A New Species of *Rhabdias* (Nematoda: Rhabditida: Rhabdiasidae) from Miyakojima Island, Okinawa, Japan

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A new nematode species that is lung parasitic to a bufonid toad, *Rhabdias kafunata* sp. nov., is described and illustrated from Miyakojima island, Okinawa Prefecture, Japan. Although the new species was previously identified as *R. incerta* Wilkie, 1930, it is clearly distinguishable from that species by the characteristics of body proportion, buccal capsule, and esophagus. Additionally, the new species differs from other congeners inhabiting East Asia and the Russian Far East in the following features: longer body length, cup-like buccal capsule, longer esophagus length, presence of thickened anterior muscular part in esophagus, cuticular inflation less prominent in region posterior to small cephalic cuticular inflation, middle part of body and tail tip, and tail long and tapering. Partial sequences of mitochondrial cytochrome c oxidase subunit I and 12S rDNA genes were provided as DNA barcodes for the new species.

**Key Words:** *Bufo gargarizans miyakonis*, endoparasite, misidentification, Ryukyu Islands, nematode.

**Introduction**

The genus *Rhabdias* Stiles and Hassall, 1905 is a parasitic nematode group that dwells in the lung of amphibians and reptiles, and consists of ca. 85 nominal species that are distributed worldwide (Kuzmin 2013). Seven species of this genus have been recorded from the Japanese and/or Ryukyu Archipelago, Japan, and two have also been identified from the adjacent continental region (Kung and Wu 1945; Wang et al. 1978; Moravec and Sey 1985; Kuzmin 2013).

*Rhabdias incerta* Wilkie, 1930 is a bufonid toad specific species, and was originally described from Tokyo, Japan (Wilkie 1930). This species has been recorded from the Japanese Archipelago, Miyakojima island (the sole island of the Ryukyu Archipelago where the native bufonid species is distributed) and the Russian Far East (Yamaguti 1941; Hasegawa 1984; Hasegawa and lwatsuki 1993; Goldberg and Bursey 2002; Hasegawa and Asakawa 2004; Kuzmin 2013).

The previous studies reported the morphological differences between a western Japanese Archipelago population and the original material of *R. incerta* (Yamaguti 1941). Although morphological similarities between the Miyakojima island population and Japanese mainland population were indicated by Hasegawa (1984), the measurements of the buccal capsule of the island population, a diagnostic character of *Rhabdias* species (Svitin et al. 2018), were not provided at that time. Furthermore, because a recent phylogenetic study of *Rhabdias* suggested the occurrence of multiple cryptic species-level divergences (Müller et al. 2018), the true species diversity of *Rhabdias* should be revealed by careful morphological observation and a molecular phylogenetic approach.

In the present study, *Rhabdias* specimens collected from the Miyako toad, *Bufo gargarizans miyakonis* Okada, 1931, obtained from Miyakojima island were inspected, and a new species is described. Additionally, mitochondrial DNA sequences of the new species are provided for future phylogenetic analyses.

**Materials and Methods**

**Sampling and morphological examination.** Specimens of *Rhabdias* examined in this study were obtained from six specimens of *B. g. miyakonis* which were collected from Hira-Rashinikakasone (24°48′48.0″N, 125°18′44.0″E), Miyakojima island, Okinawa Prefecture, Japan. All captured toads were euthanized by immediate preservation at −20°C. The body cavity of each anuran was dissected by a longitudinal incision, and then the lungs were removed. The excised organs were dissected longitudinally, and the lumens were investigated. Most of the obtained nematode individuals were fixed with a hot 5% solution of glyceralin in 70% ethyl alcohol, and the reminders were fixed with 3% formalin and then preserved in 70% ethyl alcohol. To clear the nematode specimens, they were placed in 5% solution of glyceralin in 30% ethyl alcohol, then incubated 2–3 days at 60°C to
gradually evaporate the ethyl alcohol. The cleared specimens were observed with a light microscope (OLYMPUS BX53).

