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Research article

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Two atypical new species of the genus *Sectonema* Thorne, 1930 (Nematoda, Dorylaimida, Aporcelaimidae) from Vietnam

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Abstract. Two new species of the genus *Sectonema* from natural habitats of northern Vietnam are studied. This paper includes their descriptions, measurements, line illustrations, and light microscope (LM) and scanning electron microscope (SEM) pictures. *Sectonema tropicum* sp. nov. is characterized by a 2.56–3.24 mm long body, 19–21 µm broad lip region, odontostyle 20–21 µm long at its ventral side, 730–834 µm long neck, pharyngeal expansion occupying 52–59% of total neck length, uterus a simple tube-like structure 150–242 µm long or 1.2–2.5 times the body diameter, *pars refringens vaginae* present, V = 48-52, short (31–40 µm, c = 70-91, c' = 0.5-0.6) and rounded tail, 91–97 µm long spicules, and only one weakly developed ventromedian supplement. *Sectonema vietnamense* sp. nov. is characterized by its slender (a = 33-49) and 2.71–4.25 mm long body, 14–16 µm broad lip region, odontostyle 8–9 µm long at its ventral side, 716–918 µm long neck, pharyngeal expansion occupying 63–67% of total neck length, uterus simple and 209–242 µm long or 2.5–2.9 times the corresponding body diameter, *pars refringens vaginae* absent, V = 54, short (34–39 µm, c = 70-115, c' = 0.6-0.8) and rounded tail, 59–75 µm long spicules, and three or four irregularly spaced ventromedian supplements bearing hiatus. Both species are also characterized by their nearly continuous lip region, an atypical feature in this genus. Molecular analysis of *S. tropicum* sp. nov. confirms that *Sectonema* is a natural (monophyletic) taxon, very close to *Metaporcelaimus*.

Keywords. D2–D3 expansion segments of LSU ribosomal DNA, morphology, morphometrics, taxonomy, SEM.

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Introduction

This contribution presents partial results of two nematological projects that overlap in their objectives. On the one hand, it is the second in a series of papers devoted to studying the nematode diversity of aporcelaims (family Aporcelaimidae Heyns, 1965) from natural areas of Vietnam, the first one (Álvarez-Ortega *et al.* 2015) dealing with the representatives of the genus *Aporcelaimoides* Heyns, 1965. On the other hand, it is part of a general revision of the taxonomy and systematics of this dorylaimid group, in which the genera *Aporcelaimellus* Heyns, 1965, *Metaporcelaimus* Lordello, 1965 and *Sectonema* Thorne, 1930 have already received some attention (see for instance, and respectively, Álvarez-Ortega & Peña-Santiago 2013; Álvarez-Ortega *et al.* 2013a; Peña-Santiago & Álvarez-Ortega 2014a).

Interesting material belonging to the genus *Sectonema* was collected by the first author during several nematological surveys conducted in natural areas of Vietnam over the last years. The specimens are characterized by a nearly continuous lip region, indeed a rather atypical feature in *Sectonema*. Its detailed study revealed it to belong to two unknown forms, which are herein described.

Material and methods

Nematodes

Nematodes were collected from pristine areas in Northern Vietnam, extracted from soil samples using the methods by Baermann (1917) and Flegg (1967), relaxed and killed by heat, fixed in 4% formaldehyde, and processed to anhydrous glycerine following Siddiqi's (1964) technique. Finally, the specimens were mounted on permanent glass slides to allow handling and observation under LM.

Light microscopy

Specimens were measured using a light Olympus BH-2 microscope equipped with differential interference contrast (DIC). Morphometrics included de Man's indices and the usual measurements. The location of the pharyngeal gland nuclei is expressed according to Loof & Coomans (1970) and spicule terminology follows Peña-Santiago *et al.* (2014). Some of the best preserved specimens were photographed with a Nikon Eclipse 80i microscope and a Nikon DS digital camera. Raw photographs were edited using Adobe® Photoshop® CS version 8.0.1. Drawings were made using a *camera lucida*.

Scanning electron microscopy

After examination and identification, a few of the preserved specimens were selected for observation under SEM following the protocol by Abolafia & Peña-Santiago (2005). The nematodes were hydrated in distilled water, dehydrated in a graded ethanol and acetone series, critical point dried, coated with gold, and observed with a Zeiss Merlin microscope.

DNA extraction, PCR and sequencing

DNA was extracted from individuals using the proteinase K protocol. Nematode material was transferred to an 0.5 ml Eppendorf tube containing 18 μ l of Worm Lysis Buffer (WLB) (50 mM KCL, 10 mM Tris pH 8.3, 2.5 mM MgCl₂, 0.45% NP 40, and 0.45% Tween 20) and 2 μ l proteinase K (600 μ g ml⁻¹) (Thermo Scientific). The tubes were incubated at 65°C (1 h) and then at 95°C (15 min). The PCR was performed in a 30 μ l final volume containing 24.9 μ l of sterile water, 0.6 μ l of each PCR primer, 0.6 μ l dNTP mixture, 0.3 μ l *Taq* polyerase, 3 μ l Buffer 10x Thermo Scientific Green and 1 μ l of DNA extracted solution. The PCR amplification profile consisted of four min at 94°C, 35 cycles of 1 min at 94°C, 1.5 min at 55°C and 2 min at 72°C, followed by a final step of 10 min at 72°C. The primers used for amplification were D2A (5'-ACAAGTACCGTGAAGGAAAGTTG-3') and D3B (5'-TCGGAAGGAACCAGCTACTA-3') for amplification of D2–D3 region of 28S (Subbotin *et al.* 2006).

