elongate, length 569 µm (445–708), width 159 µm (118–249), anterior region with 2 pairs of pigmented eye spots. Mouth ventral; intestine bifurcates posterior to pharynx. Haptor set off from body, almost rectangular; length 90 µm (68–113) width 123 µm (86–181), 14 similar sized marginal hooks, length 12 µm (10–14); two pairs of anchors; two dorsal anchors with inner root 24 µm (20–28); outer root 10 µm (7–12); inner length 59 µm (50–65); outer length 47 µm (43–50); point 13 µm (11–15); two ventral anchors with inner root 22 µm (18–25); outer root 11 µm (8–13); inner length 54 µm (48–61); outer length 46 µm (41–49); point 13 µm (10–16); two connecting bars; slightly bent dorsal bar, 44 µm (28–51) wide; V-shaped ventral bar with enlarged ends, 46 µm (37–56).

Testis single, roundish; vas deferens sinistral, loops around left caecal branch, ascends diagonally in intercaecal region, distends to form seminal vesicle, narrows to enter copulatory tube. An elongated prostatic reservoir, multiple prostatic gland ducts deplete into the anterior part of prostatic reservoir; prostatic duct leaves prostatic reservoir to enter initial part of copulatory tube. Copulatory organ consists of copulatory tube, no accessory piece, length 88 µm (76–101) with

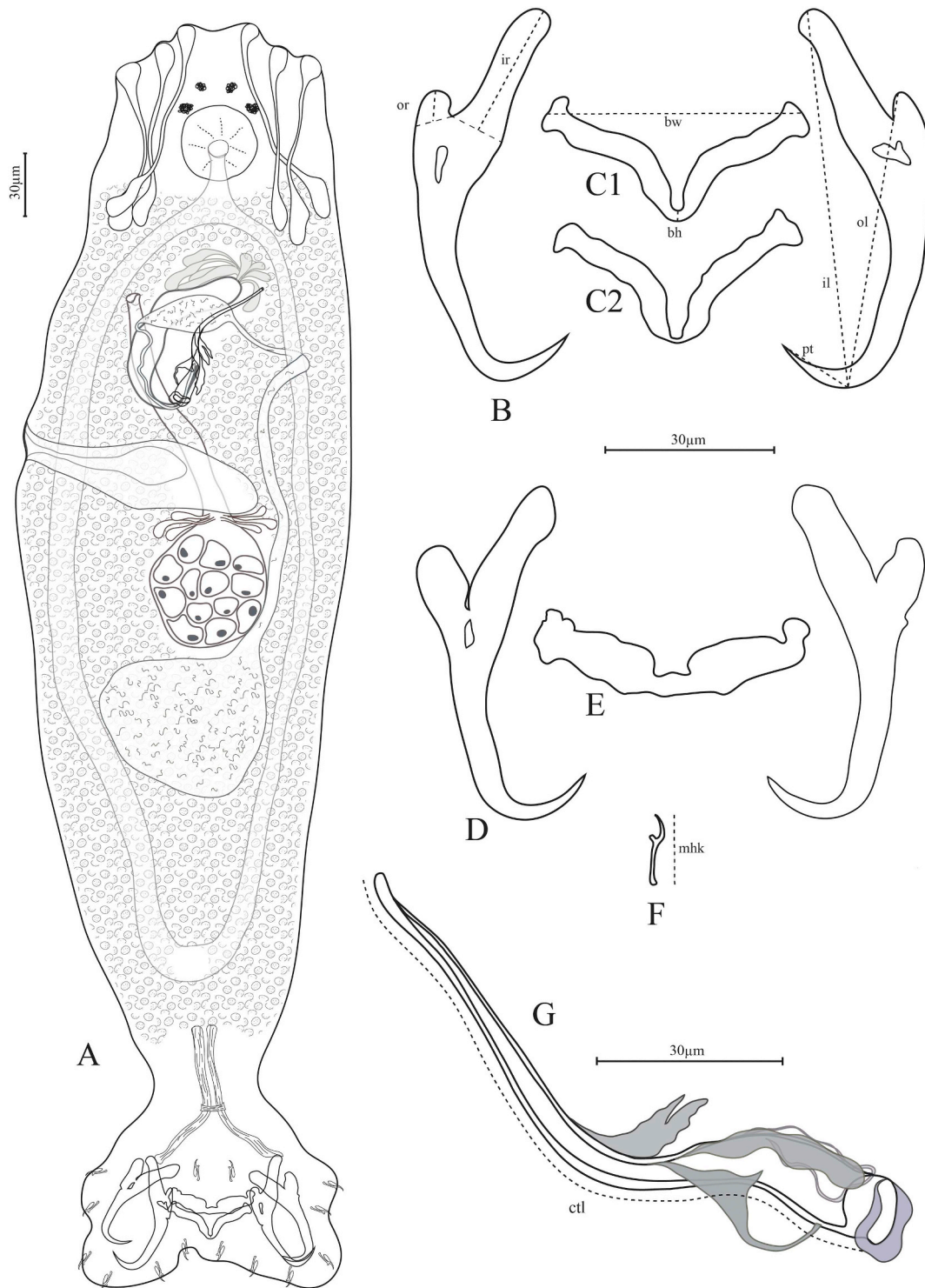


Fig. 1. *Haliotrema susanae* sp. nov. A, Whole worm (ventral view); B, Ventral anchors; C, Ventral bars (two different versions); D, Dorsal anchors; E, Dorsal bar; F, Marginal hook; G, Male copulatory organ. Abbreviations: or, outer root; ir, inner root; ol, outer length; il, inner length; pt., point; bw, bar width; bh, bar height; mhk, marginal hook length; ctl, copulatory tube length.

