Sphaerularia vespae sp. nov. (Nematoda, Tylenchomorpha, Sphaerularioidea), an Endoparasite of a Common Japanese Hornet, Vespa simillima Smith (Insecta, Hymenoptera, Vespidae)

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Kyoto University
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*Sphaerularia vespae* sp. nov., an endoparasite of a common Japanese hornet, *Vespa simillima* is described from Hokkaido, Japan, and its molecular sequence profiles are given. This newly discovered nematode appears to belong to the genus *Sphaerularia*, judging from its characteristic parasitic form, the uterium, which looks like a sausage with many verrucae on its surface. *Sphaerularia vespae* sp. nov. is distinguished from *S. bombi*, the only other nominal species of the genus, by the morphology of the male bursa, female tail, and anterior end of parasitic juveniles. SSU, ITS, and D2D3 (LSU) DNA sequences were determined for *S. vespae* sp. nov. and compared with those from related nematodes obtained from the GenBank database. The sequences from *S. vespae* sp. nov. were close to those of *S. bombi* and several another tylenchid entomoparasitic nematodes. *Sphaerularia vespae* sp. nov. parasitizes overwintering gynes of the hornet and practically sterilizes them, as *S. bombi* does for bumblebee gynes.

**Key words:** hornet, Japan, new species, *Sphaerularia*, taxonomy

**INTRODUCTION**

The genus *Sphaerularia* was established by Dufour (1837), as an extraordinary and unusual parasite of bumblebees, and is currently a monospecific genus comprised of only *S. bombi* Dufour, 1837, after generic revisions (Siddiqi, 2000). However, 270 years after the first identification of sphaerularid nematodes (Réaumur, 1742), we have discovered another *Sphaerularia* that is an endoparasite of the overwintered gynes (potential queens) of the hornet, *Vespa simillima* Smith (Sayama et al., 2007). The new species clearly belongs to the genus *Sphaerularia*, based on its characteristic morphology of the “uterium” female, the parasitic form, but is distinguished from *S. bombi* by several morphological features and its molecular profiles using SSU, ITS, and D2/D3 (LSU) rRNA gene sequences. The nematode is described and figured herein as *Sphaerularia vespae* sp. nov. Preliminary observations on its biological features are also given.

**MATERIALS AND METHODS**

**Collection of hornets**

Overwintered gynes of *V. simillima* were caught using transparent plastic traps baited with clear liquor (distilled spirit) and orange juice from May to August 2005 at Hitsujigaoka, Sapporo, Hokkaido, Japan (E 141°23’, N 42°59’). Most trapped hornets were fixed in Carmon’s solution (ethanol : chloroform : glacial acetic acid=6 : 3 : 1) and preserved in 80% ethanol, and were later dissected for examination of parasitism, though some individuals found alive in the traps were immediately dissected to obtain living nematodes.

**Rearing of the nematodes and morphological observation**

Parasitic juveniles and uteria of the nematode were isolated from the haemocoel of a gyne of *V. simillima* caught on 21 July 2005. A water suspension containing the juvenile nematodes was pipetted onto autoclaved sawdust in a Petri dish (diameter 6.5 cm, height 9.0 cm). Infective adults of the nematode obtained from a 6-month-old culture and parasitic juveniles isolated from a gyne hor-
net caught on 20 July 2005 were heat-relaxed at 65°C, fixed in TAF (triethanolamine-formalin), and processed through a glycerol-ethanol series by using Seinhorst’s method (see Hooper, 1986). The glycerol specimens of the parasitic juveniles and infective adults were mounted in glycerin following the method of Maeseneer and d’Herde (see Hooper, 1986), and the uteria were stored in a small glass vial.