The PCR products were purified using Exo-SAP PCR cleanup containing 7 μ l of PCR product, 0.15 μ l Exonuclease I (Exo), 0.9 μ l Shrimps Phosphatase Alkali (SAP) and 1.95 μ l of sterile water. They were incubated at 37°C for 20 min and then heated up to 85°C for 15 min; then the purified solution was diluted ten times.

The sequencing reaction was performed with 1 μ l of DNA purified template, 0.25 μ l BigDye v3.1, 2.25 μ l 5x BigDye sequencing buffer, 0.25 μ l of one primer. The mixture was heated up for 10 s at 96°C, 5s at 55°C repeated for 32 cycles followed by 4 min at 60°C. The sequencing was performed at the Cologne Center for Genomics (CCG). The sequences obtained were submitted to the GenBank database under accession numbers KT868957 and KT868958.

Phylogenetic analyses

The newly obtained sequences were aligned with other forty two D2–D3 expansion segments of 28S rRNA gene sequences available in GenBank using ClustalX 1.83 (Thompson *et al.* 1997). Outgroup taxa were chosen according to the results of previously published data (Holterman *et al.* 2008; Álvarez-Ortega *et al.* 2013b). Sequence alignments were manually edited using GenDoc 2.6.002 (Nicholas *et al.* 1997). The sequence dataset was analysed with Bayesian inference (BI) and Maximum Likelihood (ML) using MrBayes 3.1.2 (Huelsenbeck & Ronquist 2001; Ronquist & Huelsenbeck 2003) and MEGA 6 (Tamura *et al.* 2013), respectively. The best fit model of DNA evolution for BI was obtained using the program MrModeltest 2.2 (Nylander 2002) with the Akaike Information Criterion in conjunction with PAUP* 4b10 (Swofford 2003). BI analysis under the GTR + G + I model was initiated with a random starting tree and run with the four Metropolis-coupled Markov chain Monte Carlo (MCMC) for 10⁶ generations. ML analysis was implemented under the best-fitting evolutionary model (GTR + I + G), obtained using the program MEGA 6, and 1000 bootstrap replications. The topologies were used to generate a 50% majority rule consensus tree. Posterior probabilities (PP) are given on appropriate clades. The trees were visualised with the program FigTree v1.4.0 and drawn with Adobe Acrobat XI Pro 11.0.1

List of abbreviations

- a = body length / greatest body diameter
- b = body length / distance from anterior end to pharyngo-intestinal junction
- c = body length / tail length
- c' = tail length / tail diameter at anus or cloaca
- DN = distance from body anterior end to the nucleus of pharyngeal dorsal gland expressed as percentage (%) of total neck length
- DO = distance from body anterior end to the outlet of pharyngeal dorsal gland expressed as percentage (%) of total neck length
- L = overall body length
- n = number of specimens on which measurements are based
- S_1N_1 = distance from body anterior end to the anterior nucleus of first pair of ventro-sublateral pharyngeal glands expressed as percentage (%) of total neck length
- S_1N_2 = distance from body anterior end to the posterior nucleus of the first pair of ventro-sublateral pharyngeal glands expressed as percentage (%) of total neck length
- S_2N = distance from body anterior end to the nuclei of the second pair of ventro-sublateral pharyngeal glands expressed as percentage (%) of total neck length
- V = distance from body anterior end to vulva expressed as percentage (%) of the body length.

Results

Phylum Nematoda Cobb, 1932 Class Enoplea Inglis, 1983 Order Dorylaimida Pearse, 1942 Family Aporcelaimidae Heyns, 1965 Genus *Sectonema* Thorne, 1930

Sectonema tropicum sp. nov. urn:lsid:zoobank.org:act:014CA9CB-57EE-43BB-BBFF-0C1A7EBD522D Figs 1–4; Table 1

Diagnosis

The new species is characterized by a 2.56–3.24 mm long body, lip region hardly offset by very weak depression and 19–21 μ m broad, odontostyle 20–21 μ m long at its ventral side and 6.6–7.1 times as long as wide, 730–834 μ m long neck, pharyngeal expansion 403–470 μ m long or occupying 52–59% of total neck length, uterus a simple tube-like structure 150–242 μ m long or 1.2–2.5 times the body diameter, *pars refringens vaginae* present, *V* = 48–52, short (32–39 μ m, *c* = 70–88, *c'* = 0.5–0.6) and rounded female tail, male tail similar to that of female (31–40 μ m, *c* = 73–91, *c'* = 0.5–0.6), 91–97 μ m long spicules, and only one weakly developed ventromedian supplement.

Etymology

The specific epithet refers to the tropical area where the new species was collected.

Type material examined

Holotype

VIETNAM: \bigcirc , in acceptable state of preservation, Northern Vietnam, Cao Bang Province, Pia Oac Natural Reserve, 22° 36'28'' N, 105° 52'15'' E, tropical evergreen forest soil associated with *Machilus* sp. and *Dimocarpus* sp., deposited in the nematode collection of the University of Jaén, Spain.