The measurements were given for a holotype, followed by the range of paratypes in parentheses except for eggs. The measurements of egg were given as mean value, followed by range of holotype. All measurements were described in micrometers (µm) unless otherwise stated in mm.

Both the nematode and toad specimens examined in this study have been deposited in the Zoological Collection of Kyoto University (KUZ).

**PCR and DNA sequencing.** Genomic DNA of nematode specimens was extracted following the method described by Sata (2018). The partial sequence of cytochrome c oxidase subunits I (COI) and 12S ribosomal DNA (12S) were amplified by polymerase chain reaction (PCR) using a TaKaRa Ex Taq kit (Takara Bio, Japan) and a GeneAmp PCR Systems 2700 instrument (Applied Biosystems, USA; ABI). The primer sets used for PCR of COI was primer cocktail of NemFi-t1–NemF3-t1 and NemR1-t1–NemR3-t1, as described in Prosser et al. (2013), of 12S was 12SF and 12SR, as described in Casiraghi et al. (2004). PCR conditions were as follows: 94°C for 3 min, followed by five cycles of 40 s at 94°C, 40 s at 51°C, and 1 min at 72°C; 35 cycles of 40 s at 94°C, 40 s at 51°C, and 1 min at 72°C and then a final extension at 72°C for 5 min for COI; 94°C for 3 min, followed by 35 cycles of 45 s at 94°C, 45 s at 48°C, and 1 min at 72°C and then a final extension at 72°C for 5 min for 12S. The PCR products were purified with ExoSAP-IT reagent (Affymetrix, USA). Sequencing reactions were performed using a BigDye Terminator Cycle Sequencing Kit v. 3.1 (ABI). The primer set used for cycle sequencing (CS) reaction of COI was M13F and M13R, as described in Messing (1983); and the primer set for CS reaction of 12S corresponded to that used for PCR reaction. The products were cleaned by ethanol precipitation and sequenced with an ABI 3130xl Genetic Analyzer. In total, four DNA sequences from the present specimens of *Rhabdias* were determined, and deposited with the International Nucleotide Sequence Database Collaboration (INSDC) through the DNA Data Bank of Japan.

**Results**

**Genus *Rhabdias* Stiles and Hassall, 1905**

*Rhabdias kafunata* sp. nov. (Fig. 1)


**Diagnosis.** The East Asian *Rhabdias*, with long and slender body. Buccal capsule small and cup-like. Esophagus 694–878 long, anterior muscular region thickened. Cicutular inflation less prominent in region posterior to small cephalic cuticular inflation, middle part of body and tail tip. Tail long and tapering sensu Wilkie (1930) and Yuen (1965).

**Material examined.** *Holotype*: KUZ Z2572, whole specimen, adult female, obtained from the lung of a specimen of *Bufo gargarizans miyakonis* (KUZ R77141), collected from Hirara-Nishinakasone (24°48′48.0″N, 125°18′44.0″E), Miyakojima island, Okinawa Prefecture, Japan on 14 November 2016. *Paratypes*: KUZ Z2573–Z2579, whole specimens obtained from the lung of the same specimen of holotype’s host and other *B. g. miyakonis* specimens (KUZ R77142, R77143, R77147, R77149), collected from the same locality and data as the holotype’s host-specimen; KUZ Z2258, one prepared slide of a section of anterior end of a female; and KUZ Z2580, remaining body specimen of KUZ Z2258. KUZ Z2258 and Z2258 were obtained from the lung of a *B. g. miyakonis* specimen (KUZ R77148), collected from the same locality and data as the holotype’s host-specimen.