a short base and tapering tube, lightly sclerotized bract-like extensions at initial (proximal) region of copulatory tube. Ovary rounded, in mid body; oviduct arises from anterior of ovary, proceeds anteriorly to form ootype, receiving ducts from surrounding Mehlis' gland (not discernable in all specimens), ascends as uterus to open near male copulatory organ; vaginal duct not observed. Dextral vaginal opening with prominent large elongate gourd-shaped vaginal chamber, thick-walled, lightly sclerotized with distal end traversing into midportion of body. Vitelline

system confluent anterior to haptor.

### 3.2. Taxonomic summary

*Type-host*: *Myripristis murdjan* Forsskål, 1775 (Beryciformes: Holocentridae: Myripristinae).

*Site of infection*: Gill filaments.

*Type-locality*: Langkawi Island, Malaysia.

**Table 2**  
Species used for molecular analysis in this study with their host species, locality and GenBank accession numbers.

Monogenean species	Host species	Locality	GenBank number	GenBank authority
<i>Dactylogyridea</i> Bychowsky 1937				
<i>Ancyrocephalidae</i> Bychowsky, 1937				
<i>Haliotrema susanae</i> sp. nov.	<i>Myripristis murdjan</i>	Langkawi Island, Kedah, Malaysia	MG518632	Present study
<i>Euryhaliotrema lisae</i> Kritsky & Diggles, 2014	<i>Lufjanus johnii</i>	Pulau Ketam, Selangor, Malaysia	MG593832	Present study
<i>Euryhaliotrema longibaculoides</i> Kritsky & Diggles, 2014	<i>Lufjanus johnii</i>	Pulau Ketam, Selangor, Malaysia	MG593833	Present study
<i>Haliotrema bilobatus</i> (Yamaguti, 1953) Bychowsky & Nagibina, 1970	<i>Drepane punctata</i>	Off Johor, Malaysia	MG593837	Present study
<i>Haliotrema magnihamus</i> Bychowsky & Nagibina, 1970	<i>Drepane punctata</i>	Off Johor, Malaysia	MG593838	Present study
<i>Haliotrema neoblobatus</i> Bychowsky & Nagibina, 1970	<i>Drepane punctata</i>	Carey Island, Selangor, Malaysia	MG593835	Present study
<i>Haliotrema parvihamus</i> Bychowsky & Nagibina, 1970	<i>Drepane punctata</i>	Carey Island, Selangor, Malaysia	MG593834	Present study
<i>Haliotrema spinnicirrus</i> (Yamaguti, 1953) Bychowsky & Nagibina, 1970	<i>Drepane punctata</i>	Carey Island, Selangor, Malaysia	MG593836	Present study
<i>Euryhaliotrema johnii</i> (Tripathi, 1959) Kritsky & Boeger, 2002	<i>Lufjanus rhodopterus</i> (this host name is stated in the publication but is invalid, most likely <i>Lethrinus rhodopterus</i> )	Yangjiang, China	DQ157657.1	Wu et al., 2006
<i>Euryhaliotrema perezponcei</i> Garcia-Vargas, Fajer-Avila & Lamothe-Argumedo, 2008	Not available	Not available	HQ615996.1	Soler-Jimenez et al., unpublished
<i>Euryhaliotrema annulocirrus</i> (Yamaguti, 1968) Kritsky, 2012 [previously known as <i>Euryhaliotrematoides annulocirrus</i> ]	Not available	Not available	EU836195.1	Sun & Yang, unpublished
<i>Euryhaliotrematoides aspis</i> (Plaisance & Kritsky, 2004) Kritsky, 2012 [previously known as <i>Euryhaliotrematoides aspis</i> ]	<i>Chaetodon vagabundus</i>	Australia	AY820614.1	Plaisance et al., 2005
<i>Euryhaliotrematoides berenguelae</i> (Plaisance & Kritsky, 2004) Kritsky, 2012 [previously known as <i>Euryhaliotrematoides berenguelae</i> ]	<i>Chaetodon citrinellus</i>	French Polynesia	AY820615.1	Plaisance et al., 2005
<i>Euryhaliotrematoides grandis</i> (Mizelle & Kritsky, 1969) Kritsky, 2012 [previously known as <i>Euryhaliotrematoides grandis</i> ]	<i>Chaetodon vagabundus</i>	Palau	AY820616.1	Plaisance et al., 2005
<i>Euryhaliotrematoides microphallus</i> (Yamaguti, 1968) Kritsky, 2012 [previously known as <i>Euryhaliotrematoides microphallus</i> ]	<i>Heniochus chrysostrabus</i>	Palau	AY820617.1	Plaisance et al., 2005
<i>Euryhaliotrematoides pirulum</i> (Plaisance & Kritsky, 2004) Kritsky, 2012 [previously known as <i>Euryhaliotrematoides pirulum</i> ]	<i>Chaetodon lunula</i>	French Polynesia	AY820618.1	Plaisance et al., 2005
<i>Euryhaliotrematoides triangulovagina</i> (Yamaguti, 1968) Kritsky, 2012 [previously known as <i>Euryhaliotrematoides triangulovagina</i> ]	<i>Chaetodon kleinii</i>	Palau	AY820619.1	Plaisance et al., 2005
<i>Haliotrema angelepterus</i> Plaisance, Bouamer & Morand, 2004	<i>Chaetodon kleinii</i>	Palau	AY820620.1	Plaisance et al., 2005
<i>Euryhaliotrematoides anguiformis</i> (Zhang in Zhang, Yang & Liu, 2001) Kritsky, 2012 [previously known as <i>Haliotrema anguiforme</i> ]	<i>Lufjanus monostigma</i>	Guangdong, China	DQ537375.1	Wu et al., 2007
<i>Haliotrema cromilepis</i> Young, 1968	<i>Upeneus quadrilimeatus</i>	Guangdong, China	DQ537378.1	Wu et al., 2007
<i>Haliotrema ctenochaeti</i> Yamaguti, 1968 (nec Young, 1968)	<i>Epinephelus coioides</i>	Nha Trang Bay, Vietnam	EU523146.1	Dang et al., 2010
<i>Haliotrema digroides</i> Zhang in Zhang, Yang & Liu, 2001	Not available	Not available	EU836199.1	Sun & Yang, unpublished
<i>Haliotrema dongshaense</i> Sun, Gibson & Yang, 2011	Not available	Not available	EU836200.1	Sun & Yang, unpublished
<i>Haliotrema epinepheli</i> Young, 1968	<i>Epinephelus fasciatus</i>	Heron Island, Australia	KJ571014.1	Sun et al., unpublished
<i>Lethrinurema flet</i> (Young, 1968) Lim & Justine, 2011 [previously known as <i>Haliotrema flet</i> ]	<i>Lethrinus nebulosus</i>	Yangjiang, Guangdong Province, China	EU836201.1	Dang et al., 2010
<i>Lethrinurema flet</i> (Li & Chen, 2005) Sun et al. 2014 [previously known as <i>Haliotrema grossecuvitubus</i> ]	Not available	Not available	DQ157661.1	Wu et al., 2006
<i>Haliotrema johnstoni</i> Bychowsky & Nagibina, 1970	<i>Upeneus luzonius</i>	Haikou Hainan Province, China	EU836204.1	Sun & Yang, unpublished
<i>Haliotrema macassarense</i> (Yamaguti, 1963) Bychowsky & Nagibina, 1970 [previously known as <i>Haliotrema macassarensis</i> ]	Not available	Not available	DQ157664.1	Wu et al., 2006
<i>Haliotrema macracantha</i> Yamaguti, 1968	Not available	Not available	EU836207.1	Sun & Yang, unpublished
<i>Euryhaliotrema nanaensis</i> (Li, Yan, Yui, Lan & Huang, 2005) Kritsky, 2012 [previously known as <i>Haliotrema nanaense</i> ]	Not available	Not available	EU836208.1	Sun & Yang, unpublished
	<i>Lufjanus argentimaculatus</i>	Guangdong, China	DQ537373.1	Wu et al., 2007