Paratypes

VIETNAM: $4 \ \bigcirc \ \bigcirc, 3 \ \oslash \ \oslash, 1 \ \cup, 1 \$

Description

Adult

Moderately slender to slender nematodes of medium to big size, 2.56–3.24 mm long. Body cylindrical, distinctly tapering towards the anterior end, less so towards the posterior end as the caudal region is short and rounded. Habitus curved ventrad after fixation, especially in posterior body region, C-, G- or spiral-shaped. Cuticle $3.0-4.5 \mu$ m thick at anterior region, $5-6 \mu$ m in mid-body and $8-10 \mu$ m on tail; consisting of three layers, especially distinguishable at caudal region: thinner outer layer bearing very fine transverse striation across the entire body, thicker intermediate layer with radial striation, and thin inner layer. Lateral chords $13-21 \mu$ m wide at mid-body, occupying about one-sixth (13-18%) of mid-body diameter. Two ventral and two dorsal body pores often present at level of odontostyle-odontophore. Lip region hardly offset from the adjacent body by weak, but perceptible depression, $2.8-3.0 \mu$ m softy amalgamated, but their perioral part distinctly separated by the existence of six radial, interlabial, deep incisures delimiting six perceptible liplets; button-like labial papillae, the inner ones located at the margin of the oral field and surrounded by two or three concentric annuli, whereas the outer papillae, located



Fig. 1. *Sectonema tropicum* sp. nov. (Line). **A**. Lip region in lateral surface view. **B**. Anterior region in mid-optical level, lateral view. **C**. Neck region. **D**. \bigcirc , anterior genital branch. **E**. Vagina. **F**. \bigcirc , posterior body region. **G**. \bigcirc , posterior body region. **H**. \bigcirc , entire. **I**. \bigcirc , entire. **J**. Lateral guiding piece. **K**. Spicule.



Fig. 2. Sectonema tropicum sp. nov. (LM). **A**, **C**, **E**. Anterior region in lateral, median view. **B**. Lip region in lateral, surface view. **D**. Neck region. **F**. Pharyngo-intestinal junction. **G**. Vagina. **H**. \bigcirc , anterior genital branch. **I**. Vagina and anterior uterus containing sperm cells inside. **J**. Uterine egg. Scale bars: A–B, E, G = 10 µm; C, F, I–J = 20 µm; D, H = 50 µm.

a little behind the inner papillae, are surrounded by only one annulus; cephalic papillae pore- rather than button-like, also surrounded by only one ring-like annulus; oral aperture a dorso-ventral, slightly hexagonal orifice, the lip region hence showing a bi-radial symmetry. Amphid fovea cup-shaped, its opening occupying 9–10 μ m or less than one-half (43–49%) of lip region diameter. Cheilostom nearly cylindrical, without any differentiation. Stomatal protruding structure apparently a reduced odontostyle 6.6–7.1 times longer than wide (see: Remarks), its ventral side 1.0–1.1 times longer than lip region diameter and 0.65–0.83% of body length. Guiding ring simple, plicate, at 0.8–0.9 lip region diameters from the anterior end. Odontophore linear, rod-like, 1.7–1.9 times the odontostyle length, somewhat irregular at its base and with (in lateral view) the ventral side slightly longer than the dorsal one (Fig. 2A). Anterior region of pharynx enlarging very gradually; basal expansion 7.1–10.7 times longer than wide, 3.6–5.6 times as long as body diameter, and occupying 52–59% of total neck length; gland nuclei



Fig. 3. Sectonema tropicum sp. nov. (LM). **A**. \Diamond , posterior body region. **B**. \bigcirc , posterior body region. **C**. \bigcirc , caudal region. **D**. \Diamond , caudal region. **E**–**G**. Spicules, E–F also showing the lateral guiding piece. Scale bars: A–B = 20 µm; C–G = 10 µm.

often obscure, located as follows: DN = 60-62 (n = 4); $S_1N_1 = 75$ (n = 2); $S_1N_2 = obscure$; $S_2N = obscure$. Nerve ring located at 182–216 µm from anterior end or 25–26% of total neck length. Cardia rounded conoid, $13-17 \times 13-17$ µm; a weak ring-like structure is present surrounding its junction to pharyngeal base.

Female

Genital system didelphic-amphidelphic, with almost equally and well developed branches, the anterior 320–442 μ m or 12–15% of body length and the posterior 348, 440 μ m (n = 2) or 14, 16% of body length (447, 509 μ m or 15, 17% bearing one uterine egg inside). Reflexed ovaries well developed, often surpassing the sphincter level, the anterior 182–319 μ m, the posterior 207–380 μ m long; oocytes arranged first in two or more rows, then in a single row. Oviduct 124–174 μ m long or 1.1–1.8 times the



Fig. 4. *Sectonema tropicum* sp. nov. (SEM, \mathcal{J}). **A**. Lip region in ventral view. **B**. Lip region in frontal view. **C**. Lip region in lateral view. **D**. Lip region in sublateral view. **E**. Posterior body region in ventral view. **F**. Posterior body region in lateral view. **G**. Caudal region in sublateral view. Scale bars: $A-D = 5 \mu m$; $E-F = 20 \mu m$; $G = 10 \mu m$

corresponding body diameter, and consisting of a slender part made of prismatic cells and a moderately developed *pars dilatata*, with visible lumen and often containing sperm cells inside. Oviduct-uterus junction marked by a sphincter. Uterus a simple, tube-like structure 150–190 µm long or 1.2–1.7 times the corresponding body diameter [223, 242 µm (n = 2) long or 2.2, 2.5 times the corresponding body diameter, with one uterine egg inside; and 231, 240 µm (n = 1) long, with two uterine eggs], always containing abundant sperm cells inside. Vagina extending inwards 52–57 µm or one-half to foursevenths (48–57%) of body diameter: *pars proximalis* 33–43 × 23–28 µm, with somewhat sigmoid walls surrounded by weak musculature; *pars refringens* with two trapezoidal pieces measuring (in optical section) 13–16 × 7–9 µm and with a combined width of 24–28 µm; *pars distalis* 5.5–7.0 µm long. Vulva a nearly equatorial transverse slit. Prerectum 2.0–2.4, rectum 0.7–1.0 anal body diameters long. Tail short and rounded; two pairs of caudal pores, sublateral and close together.