Molecular profiles

The molecular profiles, i.e., nearly full-length SSU, ITS (ITS 1, 5.8S, and ITS 2) and D2D3 (expansion segment of the LSU) rRNA gene sequences, of the new species were determined from a single free-living female, obtained from the sawdust medium. We prepared a DNA sample following the methods described in Ye et al. (2007), and DNA fragments were amplified with the primer sets SSU_F4 (5’-GCT TGT CTC AAA GAT TAA GCC -3’) and SSU_R81 (5’-TGA TCC WKC YGC AGG TTC AC -3’) (SSU: provided at Dr. M. Blaxter’s website, http://www.nematodes.org/barcoding/sourhope/nemoprimers.html), 18SF (5’-CGT AAC AAG GTA GCT GTA G -3’) and 28SR (5’-TTT CAC TCG CCG TTA CTA AGG -3’) (ITS: Ferris et al., 1993; Vrain, 1993), and D2a (5’-ACA AGT ACC GTC AGG GAA AGT TG -3’) and D3b (5’-TCA GGA GGA ACC AGC TAC TA -3’) (D2D3: Nunn, 1992). The amplified fragments were sequenced by PCR direct sequencing by using a Big Dye

Table 1. Morphometric values of Sphaerularia vespae sp. nov. All measurements are in micrometers. The abbreviations are defined as: V=vulval position relative to body length (%); T=testis length relative to body length (%).

<table>
<thead>
<tr>
<th></th>
<th>Holotype (female)</th>
<th>Patatypes (females)a</th>
<th>Patatypes (males)b</th>
<th>Parasitic juvenilesa</th>
</tr>
</thead>
<tbody>
<tr>
<td>n (number of specimens)</td>
<td>–</td>
<td>12</td>
<td>2</td>
<td>10</td>
</tr>
<tr>
<td>L (body length)</td>
<td>1338</td>
<td>1392 ± 61 (1300–1500)</td>
<td>1162, 905</td>
<td>959 ± 54 (871–1024)</td>
</tr>
<tr>
<td>a (body length / maximum body width)</td>
<td>55.6</td>
<td>52.7 ± 6.9 (39.0–59.6)</td>
<td>43.3, 39.1</td>
<td>26.8 ± 3.4 (21.7–32.2)</td>
</tr>
<tr>
<td>c (body length / tail length)</td>
<td>8.2</td>
<td>8.2 ± 0.4 (7.8–8.8)</td>
<td>21.5, 18.8</td>
<td>14.8 ± 0.7 (13.9–15.5)</td>
</tr>
<tr>
<td>c’ (tail length / anal body width)</td>
<td>5.0</td>
<td>5.2 ± 0.5 (4.2–5.7)</td>
<td>3.3, 3.2</td>
<td>2.7 ± 0.3 (2.3–3.1)</td>
</tr>
<tr>
<td>V or T</td>
<td>84.2</td>
<td>83.9 ± 0.7 (82.4–85.0)</td>
<td>68.0, 57.2</td>
<td>–</td>
</tr>
<tr>
<td>M (conus length relative to stylet length (%))</td>
<td>25.6</td>
<td>27.9 ± 1.6 (25.6–30.6)</td>
<td>30.0, 26.7</td>
<td>–</td>
</tr>
<tr>
<td>Lip width</td>
<td>10.5</td>
<td>10.5 ± 0.4 (9.5–11.0)</td>
<td>9.0, 8.0</td>
<td>–</td>
</tr>
<tr>
<td>Lip height</td>
<td>4.5</td>
<td>4.5 ± 0.4 (4.0–5.5)</td>
<td>4.0, 3.0</td>
<td>–</td>
</tr>
<tr>
<td>Stylet cone</td>
<td>4.5</td>
<td>4.5 ± 0.2 (4.0–5.0)</td>
<td>4.0, 4.0</td>
<td>–</td>
</tr>
<tr>
<td>Stylet length</td>
<td>18.0</td>
<td>16.5 ± 0.8 (15.5–18.0)</td>
<td>14.0, 14.0</td>
<td>6.5 ± 0.6 (5.0–7.0)</td>
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<tr>
<td>Nerve ring</td>
<td>100</td>
<td>118 ± 7.7 (100–126)</td>
<td>108, 94</td>
<td>101 ± 3.8 (98–108)</td>
</tr>
<tr>
<td>Excretory pore</td>
<td>135</td>
<td>147 ± 7.4 (135–158)</td>
<td>122, 107</td>
<td>114 ± 4.2 (109–121)</td>
</tr>
<tr>
<td>Maximum body width</td>
<td>24.0</td>
<td>26.5 ± 3.8 (24.0–34.5)</td>
<td>27.0, 23.0</td>
<td>36.0 ± 3.4 (30.5–40.0)</td>
</tr>
<tr>
<td>Ovary+oviduct or testis length</td>
<td>158</td>
<td>165 ± 20 (129–188)</td>
<td>790, 517</td>
<td>–</td>
</tr>
<tr>
<td>Spermatheca</td>
<td>198</td>
<td>412 ± 141 (198–691)</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>Uterus+ quadricolumella</td>
<td>115</td>
<td>94 ± 24 (38–122)</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>Post uterine sac</td>
<td>11.5</td>
<td>14 ± 5.9 (7–28)</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>Testis reflection</td>
<td>–</td>
<td>0, 41.0</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>Vulva-anus length</td>
<td>136.0</td>
<td>147 ± 17 (121–175)</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>Anal body width</td>
<td>15.0</td>
<td>15.0 ± 1.8 (13.0–18.5)</td>
<td>16.5, 15.0</td>
<td>23.5 ± 2.9 (21.0–28.5)</td>
</tr>
<tr>
<td>Tail length</td>
<td>76</td>
<td>79 ± 4.8 (74–89)</td>
<td>54, 48</td>
<td>64 ± 2.9 (58–67)</td>
</tr>
<tr>
<td>Spicule length</td>
<td>–</td>
<td>–</td>
<td>24.0 ± 2, 27.0</td>
<td>–</td>
</tr>
<tr>
<td>Gubernaculum length</td>
<td>–</td>
<td>–</td>
<td>6.0 ± 8.5</td>
<td>–</td>
</tr>
</tbody>
</table>