(continued on next page)

Table 2 (continued)

Monogenean species	Host species	Locality	GenBank number	GenBank authority
<i>Pseudohaliotrema platicephali</i> (Yin & Sproston, 1948) Young, 1968 [previously known as <i>Haliotrema playicephali</i> ]	Not available	Not available	FJ757866.1	Su, unpublished
<i>Haliotrema pratense</i> Sun et al., 2007	Not available	Not available	EU836209.1	Sun & Yang, unpublished
<i>Haliotrema scyphovagina</i> Yamaguti, 1968	<i>Forcipiger flavissimus</i>	French Polynesia	AY820622.1	Plaisance et al., 2005
<i>Haliotrematoides shenzhenensis</i> (Wang, Liu & Zhou, 2003) Kritsky, Yang & Sun, 2009 [previously known as <i>Haliotrema shenzhenensis</i> ]	<i>Lujitanus argentimaculatus</i>	Guangdong, China	DQ537372.1	Wu et al., 2007
<i>Euryhaliotrema spirotaubiforum</i> (Zhang in Zhang, Yang & Liu, 2001) Wu, Zhu, Xie & Li, 2006 [previously known as <i>Haliotrema spirotaubiforum</i> ]	<i>Lujitanus stellatus</i>	Yangjiang, Guangdong Province	DQ157656.1	Wu et al., 2006
<i>Haliotrema subancistroides</i> Zhang in Zhang, Yang & Liu, 2001	<i>Gerres filamentosus</i> , <i>G. abbreviatus</i>	Dayawan, Guangdong Province	DQ157648.1	Wu et al., 2006
<i>Haliotrematoides guttati</i> (Garcia-Vargas, Fajer-Avila & Lamothe-Argumedo, 2008) Kritsky, Yang & Sun, 2009	Not available	Not available	HQ615993.1	Soler-Jiminez et al., unpublished
<i>Haliotrematoides plectridium</i> Kritsky & Mendoza-Franco in Kritsky, Yang & Sun, 2009	Not available	Not available	HQ615994.1	Soler-Jiminez et al., unpublished
<i>Haliotrematoides spinatus</i> Kritsky & Mendoza-Franco in Kritsky, Yang & Sun, 2009	Not available	Not available	HQ615995.1	Soler-Jiminez et al., unpublished
<i>Ligophorus bantingensis</i> Soo & Lim, 2012	<i>Planitiza subviridis</i>	Carey Island, Selangor, Malaysia	KM221909	Soo & Lim, 2012
<i>Ligophorus chelanus</i> Soo & Lim, 2012	<i>Planitiza subviridis</i>	Carey Island, Selangor, Malaysia	KM221912	Soo & Lim, 2012
<i>Ligophorus funnelus</i> Soo & Lim, 2012	<i>Planitiza subviridis</i>	Carey Island, Selangor, Malaysia	KM221914	Soo & Lim, 2012
<i>Ligophorus najiosodhii</i> Soo & Lim, 2012	<i>Planitiza subviridis</i>	Carey Island, Selangor, Malaysia	KM221920	Soo & Lim, 2012
<i>Ligophorus parvicopulatrix</i> Soo & Lim, 2012	<i>Planitiza subviridis</i>	Carey Island, Selangor, Malaysia	KM221921	Soo & Lim, 2012
<i>Metahaliotrema scaophagi</i> Yamaguti, 1953 [previously known as <i>M. geminatahamula</i> Pan et al., 1995]	<i>Scatophagus argus</i>	Panyu, Guangdong Province, China	DQ157646.1	Wu et al., 2006
<i>Metahaliotrema mizellei</i> Venkatanarasiah, 1981	<i>Scatophagus argus</i>	Panyu, Guangdong Province, China	DQ157647.1	Wu et al., 2006
Outgroup				
Tetraonchidae				
<i>Calydiscoides indianus</i>	<i>Nemipterus japonicus</i>		EF100557.1	Wu et al., unpublished