Male

Genital system diorchic, with opposite testes. In addition to the ad-cloacal pair, situated at $18-21 \mu m$ from cloacal aperture, one weakly developed ventromedian supplement, lying out the range of retracted spicules and located at $80-131 \mu m$ from the ad-cloacal pair. Spicules distinctly robust and massive, especially in its posterior half, 3.6-4.2 times its maximum width, 1.4-1.6 times the body diameter at level of the cloacal aperture: dorsal contour regularly convex, ventral contour slightly concave, with distinct hump and hollow; curvature $121-124^{\circ}$; head occupying 11-16% of spicule total length, its dorsal side visibly curved at its anterior end and longer than the ventral one, which is shorter and almost straight; median piece 5.4-6.2 times longer than wide, occupying 51-59% of spicule maximum width, reaching the posterior tip; posterior end $7-8 \mu m$ wide. Lateral guiding pieces $23-29 \mu m$ long, 5.3-7.1 times longer than wide. Prerectum 3.2-4.6, cloaca 0.9-1.3 times the corresponding body width long. Cloacal aperture, as seen under SEM, a curved anteriad, transverse slit. Tail similar to that of female; caudal pores two pairs, one lateral, another subdorsal.

Relationships

In having a long stomatal protruding structure (20–21 µm at its ventral side) and short (c' up to 1.0) and rounded tail, S. tropicum sp. nov. is morphologically very similar to S. demani Altherr, 1965, S. heynsi Altherr, 1968, S. rotundicauda Goodey, 1951, S. septentrionale Peña-Santiago & Álvarez-Ortega, 2015 and S. sica Clark, 1964, but it can be distinguished from all of them in its smaller general size (body length 2.56–3.24 mm, neck 730–834 µm long vs more than 5 mm, more than 1000 µm long, respectively), nearly continuous lip region (vs offset by a more or less deep constriction) and stomatal protruding structure of different nature (a reduced odontostyle vs a mural tooth; see Remarks). Besides, the new species differs from S. demani (see recent re-description by Peña-Santiago & Álvarez-Ortega 2014b) in its less slender (a = 21-36 vs a = 48-56) body, narrower (19–21 vs 27–28 µm broad) lip region, shorter pharyngeal expansion (occupying 52–59 vs 68–69% of total neck length), shorter female tail (32–39 vs 51–55 μ m, c' = 0.5-0.6 vs c' = 0.7-0.8), and male present (vs absent). It differs from S. heynsi (see recent re-description by Peña-Santiago & Álvarez-Ortega 2014b) in its narrower (19-21 vs 28 µm broad) lip region, circumoral area lacking (vs bearing) cilia- or setae-like structures, shorter pharyngeal expansion (occupying 52–59 vs 68% of total neck length), shorter (32–39 vs 54 μ m, c' = 0.5–0.6 vs c' = 0.7) female tail, and male present (vs absent). It differs from S. rotundicauda in its less slender (a = 21-36 vs a = 80) body, narrower (19–21 vs about 28 µm broad) lip region, shorter (31–40 vs 64 um) male tail, fewer (one vs four) ventromedian supplements, and shorter (91–97 vs about 155 µm) spicules. It differs from S. septentrionale in its less slender (a = 21-36 vs a = 47-64) body, narrower (19-21 vs 25-29 µm broad) lip region, shorter (occupying 52-59 vs 60-73% of total neck length) pharyngeal expansion, uterus simple (vs tripartite) and shorter (150-242 µm or 1.2-2.5 times the body diameter vs 370–493 µm or 2.9–4.3 times the corresponding body diameter), shorter (31–40 vs 40–62 μ m, c' = 0.5-0.6 vs c' = 0.6-0.8) tail, fewer (one vs 5–11) ventromedian supplements, and shorter (91-97 vs 100-145 µm) spicules. Finally, it differs from S. sica in its narrower (19-21 vs about 24 µm