**Additional materials:** in total, 164 adult and young females were obtained from the lung of *B. g. miyakonis*; KUZ Z2702 and Z2703, 57 whole specimens, and six non-deposited specimens for the present and future molecular studies, obtained from KUZ R77149; KUZ Z2704 and Z2705, eight whole specimens, and one non-deposited specimen for a future molecular study, from KUZ R77147; KUZ Z2706, five whole specimens, and three non-deposited specimens for a future molecular study, from KUZ R77143; KUZ Z2707 and Z2708, 33 whole specimens from KUZ R77142; KUZ Z2709 and Z2710, 44 whole specimens, and five non-deposited specimens for the present and future molecular studies, from KUZ R77141; and KUZ Z2711, two whole specimens, from KUZ R77148.

**Type locality.** Hirara-Nishinakasone (24°48′48.0″N, 125°18′44.0″E), Miyakojima island, Okinawa Prefecture, Japan.

**Type host.** *Bufo gargarizans miyakonis* Okada, 1931 (Amphibia: Bufonidae); site of infection: lung. A total of 173 individuals were obtained from six specimens of *B. g. miyakonis*; mean intensity (mean ± standard error, followed by ranges): 28.8 ± 10.5, 3–66.

**Description.** General. Body long and slender with tapered extremities, body becoming wider at region between anterior end of genital system and vulva. Cephalic end with six lips, each lip with a minute papilla. Amphids situated on anterior end of genital system and vulva. Cephalic end with six lips, each lip with a minute papilla. Amphids situated on anterior end of genital system and vulva. Cephalic end with six lips, each lip with a minute papilla. Amphids situated on anterior end of genital system and vulva. Cephalic end with six lips, each lip with a minute papilla. Amphids situated on anterior end of genital system and vulva.
A new *Rhabdias* from Miyakojima island

208 (166–208) [27.4% (20.8–27.4%) of esophagus length] and 247 (195–259) [32.6% (26.2–32.6%) of body length], respectively, from anterior end of esophagus. Intestine thick walled. Rectum short, with thin wall. Vulva 6.18 (5.97–8.32) mm from cephalic end, and located at middle to posterior of body [55.9% (49.6–55.9%) of body length], having slightly salient lips. Genital system amphidelphic, anterior limb turned posteriorly at 2.18 (2.18–3.28) mm from cephalic end, and posterior limb turned anteriorly at 2.48 (1.45–3.25) mm from tail end. Uteri long, tubular, filled with numerous eggs. Tail long, gradually tapering from anus, and 477 (376–493) long [4.3% (2.8–4.3%) of body length]. Most specimens possessing few numbers of lateral pores. Phasmids located both lateroventral sides, 166 (144–200) from tail end (n=4).

**Etymology.** The specific name *kafunata* is a noun from the word locally used in the Miyako Islands for *B. g. miyanokonis*, the host toad of the new species (Okinawa Center of Language Study 1999–2003); thus not a Latin or Latinized
DNA Sequences. In total, 4 sequences were determined: one specimen obtained from a same host toad as holotype (KUZ R77141), 2 sequences—COI (LC496790; 655 bp), and 12S (LC496792; 475 bp); one specimen obtained from a same host toad as paratypes (KUZ R77148), 2 sequences—COI (LC496791; 655 bp), and 12S (LC496793; 475 bp).