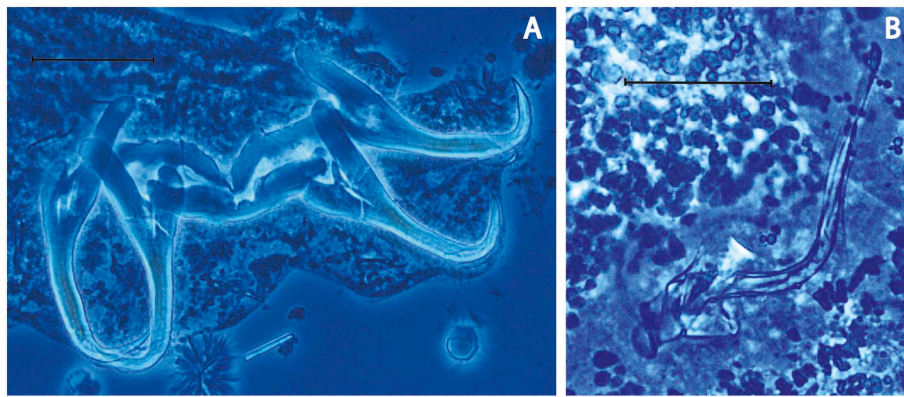


Fig. 2. Photomicrographs of sclerotized hard parts of *Haliotrema susanae* sp. nov., captured using Leica digital camera and an image analysis software (QWin Plus): (A) anchors and bars; (B) male copulatory organ. Scale = 30  $\mu$ m.

**Other host and locality:** *Myripristis hexagona* Lacépède, 1802 from Langkawi Island, Malaysia.

**Type-specimens:** Holotype (ZRC.PLA.0367) and 1 paratype (ZRC.PLA.0368) in Zoological Reference Collection, Lee Kong Chian Natural History Museum, Singapore; 48 paratypes (NHMUK 2018.9.27.1 - 48) in the Natural History Museum London (NHMUK). **Specimens studied:** 50 specimens studied; 46 specimens measured.

**Etymology:** This new species is named in honor of the late Dr. Susan Lim Lee Hong who pioneered taxonomic studies of monogeneans from Malaysian waters.

**Representative DNA sequence:** MG518632 (partial 28S rDNA).

### 3.3. Differential diagnosis

*Haliotrema susanae* sp. nov. differs from previously described *Haliotrema* species from holocentrids in the morphology of the male copulatory organ and ventral bar. The general morphology of the male copulatory organ is similar for all *Haliotrema* species from holocentrids: a long copulatory tube with a broadened distal region and prominent initial. But the new species is different in having a bulbous base with bract-like expansions near the initial of the copulatory tube (Fig. 2B). Upon comparison with previous studies [10,11], it was observed that the copulatory tube of *H. susanae* sp. nov. is long, similar to *H. curvicirrus* Zhukov, 1980. However, the copulatory tube of *H. susanae* sp. nov. has bract-like extensions from mid length to the initial region compared to none in *H. curvicirrus*. The ventral bar of the new species is V-shaped, similar to *H. longirectocirrus* Zhukov, 1980 and *H. myripristisi* Zhukov, 1980 but differs from both mentioned species in being enlarged at the ends (Fig. 2A). The dorsal bars of the new species are similar to *H. curvicirrus* Zhukov, 1980 in being narrow and slightly bent in the center but differs in length; 43  $\mu$ m in *H. susanae* sp. nov. and 33  $\mu$ m in *H. curvicirrus*. Single grooves were observed on the base of the ventral and some dorsal anchors of *H. susanae* sp. nov. The presence of grooves were observed on the dorsal anchors of *H. tenuihamus* Zhukov, 1980 and *H. myripristisi* but this feature is not a distinguishing one for *H. susanae* sp. nov. as not all of its dorsal anchors had grooves. Apart from grooves, the dorsal anchors of *H. susanae* sp. nov. has similar shaped bases, shafts and points with *H. papillibaculum* Zhukov, 1980 but differs from *H. papillibaculum* in the morphology of the copulatory organ. The copulatory tube is bent and shorter (32  $\mu$ m) in *H. papillibaculum* compared to the new species where the copulatory tube is not bent and is longer (88  $\mu$ m).