		Cao Bằng Province (Pia Oac Natural Reserve)	
Population Character	Holotype	Paratypes	
	Ŷ	5 ♀♀	5
L	3.04	2.81 ± 0.19 (2.57–3.05)	2.87 ± 0.31 (2.56–3.24)
a	31	24.9 ± 4.7 (21–31)	29.0 ± 5.9 (23–36)
b	?	3.7 ± 0.2 (3.4–3.9)	3.7 ± 0.3 (3.4–3.9)
c	87	79.4 ± 6.5 (70-88)	81.3 ± 7.0 (73–91)
с'	0.6	$0.6 \pm 0.0 \ (0.5 - 0.6)$	$0.6 \pm 0.1 \ (0.5 - 0.6)$
V	50	50.6 ± 1.5 (48-52)	_
Lip region diam.	20	20.1 ± 0.6 (20-21)	20.1 ± 0.6 (19–21)
Odontostyle length at ventral side	21	20.6 ± 0.6 (20-21)	20.7 ± 0.5 (20–21)
Odontostyle length at dorsal side	23	$22.5 \pm 0.5 (22 - 23)$	$22.5 \pm 0.4 (22 - 23)$
Odontophore length	36	36.0 ± 1.0 (35–38)	35.3 ± 0.7 (34–36)
Guiding ring from ant. end	17	17.4 ± 0.6 (17–18)	18.1 ± 0.1 (18–18)
Neck length	?	764 ± 23 (730–790)	801 ± 49 (745–834)
Pharyngeal expansion length	?	427 ± 29 (403–468)	448 ± 19 (436–470)
Diam. at neck base	99	103 ± 12 (91–116)	95 ± 11 (84–111)
at midbody	99	114 ± 14 (98–130)	101 ± 12 (90–118)
at anus	60	64.4 ± 4.5 (61–72)	61.2 ± 2.1 (60–65)
Prerectum length	143	136 ± 9 (126–150)	228 ± 37 (189–277)
Rectum/cloacla length	54	53.8 ± 3.0 (52–59)	66.3 ± 8.8 (53–77)
Tail length	35	35.5 ± 3.3 (32–39)	35.4 ± 3.0 (31–40)
Spicule length	_	_	$92.5 \pm 2.6 \ (91-97)$
Ventromedian supplements	-	_	$1 \pm 0 (1-1)$

Table 1. Morphometrics of *Sectonema tropicum* sp. nov. Measurements in μ m (except L, in mm), and in the form: mean \pm standard deviation (range).

broad) lip region, shorter (occupying 52–59% vs two-thirds of total neck length) pharyngeal expansion, more posterior (V = 48-52 vs V = 40) vulva, shorter (32–39 vs 54 µm, c' = 0.5-0.6 vs c' = 0.7) female tail, and male present (vs absent).

Moreover, in having a lip region nearly continuous with the adjacent body, the new species resembles *S. mucrodens* Siddiqi, 1984 and *S. truxum* Siddiqi, 1984, but it can be easily distinguished from these by the more anterior position of the stomatal protruding structure (its anterior tip in front of the level of guiding ring, just behind the base of lip region vs its anterior tip distinctly behind the level of guiding ring, far from the lip region base) and *pars refringens vaginae* well (vs faintly) developed. Moreover, it differs from *S. mucrodens* by its shorter (730–834 vs 1086–1157 μ m) neck, shorter (occupying 52–59 vs 69% of total neck length) pharyngeal expansion, shorter (32–39 μ m, *c*' = 0.5–0.6 vs 44–50 μ m, *c*' = 0.7) female tail, and male present (vs absent). It differs from *S. truxum* in its narrower (19–21 vs 24 μ m broad) lip region, shorter (730–834 vs 837–1070 μ m) neck, more anterior (*V*= 48–52 vs *V*= 54–56) vulva, and male present (vs absent).

Molecular characterisation

Two sequences of the D2–D3 28S rRNA gene were obtained from one female and one male. Both sequences are very similar (99%) and differ by two nucleotides only. The evolutionary relationships of the new species with representatives of the order Dorylaimida are presented in Figs 5–6. The two *S. tropicum* sp. nov. sequences are clustered together with other available sequences of *Sectonema* representatives and form a highly supported clade together with members of the genus *Metaporcelaimus*, thus confirming previous results (Álvarez-Ortega *et al.* 2013a, b).

Remarks

The nature of the stomatal protruding structure in the new species is difficult to interpret and deserves further explanation. The specimens examined bear a reduced odontostyle (comparable to that described in the type species *S. ventrale* Thone, 1930; see recent description by Peña-Santiago & Álvarez-Ortega



Fig. 5. Bayesian 50% majority rule consensus trees as inferred from D2–D3 expansion segments of 28S rRNA gene sequence alignments under the GTR + I + G model. Posterior probabilities are given for appropriate clades. Newly obtained sequences are indicated by bold letters.

2014a) rather than a mural tooth as the base of its dorsal side is visibly sclerotized and seems to join the dorsal stomatal wall (vs dorsal side not scletorized at the base, somewhat sigmoid and not joining the dorsal stomatal wall). Nonetheless, some doubt persists in this matter.



Fig. 6. Maximum Likelihood tree as inferred from D2–D3 expansion segments of 28S rRNA gene sequence alignments under the GTR + I + G model. Bootstrap values are given for appropriate clades. Newly obtained sequences are indicated by bold letters.

Sectonema vietnamense sp. nov.

urn:lsid:zoobank.org:act:AF58451E-D7A7-4E49-8A18-9F3998EF4A87

Figs 7–9, Table 2

Diagnosis

The new species is characterized by a slender (a = 33-49) 2.71–4.25 mm long body, lip region offset by depression and 14–16 µm broad, odontostyle 8–9 µm long at its ventral side, 716–918 µm long neck, pharyngeal expansion 463–618 µm long or 63–67% of total neck length, uterus simple and 209–242 µm long or 2.5–2.9 times the corresponding body diameter, *pars refringens vaginae* absent, V = 54, short (36 µm, c = 105, c' = 0.7) and rounded tail in female and slightly more conoid (34–39 µm, c = 70–115, c' = 0.6–0.8) in male, 59–75 µm long spicules, and three or four irregularly spaced ventromedian supplements bearing hiatus.

Etymology

The specific epithet refers to the geographical origin of the species.