aAverage ± SD (range).
bFirst and second individual, respectively.
cSubventral view.
The SSU and D2D3 sequences from a uterium were also determined and compared with those obtained from a single female to confirm species identification. A DNA sample was prepared from an ethanol-fixed uterium using a DNeasy Mini Kit® (Qiagen Japan, Tokyo) following the manufacturer’s instructions. PCR amplification was conducted as above. To separate the nematode’s amplicons from the host’s amplicons, the PCR fragments were cloned with a pCR8/GW/TOPO TA cloning Kit® (Invitrogen, Tokyo), and sequenced using M13 sequencing primers 5‘- GTA AAA CGA CGG CCA G -3’ and 5‘- GTA AAA CGG CGG CCA G -3’, following the manufacturer’s manual.

**TAXONOMY**

*Sphaerularia vespae* sp. nov.  
(Figs. 1–6)

**Measurements.** See Table 1.

**Description.** Female (Figs. 1, 3). Body slender, annulated, relaxed or slightly ventrally bent when killed by heat. Lateral field with four conspicuous incisions, originating in the region of procorpus extending to close to the tail tip. Lip fused, smooth, slightly constricted, difficult to distinguish. Stylet conspicuous, conus and shaft separated. Stylet conus short, stout, triangular. Stylet shaft long, slightly curved, basal thickening absent. Procorpus tube-like, about two body diameters long. Two esophageal glands observed, but their openings very obscure. Esophageal glands approx. 4–6 body diameters long, constricted by a nerve ring at about anterior 1/4 position, posterior part overlapping intestine. Esophageal-intestinal junction very vague, not observed. Nerve ring conspicuous, about 120 μm behind anterior end. Rectum and anus may be nonfunctional. Tail short, conical, possessing one to three constrictions. Tail tip has a rounded, finger-like shape.