### 3.4. Molecular results

The partial 28S rDNA sequence data for *Haliotrema susanae* sp. nov. was generated and aligned alongside 47 monogeneans from closely related genera with *Calydiscoides indianus* as the outgroup species. The

aligned dataset consisted of 48 sequences and comprised of 633 positions after trimming the ends to match the shortest aligned sequence. The maximum likelihood (ML) tree constructed consists of five clades (Fig. 3). The species sequences used in this analysis were specifically chosen for their relationship to the genus *Haliotrema* or those reassigned from the genus *Haliotrema* (see Table 2). Clade 1 consists of only *Euryhaliotrema* species (Fig. 3). Clade 2 and 3 consists of single genera: Clade 2 with only *Haliotrematoides* species (*H. spinatus*, *H. plectridium*, *H. guttati* and *H. shenzhenensis*) and Clade 3 with only *Ligophorus* species (*L. parvicopulatrix*, *L. bantingensis*, *L. chelatus*, *L. funnelus* and *L. navjotsodhii*). Clade 4 consists of four species: two *Metahaliotrema* species (*M. geminatohamula* and *M. mizellei*) and two *Haliotrema* species (*H. subancistroides* and *H. digyroides*). Clade 5 consists of the new species, *H. susanae* sp. nov. and other *Haliotrema* species (Fig. 3), reaffirming the designation of the new species as a member of the genus *Haliotrema*.

## 4. Discussion

The genus *Haliotrema* has been termed the ‘waste-basket’ genera because of the assignment of monogeneans with the following general characteristics: four anchors, 14 marginal hooks and two bars [12]. Forty three species, originally described as *Haliotrema* species were reassigned to different or newly-erected genera: eighteen species reassigned as *Euryhaliotrema* Kritsky & Boeger, 2002; fourteen as *Haliotrematoides* Kritsky, Yang & Sun, 2009; five as *Lethrinotrema* Lim & Justine, 2011; two as *Ligophorus* Euzet & Suriano, 1977 and one each for the following genera; *Triacanthinella* Bychowsky & Nagibina, 1968; *Tetrancistrum* Goto & Kikuchi, 1917; *Parancylodiscoides* Caballero & Bravo Hollis, 1961 and *Pseudohaliotrema* Yamaguti, 1953. After these reassignments, there are 191 species listed as *Haliotrema* species on the World Register of Marine Species [6]. These reassignments have been based on host groups and salient morphological characters, as exemplified by the reassignments of the *Haliotrema* species from mugilids to *Ligophorus* [13]; from lutjanids, caesionids, haemulids and sparids to *Haliotrematoides* [14]; from the sciaenids, sparids, haemulids, chaetodontids and lutjanids to *Euryhaliotrema* [15] and from lethinids to *Lethrinotrema* [16].

*Haliotrema susanae* sp. nov. is the most recently described *Haliotrema* species from holocentrids since Zhukov [11]. *Haliotrema susanae* sp. nov. has the ‘classic’ morphological characteristics of the genus *Haliotrema*; four anchors, two bars, fourteen marginal hooks and a vas deferens which loops around the intestinal caeca [17]. The new species also has a large vaginal chamber which traverses across more than half the width of the monogenean’s body and is larger in comparison to the vaginal chamber of other *Haliotrema* species from holocentrids (Fig. 1). The anchors of the new species are morphologically similar to *H. papillibaculum* and *H. tenuihamus* in having long shafts and recurved points but *H. susanae* sp. nov. can be differentiated from both these species in