Type material examined

Holotype

VIETNAM: \bigcirc , in good state of preservation, Northern Vietnam, Cuc Phuong National Park, pristine tropical forest, deposited in the nematode collection of the University of Jaén, Spain.

Paratypes

Other material examined

VIETNAM: 2 3 3, in acceptable state of preservation, Northern Vietnam, Thái Bình province, intensively cultured field, deposited in the nematode collection of the University of Jaén, Spain.

Description (type population)

Adult

Very slender nematodes of big size, 3.13–4.25 mm long. Body cylindrical, distinctly tapering towards the anterior end, less so towards the posterior one as the caudal region is rounded. Habitus regularly curved ventrad after fixation, to a more or less open C, occasionally more curved at posterior body region. Cuticle 2.0–2.5 µm thick at anterior region, 3.0–4.5 µm in mid-body and 8–10 µm on tail; consisting of three layers especially distinguishable at caudal region: thinner outer layer bearing very fine transverse striation through the entire body, thicker intermediate layer with radial striation, and thin inner layer. Lateral chords $6-12 \mu m$ wide at mid-body, occupying up to one-sixth (8-15%) of mid-body diameter. Two ventral and two dorsal body pores often present at level of stoma, and four distinct lateral pores 6-20 µm apart visible behind the amphid fovea. Lip region nearly continuous (a shallow depression is, however, often perceptible), weakly angular, visibly narrower than adjacent body, 2.1–2.3 times wider than high and about one-fifth (18–23%) of body diameter at neck base; it appears (under SEM) marked by radial (oral field) or longitudinal (remaining lip region) incisures, with amalgamated lips and protruding, pore-like labial and cephalic papillae, all of them surrounded by a ring-like annulus, and the inner labial papillae visibly larger than the outer ones; oral field perceptibly hexagonal, oral aperture a dorso-ventral slit, the lip region hence showing a bi-radial symmetry. Cupshaped amphid fovea, its opening at level of cephalic depression and occupying 10-11 µm or 68-72% of lip region diameter. Nearly cylindrical cheilostom, without differentiation. Odontostyle comparatively



Fig. 7. Sectonema vietnamense sp. nov. (Line). **A**. Anterior region in median, lateral view. **B**. Lip region in surface, lateral view. **C**. Neck region. **D**. Vagina. **E**. \mathcal{Q} , anterior genital branch. **F**. \mathcal{J} , entire. **G**. \mathcal{Q} , entire. **H**. \mathcal{J} , posterior body region. **J**. Spicule. **K**. Lateral guiding piece.

short and occupying the whole stomatal lumen, its ventral side 0.5–0.6 times longer than lip region diameter and 0.20–0.29% of body length. Guiding ring simple, somewhat plicate, at 0.5 times the lip region diameter from anterior end. Odontophore linear, rod-like, 3.7–4.2 times the odontostyle length. Tripartite pharynx, consisting of an anteriorly thickened section behind the odontophore base, a perceptible narrower intermediate section with the nerve ring surrounding it, which enlarges very gradually into the basal expansion 9.9–15.0 times longer than wide, 6.7–7.3 times longer than body



Fig. 8. Sectonema vietnamense sp. nov. (LM, type population). A. \mathcal{S} , entire. **B**–C. Anterior region in lateral, median view. **D**. \mathcal{Q} , anterior genital branch. **E**. \mathcal{S} , posterior body region. **F**. Oviduct-uterus junction. **G**. Lip region in lateral, surface view. **H**. Vagina. **I**. \mathcal{S} , caudal region. **J**–K. Spicules. **L**. \mathcal{Q} , caudal region. Scale bars: A = 500 µm; B–C, H = 10 µm; D = 100 µm; E–F = 50 µm; G = 5 µm; I–L = 20 µm.

diameter and occupying 63–67% of total neck length; gland nuclei mostly obscure, only S₂N are visible, at 77–78% (n = 3) of total neck length. Nerve ring located at 147–178 μ m from anterior end or 19–22% of total neck length. Cardia rounded conoid, 15–19 × 12–17 μ m; a ring-like structure is present surrounding its junction to pharyngeal base.

Female

Genital system didelphic-amphidelphic, with almost equally and well developed branches, the anterior 473 μ m long or 13% of body length and the posterior 387 μ m long or 10% of body length, each branch bearing one uterine egg. Moderately sized ovaries, not surpassing the sphincter level, the anterior 191 μ m, the posterior 141 μ m long; oocytes arranged first in two or more rows, then in a single row. Oviduct 117, 142 μ m long or 1.4, 1.7 times the corresponding body diameter, and consisting of a slender part made of prismatic cells and a well developed *pars dilatata* with wide lumen and sperm cells inside. Oviduct-uterus junction marked by a sphincter. Uterus a simple, tube-like structure 209, 242 μ m long or 2.5, 2.9 times the corresponding body diameter, containing abundant sperm cells and one uterine egg. Vagina extending inwards 36 μ m or about three-sevenths (44%) of body diameter: *pars proximalis* 23 × 28 μ m, somewhat sigmoid walls and surrounded by weak musculature; *pars refringens* absent; *pars*



Fig. 9. Sectonema vietnamense sp. nov. (SEM, A = juvenile and B–J = 3). A. Lip region in ventral view. **B**–C. Lip region in lateral view. **D**. Posterior body region in ventral view. **E**. Lip region in frontal view. **F**. Detail of a ventromedian supplement. **G**. Caudal region in lateral view. **H**. Caudal region in subventral view. **I**. Detail of the ad-cloacal pair of supplements. **J**. Caudal region in ventral view. Scale bars = A–C, E = 2 µm; D = 20 µm; F, I = 1 µm; G–H = 10 µm; J = 5 µm.