_Uterium (mature parasitic female) _(Figs. 5, 6). Found in haemocoel of the host insect. Body stout, smoothly curved ventrally. Cuticle soft, thick, possessing annulation on its surface except for anterior and posterior end. Lateral field probably present, but not conspicuous. Lip fused, square in lateral view. Stylet short, fine, without obvious conus/shaft separation. Procorpus tube-like. Two esophageal glands observed, but openings very obscure. Esophageal glands approx. 4 body diameters long, constricted by nerve ring at about anterior 1/3 position, posterior part overlapping intestine. Esophageal-intestinal junction not observed. Nerve ring conspicuous, about 101 μm from anterior end. Excretory pore very obvious, about one body diameter behind nerve ring, Hemizonid just anterior to excretory pore. Rectum present but very obscure, approx. 1/2 body diameter long, not muscular. Anus present, not protuberant, sometimes very difficult to observe. Rectum and anus may be nonfunctional.

**Diagnosis and relationships.** *Sphaerularia vespae* sp. nov. is characterized by the rounded and finger-like tail tip of females and the conspicuous male bursa. Only one species, *S. bombi* Dufour was previously known in the genus *Sphaerularia*. The new species is distinguished from *S. bombi* by the length and shape of the female tail and male bursa, i.e., *S. bombi* has a shorter (47–68 μm) tail with a bluntly pointed tail tip in females and an inconspicuous male bursa, while *S. vespae* sp. nov. has a longer (74–89 μm) tail, with a finger-like and mostly rounded tail tip in females and a large, conspicuous male bursa. The head shape of parasitic juveniles also distinguishes the new species from *S. bombi*. The anterior end of *S. vespae* sp. nov. is square in lateral view, while that of *S. bombi* is rounded and dome-like.

**Type host and locality.** The culture from which the type specimens were obtained was made from parasitic juveniles of *Sphaerularia vespae* sp. nov. isolated from the body cavity of an overwintered gyne of *Vespa simillima* Smith, caught with a bait trap at Hitsujigaoka, Sapporo, Hokkaido, Japan (42°59′N, 141°23′E), 21 July 2005.

**Type material.** Type specimens were obtained from a 6-month-old culture on wet sawdust media on 10 February 2006. Holotype (female), slide number Sphaerularia vespae F-01 (USDANC Number: T-613t). Paratypes, slide numbers Sphaerularia vespae F-02-05 (USDANC Number: T-5635p-T-5638p) (four females); M-01 (USDANC Number: T-5639p) (one male) and *Sphaerularia vespae* P-01-04 (USDANC Number: T-5640p-T-5643p) (four parasitic juveniles) deposited in the USDA Nematode Collection, Beltsville, MD, USA;
A New Endoparasite of a Common Hornet

Fig. 1. *Sphaerularia vespae* sp. nov. (A) Lateral view of female. (B) Lateral view of male. (C) Lateral view of female anterior region. (D) Body surface of female excretory pore region. (E) Lateral field at mid-body. (F) Female reproductive organ (vulva to spermatheca). (G) Female reproductive organ (spermatheca to outstretched ovary). (H) Female reproductive organ (spermatheca to out-tangled ovary). (I–K) Lateral view of female tail. (L) Subventral view of male tail (internal structure). (M) Subventral view of male tail (bursa and lateral field). (N) Lateral view of male tail. (O) Male spicule and gubernaculum. Abbreviations: ep, excretory pore; ut, uterus; qc, quadricolumella; sp, spermatheca; od, oviduct; ov, ovary.
Fig. 2. Parasitic juvenile of *Sphaerularia vespa* sp. nov. (A) Lateral view. (B) Lateral view of anterior region. (C) Body annulation of anterior end. (D) Body annulation of mid-body. (E) Body annulation of posterior end. (F) Lateral view of tail. (G–O) Variety of tail tip (left lateral view). Arrowhead, constriction.
A New Endoparasite of a Common Hornet 1139

slide numbers *Sphaerularia vespae* F-06-10 (five females) and *Sphaerularia vespae* P-05-08 (four parasitic juveniles), deposited in The Herbarium and Insect Museum of the National Institute of Agro-Environmental Science, Tsukuba, Ibaraki, Japan; slide number *Sphaerularia vespae* F-11-13 (three females), M-02 (one male) and P-09-10 (two parasitic
juveniles), deposited in the Forest Pathology Laboratory Collection, Forestry and Forest Product Research Institute, Tsukuba, Japan.