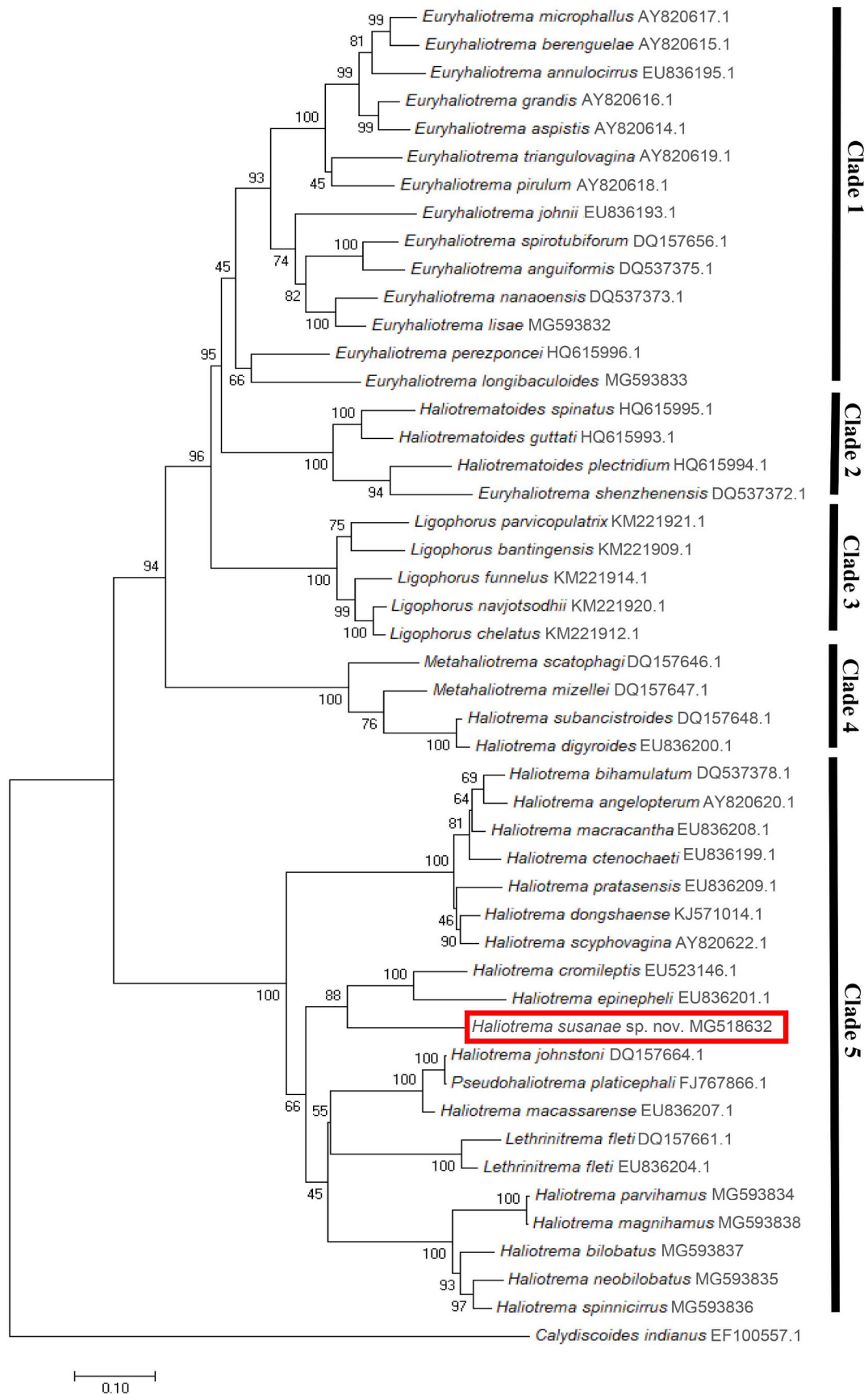


Fig. 3. Maximum likelihood (ML) tree generated from partial 28S rDNA sequence data using MEGA7 with *Calydiscoides indianus* as the outgroup. Bootstrap values are shown along the branches. The new species is indicated in the red box. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

having a morphologically different copulatory organ.

The molecular analysis consists of selected four-anchor monogeneans from previous studies and GenBank (Table 2), which serves to specifically reflect certain findings. A molecular analysis with all available *Haliotrema*-like species would have been more conclusive but this couldn't be attempted because partial 28S rDNA molecular sequence information is not available for all species on the GenBank. The present study showed that the reassignment of certain *Haliotrema*-like species based on host group and morphology is in agreement with the molecular analysis as exemplified by grouping of species in Clades 1 to 4 (Fig. 3). Clade 1 consists of only *Euryhaliotrema* species where all members of the clade have slightly overlapping gonads, a dextral vaginal pore, coiled male copulatory organ and a pretesticular germarium [15]. Clade 2 and 3 consist of single genera: Clade 2 consists only of *Haliotrematoides* species and Clade 3 only of *Ligophorus* species. Members of the genus *Haliotrematoides* were reassigned from *Haliotrema* based on a number of characteristics, some of which were marginal hook distribution, dorsal anchors lacking well-developed deep roots, ventral bar with two submedial pockets along the anterior margin and a copulatory complex lacking an accessory piece [14]. Some of the species reassigned were *Haliotrema longihamus* Zhukov, 1976 to *Haliotrematoides longihamus* (Zhukov, 1976) Kritsky et al., 2012 and *Haliotrema longitubocirrus* to *Haliotrematoides longitubocirrus* (Bychowsky & Nagibina, 1971) Kritsky et al., 2012. Two members of *Ligophorus* were reassigned from *Haliotrema* for having a long tubular copulatory organ with an accessory piece and a sclerotized vagina; *Ligophorus vanbenedenii* (Parona & Perugia, 1890) Euzet & Suriano, 1977 and *Ligophorus mugilinus* (Hargis, 1955) Euzet & Suriano, 1977, but subsequent new species description were also based on the host factor. Clade 4 consists of four species; *Metahaliotrema geminatomamula*, *M. mizellei*, *H. subancistroides* and *H. digyroides*. Reassignments are recommended for species within this clade, especially since all the species can be found in one common fish host group, *Gerres* spp. Clade 5 consists of 19 species, mostly from the genus *Haliotrema*. Three species have been reassigned: *Haliotrema fleti* Young, 1968 as *Lethrinotrema fleti* (Young, 1968) Lim & Justine, 2011 [16], *H. grossecurvitubus* Li & Chen, 2005 as *L. grossecurvitubum* (Li & Chen, 2005) Sun, Li & Yang, 2014 [18] and *H. platycephali* as *Pseudohaliotrema platycephali* (Yin & Sproston, 1948) Young, 1968 [19]. Five *Haliotrema* species (*H. bilobatus*, *H. magnihamus*, *H. parvihamus*, *H. neobilobatus* and *H. spinnicirrus*) collected from *Drepane punctata* off Malaysian waters formed a strongly supported clade within Clade 5. These species were proposed to be reassigned as members of *Parancylodiscoides* Caballero & Bravo-Hollis, 1961 based on possession of a dextral vaginal aperture but due to uncertainty of using the position of the vaginal aperture as a main distinguishing character, reassignments have not been carried out [20].