	C	uc Phuong National Park	Thái Bình Province
Character	Holotype	Paratypes	
	9	6 ඊඊ	2 ්්
L	3.79	3.57 ± 0.49 (3.13-4.25)	3.02, 2.71
а	46	45.9 ± 3.0 (41–49)	35, 33
b	4.6	4.5 ± 0.2 (4.3–4.7)	?
c	105	96.9 ± 11.4 (86–115)	83, 70
c'	0.7	$0.7 \pm 0.1 \ (0.6 - 0.8)$	0.8, 0.7
V	54	_	_
Lip region diam.	14	$14.9 \pm 0.5 (14 - 16)$	16 (n = 2)
Odontostyle length at ventral side	9	8.8 ± 0.3 (8–9)	9 (n = 2)
Odontostyle length at dorsal side	10	9.5 ± 0.2 (9–10)	10 (n = 2)
Odontophore length	32	34.2 ± 1.3 (33–36)	33 (n = 2)
Guiding ring from ant. end	7	7.3 ± 0.6 (7–8)	9 (n = 2)
Neck length	820	790 ± 84 (716–918)	?
Pharyngeal expansion length	534	521 ± 65 (463–618)	?
Diam. at neck base	?	72.0 ± 8.6 (63–85)	80, 74
at midbody	82	77.7 ± 8.3 (68-89)	87, 82
at anus	50	$50.6 \pm 4.3 \ (46 - 58)$	48, 55
Prerectum length	149	237 ± 40 (210–297)	199, ?
Rectum/cloaca length	45	58.8 ± 9.3 (47–71)	66, 62
Tail length	36	36.8 ± 1.8 (34–39)	36, 38
Spicule length	_	67.7 ± 5.3 (59–75)	73, 70
Ventromedian supplements	_	3.7 ± 0.5 (3–4)	4 (n = 2)

Table 2. Morphometrics of *Sectonema vietnamense* sp. nov. Measurements in μ m (except L, in mm), and in the form: mean \pm standard deviation (range).

distalis well developed, 15 µm long. Vulva a post-equatorial transverse slit. Prerectum 3.0, rectum 0.9 anal body diameters long. Short and rounded tail, its inner core with a finger-like projection at tail end; two pairs of caudal pores, one sub-lateral, another sub-dorsal.

Male

Genital system diorchic, with opposite testes. In addition to the ad-cloacal pair, situated at 14–19 μ m from cloacal aperture, there is a series of three or four irregularly spaced, 15–48 μ m apart, ventromedian supplements, the posteriormost of which lying out the range of retracted spicules, at 76–91 μ m from the ad-cloacal pair. Spicule robust and massive, especially in its posterior half, 3.6–4.6 times its maximum width, 1.3–1.4 times the body diameter at level of the cloacal aperture: dorsal contour regularly convex, ventral contour slightly concave, with distinct hump and hollow; curvature 125–136°; head occupying 13–19% of spicule total length, its dorsal contour distinct curved at its anterior end and longer than the ventral one, which is short and almost straight; median piece 5.9–7.1 times longer than wide, occupying 52–65% of spicule maximum width, reaching the posterior tip; posterior end 6–7 μ m wide. Lateral guiding pieces 15–17 μ m long, 5.3–6.5 times longer than wide. Prerectum 4.3–5.7, cloaca 1.0–1.3 times the corresponding body width long. Rectal glands and their ducts (*cf.* Coomans & Loof 1986) very well perceptible (Fig. 8K). Cloacal aperture, as seen under SEM, a somewhat curved anteriad, transverse slit; the two supplements of the precloacal pair distinctly separated. Tail a little more conoid than that of female; two pairs of caudal pores, one almost lateral, another sub-dorsal.

Relationships

The new species is easily distinguishable from its congeners in the unique combination of these features: narrow and nearly continuous lip region, very short odontostyle, absence of pars refringens vaginae and short and rounded tail. Nonetheless, it resembles S. barbatum Heyns, 1965 in its general morphology, but it can be distinguished from this because of its oral field lacking any setae-like structure (vs bearing numerous short forwardly pointing setae), narrower (14–16 vs about 22 µm broad) lip region, shorter (8–9 vs 16 μ m) odontostyle, pars refringens vaginae absent (vs present), shorter (36 μ m, c = 105, c' = $0.7 \text{ vs } 48 \mu\text{m}, c = 81, c' = 1.0$) female tail, and male present (vs absent). In having a comparatively small body size and nearly continuous lip region, S. vietnamense sp. nov. is close to S. mucrodens and S. truxum, from which it differs in its narrower (14–16 vs more than 20 µm broad) lip region, shorter stomatal protruding structure (8–9 vs more than 20 μ m) located as usual (vs the anterior tip distinctly behind the level of guiding ring), and pars refringens vaginae absent (vs consisting of two small pieces). Sectonema vietnamense sp. nov. is also similar to S. pseudoventrale Heyns, 1965 in having a short odontostyle, but can be distinguished from this by its narrower (14–16 vs 21–23 µm broad) lip region, shorter (209–242 µm or 2.5-2.9 times the corresponding body diameter vs 123 μ m or 1.3 times the corresponding body diameter) uterus, shorter (36 μ m, c = 105, c' = 0.7 vs 46–63 μ m, c = 82–130, c' = 0.8–1.0) female tail, shorter (59– 75 vs 90–95 μ m) spicules, and fewer (three or four vs six or seven) ventromedian supplements. Finally, the new species also resembles S. tropicum sp. nov., but it can be distinguished from this by its narrower (14–16 vs 19–21 µm broad) lip region, shorter (8–9 vs 20–21 µm at its ventral side) odontostyle, longer (occupying 63–67 vs 52–59% of total neck length) pharyngeal expansion, more posterior vulva (V = 54vs V =48–52), pars refringens vaginae absent (vs present), shorter (59–75 vs 91–97 μ m) spicules and higher number of ventromedian supplements (three or four vs one).