**Etymology.** Specific epithet derived from *Vespa*, the generic name of the host insect.

**Distribution.** Besides the type locality, *S. vespae* sp. nov. was found on Mt. Moiwa, Sapporo, Hokkaido, Japan; Shiraikawa, Sapporo, Hokkaido, Japan; and Nishi-Nopporo, Ebetsu, Hokkaido, Japan.

**Molecular profiles.** The profiles obtained from the uterus and those from a single infective female were identical to each other. The partial SSU+ITS and D2/D3 sequences of *S. vespae* sp. nov., which were obtained from the single female, were deposited in the GenBank database under accession numbers AB300595 and AB300596, respectively. The closest sequences to these three molecular regions were from *S. bombi* (AB250213) for SSU, *Tylenchorhynchus leviterminalis* (EF030984) for ITS, and *S. bombi* (DQ328726) for D2/D3. The sequence similarities in the SSU and D2D3 regions between *S. vespae* sp. nov. and *S. bombi* are 99% (1,709 identical bases per 1,727 bp: 4 substitutions and 14 insertions/deletions) and 90% (639 identical bases per 708

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**Fig. 4.** Parasitic juvenile of *Sphaerularia vespae* sp. nov. (A) Lateral view. (B) Lateral view of anterior region (arrowhead, excretory pore). (C) Body annulation of mid-body. (D) Lateral view of tail (arrowhead, anus). (E–G) Variety of tail tip (left lateral view). Bar=100 μm in A; 40 μm in B–G.
Fig. 5. Mature parasitic female (uterium) of *Sphaerularia vespae* sp. nov. (arrowhead) and eggs and juveniles (small graining and lines, respectively, sometimes appearing cloudy) found during dissection of abdomen of *Vespa similima* gyne. Bar=5 mm.

Fig. 6. Glycerol-processed mature parasitic female (uterium) of *Sphaerularia vespae* sp. nov. (A) Entire uterium. (B) Genital pore (arrowhead) of uterus and eggs in the genital tract (close-up of left side A). (C) Body of *S. vespae* sp. nov. (arrowhead) attached to uterium (close-up of right side A). Bar=5 mm in A; 1 mm in B and C.
enlarged uteria of no mature eggs, captured with bait traps in the spring, have summer, asitzied gynes, which continue wandering outside in early 2007). This is consistent with the presumed behavior of par- parasitic juveniles. These juveniles grow into adults and wait for new hosts that happen to enter the over- wintering sites in autumn.

The life cycle of S. bombi is summarized by Bedding (1984) and Poinar and van der Laan (1972) as follows. Mated females invade bumblebee gynes that overwinter underground. Parasitized gynes usually do not find colonies due to their undeveloped ovaries (but see Röseler, 2002) but repeatedly visit potential overwintering sites in spring and early summer, where they disperse numerous parasitic juveniles. These juveniles grow into adults and mate, and wait for new hosts that happen to enter the overwintering sites in autumn.

Although the details of the life cycle of S. vespa sp. nov. are not yet clear, they appear basically similar to those of S. bombi. Parasitized gynes of V. simillima, which carry no mature eggs, captured with bait traps in the spring, have enlarged uteria of S. vespa sp. nov. in the gaster (Sayama et al., 2007). This strongly suggests that these gynes are incapable of establishing colonies, as is the case with parasitized bumblebees. Furthermore, parasitized gynes are frequently trapped in July and August, while unparasitized gynes are rarely captured in these months (Sayama et al., 2007). This is consistent with the presumed behavior of parasitized gynes, which continue wandering outside in early summer, visiting potential overwintering sites and dispersing parasitic juveniles.

The prevalence of S. vespa sp. nov. in overwintered gynes of V. simillima is often very high: more than 60% of gynes collected in June and July were infested with the nematode in Sapporo, northern Japan (Sayama et al., 2007). This implies that the nematode is potentially effective in controlling hornet populations. We do not know, however, whether the prevalence of S. vespa sp. nov. changes among populations of hornets or varies according to geographical region, or if it occurs in vine species other than V. simillima. Future studies should reveal these points as well as the details of the life cycle of S. vespa sp. nov. in relation to its host.

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