Young (1968) found that *Haliotrema* species from the families Serranidae and Holocentridae have similar morphological characteristics, resulting in three species from his study (*H. epinepheli*, *H. cromileptis* and *H. holocentri*) grouped as members of Group 5 [21]. The present study supports this grouping, especially in the phylogenetic tree constructed (Fig. 3). *Haliotrema susanae* sp. nov. formed a clade (bootstrap value of 89%) with *H. epinepheli* and *H. cromileptis*, similar to Young's study [21], in which *H. holocentri* was also grouped with *H. epinepheli* and *H. cromileptis* based on only morphological features such as anchors with long roots, male copulatory organs lacking accessory pieces, a group of gland cells anterior to male copulatory organ and a large thick-walled vaginal chamber. There is no molecular sequence data for any monogeneans from the family Holocentridae in the GenBank except for *H. susanae* sp. nov. These results show that characterization of species based on morphology is in synchrony with results of the molecular analysis. Description and identification of new species are more conclusive when morphology and molecular results agree. Integrating different analytical methods during the taxonomic process (also known as integrative taxonomy) is especially useful in describing morphologically similar species. This was exemplified in the description

and identification of *Ligophorus* species from Malaysia [22,23] and the Atlantic-Pacific Oceans [24]. In the current study, the integration of molecular biology into the description of *H. susanae* sp. nov. has not only reinforced its position as a *Haliotrema* species but its results is shown to agree with a morphological study done years ago.

During the course of this study, species synonymy was observed for *Haliotrema curvicirrus* whereby one record is described by Yamaguti [10] and another described by Zhukov [11]. On the WoRMS website [6], the author of *H. curvicirrus* is Yamaguti [10]. The species described by Zhukov [11] has not been reassigned or redescribed to date. Upon investigation of literature from the two authors, both synonyms of *H. curvicirrus* differ in first, having different hosts (Yamaguti's species is from *Parupeneus porphyreus* whilst Zhukov's is from *Holocentrus ascensionis*) and second, in having morphologically different hard parts. No further inferences can be made regarding the true identity of *H. curvicirrus* because both hosts are endemic to waters in the Western Atlantic Ocean and there have been no subsequent studies on both species to date. Other cases of species synonymy has also been reported for *Haliotrema* species from acanthurids; *H. ctenochaeti* Yamaguti, 1968 - *H. ctenochaeti* Young, 1968 and *H. serpenticirrus* Yamaguti, 1968 - *H. dempsteri* (Mizelle & Price, 1964) Young, 1968 [25]. Further investigation and possible reassignments are proposed for these synonymic species.

## Acknowledgement

This paper was supported by the Postgraduate Research Fund (Vote F PS275/2008C) from the University of Malaya to the author. The author would like to thank Yap Fook Choy, Hazreen Abdul Jabar and Masrizal Shauri for assisting in obtaining dead host specimens from Langkawi Island, Malaysia.

## Conflict of interest

The author declares that there is no conflict of interests.

## Ethical approval

None needed as host were obtained dead from the fishermen port.