Remarks. \textit{Rhabdias kafunata} sp. nov. can be discriminated from the other nine congeners recorded from East Asia and the Russian Far East as follows. The new species, of which body length ranges from 10.99 to 16.77 mm, can be distinguished from the following four \textit{Rhabdias} species: \textit{R. globocephala} Kung and Wu, 1945 [3.18–6.88 mm in Kuzmin (2005)], \textit{R. montana} Yamaguti, 1954 (6.0–7.8 mm; Yamaguti 1954), \textit{R. nipponica} Yamaguti, 1935 (4.18–5.04 mm; Yamaguti 1935), and \textit{R. polypedatis} Yamaguti, 1941 (4.2–5.2 mm; Yamaguti 1941). \textit{Rhabdias kafunata} sp. nov. can be discriminated from \textit{R. bermani} Rausch, Rausch, and Atrashkevich, 1984 and \textit{R. rhacophori} Yamaguti, 1941 by their esophagus lengths, because latter two species possess much shorter esophagus than that of the new species (Yamaguti 1941; Kuzmin 2013). Accordingly, the taxonomic status of western Japanese-mainland populations of \textit{R. incerta} that were reported by Yamaguti (1941) may reflect genetic divergences between eastern and western Japanese-mainland populations of \textit{R. incerta} in having a shallower buccal capsule. However, the measurements provided in the previous studies cannot discriminate the Russian Far East populations from the western Japanese-mainland populations. The Russian Far East \textit{R. incerta} in Kuzmin (2013) differs from the Wilkie's \textit{R. incerta} in having a shallower buccal capsule. Unlike the new species, \textit{R. japalurae} and \textit{R. tokoyensis} have short and conical tails. \textit{Rhabdias kafunata} sp. nov. also differs from \textit{R. japalurae} in the absence of a cuticular inflation at the region middle of esophagus. The new species possesses a cup-like buccal capsule; this characteristic can discriminate \textit{R. kafunata} sp. nov. from \textit{R. bicornis} Lu, 1934, which has a funnel-like buccal capsule (Lu 1934).

Discussion

As previously identified as \textit{Rhabdias incerta} (Hasegawa 1984; Kuzmin 2013), \textit{R. kafunata} sp. nov. morphologically resembles the \textit{Rhabdias} species described by Wilkie (1930). However, the new species can be clearly discriminated from \textit{R. incerta sensu} Wilkie (1930) by its comparatively slender body, smaller buccal capsule, and shorter esophagus. The new species is also distinguishable from the western Japanese-mainland population of \textit{R. incerta} defined by Yamaguti (1941) by having comparatively slender body, and shorter esophagus. Furthermore, \textit{R. kafunata} sp. nov. differs from the Russian Far East \textit{R. incerta} (Kuzmin 2013) by its narrower width of buccal capsule and shorter esophagus. Although some key characters were not provided, the most numerical characteristics of the Miyakojima island population of “\textit{R. incerta}” reported by the previous studies (Hasegawa 1984; Kuzmin 2013) are concordant with those of the new species. Thus, the nematodes of \textit{Rhabdias} from Miyakojima island in Hasegawa (1984) and Kuzmin (2013) might belong to \textit{R. kafunata} sp. nov. described in the present study.

Considering the high host specificity of \textit{R. incerta} to native bufonid toads in Japan (Wilkie 1930; Yamaguti 1941; Goldberg and Bursey 2002; Hasegawa and Asakawa 2004), their inter-population gene flow may largely depend on the dispersal range of its bufonid hosts. The deep genetic divergence between two subspecies of \textit{Bufo japonicus} Temminck and Schlegel, 1838, which are parapatrically distributed in Honshu, Japan, was revealed by phylogenetic studies (Kawamura et al. 1990; Igawa et al. 2006). Therefore, morphological differences between eastern and western Japanese-mainland populations of \textit{R. incerta} may reflect genetic divergences between those two populations. Because Russian Far East nematodes identified as \textit{R. incerta} were found from the bufonid \textit{Pseudepidalea raddei} (Strauch, 1876) (Kuzmin 2013), it is also highly possible that the Russian population might have genetically diverged from the Japanese mainland populations. The Russian Far East \textit{R. incerta} in Kuzmin (2013) differs from the Wilkie's \textit{R. incerta} in having a shallower buccal capsule. However, the measurements provided in the previous studies cannot discriminate the Russian Far East population from the western Japanese-mainland population (Yamaguti 1941; Kuzmin 2013). Accordingly, the taxonomic status of western Japanese-mainland and Russian Far East populations, which have traditionally been deemed \textit{R. incerta}, should be clarified by further morphological investigation and molecular phylogenetic analysis.

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References


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