## References

- [1] R. Froese, D. Pauly (Eds), Fishbase. World Wide Web electronic publication. <http://www.fishbase.org>, 2018 (accessed on 9 January 2018).
- [2] L.H.S. Lim, Three new species of *Bychowskyella* Achmerow, 1952 (Monogenea) from Peninsular Malaysia, *Syst. Parasitol.* 19 (1991) 33–41.
- [3] O.Y.M. Soo, L.H.S. Lim, Eight new species of *Ligophorus* Euzet & Suriano, 1977 (Monogenea: Ancyrocephalidae) from mugilids off Peninsular Malaysia, *Raffles Bull. Zool.* 60 (2012) 241–264.
- [4] H. Hassouna, B. Michot, J.P. Bachelier, The complete nucleotide sequence of mouse 28S rRNA gene. Implications for the process of size increase of the large subunit rRNA in higher eukaryotes, *Nucleic Acids Res.* 12 (8) (1984) 3563–3583.
- [5] L.H. Qu, M. Nicoloso, J.P. Bachelier, Phylogenetic calibration of the 5' terminal domain of large rRNA achieved by determining twenty eukaryotic sequences, *J. Mol. Evol.* 28 (1988) 113–124.
- [6] WoRMS Editorial Board, World Register of Marine Species, Available from <http://www.marinespecies.org> at VLIZ. Accessed 2018-02-05 (2018), <https://doi.org/10.14284/170>.
- [7] K. Katoh, D.M. Standley, MAFFT multiple sequence alignment software version 7: improvements in performance and usability, *Mol. Biol. Evol.* 30 (2013) 772–780.
- [8] T.A. Hall, BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT, *Nucleic Acids Symp. Ser.* 4 (1999) 95–98.
- [9] S. Kumar, G. Stecher, K. Tamura, MEGA7: Molecular Evolutionary Genetics Analysis version 7.0 for bigger datasets, *Mol. Biol. Evol.* 33 (2016) 1870–1874.
- [10] S. Yamaguti, Monogenetic Trematodes of Hawaiian Fishes, University of Hawaii Press, Hawaii, 1968.
- [11] E.V. Zhukov, Monogeneans of the genus *Haliotrema* Johnston & Tiegs, 1922 from the gills of fishes from the families Holocentridae and Acanthuridae of the Gulf of Mexico, *Parazitologiya* 29 (1980) 41–52.
- [12] G.J. Klassen, Phylogeny of *Haliotrema* species from boxfishes: are *Haliotrema* species from boxfishes monophyletic? *J. Parasitol.* 80 (4) (1994) 596–610.
- [13] L. Euzet, D.M. Suriano, *Ligophorus* n. g. (Monogenea: Ancyrocephalidae) parasite des Mugilidae (Téléostéens) en Méditerranée, *Buletin du Muséum National*



- d'Histoire Naturelle, Série 3, Zoologie 472 (1977) 799–821.
- [14] D.C. Kritsky, T.B. Yang, Y. Sun, Dactylogyrids (Monogeneoidea, Polyonchoinea) parasitizing the gills of snappers (Perciformes, Lutjanidae): proposal of *Haliotrematoides* n. gen. and descriptions of new and previously described species from marine fishes of the Red Sea, the eastern and Indo-west Pacific Ocean, Gulf of Mexico and Caribbean Sea, *Zootaxa* (1970) (2012) 1–51.
- [15] D.C. Kritsky, Dactylogyrids (Monogeneoidea: Polyonchoinea) parasitizing the gills of snappers (Perciformes: Lutjanidae): revision of *Euryhaliotrema* with new and previously described species from the Red Sea, Persian Gulf, the eastern and Indo-west Pacific Ocean, and the Gulf of Mexico, *Zoologia* 29 (3) (2012) 227–276.
- [16] L.H.S. Lim, J.-L. Justine, Two new species of ancyrocephalid monogeneans from *Lethrinus rubrioperculatus* Sato (Perciformes: Lethrinidae) off New Caledonia, with the proposal of *Lethrinitrema* n. gen, *Syst. Parasitol.* 78 (2011) 123–138.
- [17] T.H. Johnston, O.W. Tiegs, New Gyrodactyloid trematodes from Australian fishes together with a reclassification of the superfamily Gyrodactyloidea, *Proc. Linn. Soc. NSW* 4 (1922) 83–131.
- [18] Y. Sun, M. Li, T. Yang, Studies on *Lethrinitrema* Lim & Justine, 2011 (Monogenea: Dactylogyridae), with the description of two new species, a key to the genus and a phylogenetic analysis based on rDNA sequences, *Syst. Parasitol.* 88 (2) (2014) 119–139.
- [19] P.C. Young, The taxonomy of some dactylogyrid Monogeneoidea from Australian fishes, *Zool. Anz.* 180 (1968) 269–279.
- [20] D.C. Kritsky, M.D. Bakenhaster, Redescription and New Host Records for *Parancylodiscoides macrobaculum* n. comb. (Monogeneoidea: Dactylogyridae) from Groupers (Serranidae: Epinephelinae) in the Gulf of Mexico, *Comp. Parasitol.* 83 (2) (2016) 260–264.
- [21] P.C. Young, Ten new species of *Haliotrema* (Monogeneoidea: Dactylogyridae) from Australian fish and a revision of the genus, *J. Zool.* 154 (1968) 41–75.
- [22] O.Y.M. Soo, L.H.S. Lim, A description of two new species of *Ligophorus* Euzet & Suriano, 1977 (Monogenea: Ancyrocephalidae) from Malaysian mugilid fish using principal component analysis and numerical taxonomy, *J. Helminthol.* 89 (2015) 131–149.
- [23] O.Y.M. Soo, W.B. Tan, L.H.S. Lim, Three new species of *Ligophorus* Euzet & Suriano, 1977 (Monogenea: Ancyrocephalidae) from *Moolgarda buchanani* (Bleeker) off Johor, Malaysia based on morphological, morphometric and molecular data, *Raffles Bull. Zool.* 63 (2015) 49–65.
- [24] V. Sarabeev, D. Desdevises, Phylogeny of the Atlantic and Pacific species of *Ligophorus* (Monogenea: Dactylogyridae): Morphology vs. molecules, *Parasitol. Int.* 63 (1) (2014) 9–20.
- [25] Y. Sun, C. Yang, T. Yang, Two new species of *Haliotrema* Johnston & Tiegs, 1922 (Monogenea: Dactylogyridae) from *Acanthurus nigrofuscus* (Forsskal) and *A. triostegus* (Linnaeus) (Teleostei: Acanthuridae) in the South China Sea, *Syst. Parasitol.* 91 (3) (2015) 253–259.