Remarks

The two males from Thái Bình are very similar to those of the type population, but some minor morphometric differences have been also noted: somewhat smaller (2.71, 3.02 vs 3.13-4.25 mm long) and more obese (a = 33, 35 vs a = 41-49) body. Such differences have been regarded as instraspecific variation as the number of specimens examined is low and the main morphological and morphometric features are coincident.

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References

Abolafia J. & Peña-Santiago R. 2005. Nematodes of the order Rhabditida from Andalucía Oriental: *Pseudacrobeles elongatus* (de Man, 1880) comb. n. *Nematology* 7 (6): 917–926. <u>http://dx.doi.org/10.1163/156854105776186415</u>

Álvarez-Ortega S., Nguyen T.A.D., Abolafia J., Vu T.T.T. & Peña-Santiago R. 2015. Three new species of the genus *Aporcelaimoides* Heyns, 1965 from Vietnam (Nematoda, Dorylaimida, Aporcelaimidae), and an updated taxonomy of the genus. *ZooKeys* 516: 1–26. <u>http://dx.doi.org/10.3897/zookeys.516.10087</u>

Álvarez-Ortega S. & Peña-Santiago R. 2013. Taxonomy of the genus *Aporcelaimellus* Heyns, 1965 (Nematoda, Dorylaimida, Aporcelaimidae). *Zootaxa* 3669: 243–260. <u>http://dx.doi.org/10.11646/zootaxa.3669.3.3</u>

Álvarez-Ortega S., Subbotin S.A. & Peña-Santiago R. 2013a. Morphological and molecular characterization of Californian species of the genus *Metaporcelaimus* Lordello, 1965 (Dorylaimida, Aporcelaimidae), with a new concept of the genus. *Nematology* 15 (3): 251–278. <u>http://dx.doi.org/10.1163/15685411-00002674</u>

Álvarez-Ortega S., Subbotin S.A. & Peña-Santiago R. 2013b. Morphological and molecular characterization of Californian species of the genus *Aporcelaimellus* Heyns, 1965 (Dorylaimida: Aporcelaimidae). *Nematology* 15 (4): 431–439. <u>http://dx.doi.org/10.1163/15685411-00002691</u>

Baermann G. 1917. Eine einfache Methode zur Auffindung von *Ankylostomum* (Nematoden) Larven in Erdproben. *Geneeskunding Tijdschrift voor Nederlandsch-Indië* 57: 131–137.

Coomans A. & Loof P.A.A. 1986. Observations on the glands of the male reproductive system in dorylaims and its phylogenetic importance. *Revue de Nématologie* 9 (3): 261–265.

Flegg J.J.M. 1967. Extraction of *Xiphinema* and *Longidorus* species from soil by a modification of Cobb's decanting and sieving technique. *Annals of Applied Biology* 60 (3): 429–437. <u>http://dx.doi.org/10.1111/j.1744-7348.1967.tb04497.x</u>

Holterman M., Rybarczyk K., van den Elsen S., van Megen H., Mooyman P., Peña-Santiago R., Bongers T., Bakker J. & Helder J. 2008. A ribosomal DNA-based framework for the detection and quantification of stress-sensitive nematode families in terrestrial habitats. *Molecular Ecology Resources* 8 (1): 23–34. http://dx.doi.org/10.1111/j.1471-8286.2007.01963.x

Huelsenbeck J.P. & Ronquist F. 2001. MRBAYES: Bayesian inference of phylogenetic trees. *Bioinformatics* 17 (8): 754–755. <u>http://dx.doi.org/10.1093/bioinformatics/17.8.754</u>

Loof P.A.A. & Coomans A. 1970. On the development and location of the oesophageal gland nuclei in the Dorylaimina. *Proceedings of the IXth International Nematology Symposium (Warsaw, Poland, 1967)*: 79–161.

Nicholas K.B., Nicholas Jr H.B. & Deerfield II D.W. 1997. GeneDoc: analysis and visualization of genetic variation. *EMBnet News* 4: 1–14.

Nylander J.A.A. 2002. MrModeltest v1.0b. Department of Systematic Zoology, Uppsala University.

Peña-Santiago R., Abolafia J. & Álvarez-Ortega S. 2014. New proposal for a detailed description of the dorylaim spicule (Nematoda: Dorylaimida). *Nematology* 16 (9): 1091–1095. <u>http://dx.doi.org/10.1163/15685411-00002834</u>

Peña-Santiago R. & Álvarez-Ortega S. 2014a. Studies on the genus *Sectonema* Thorne, 1930 (Dorylaimida: Aporcelaimidae). Re-description of *S. ventrale*, the type species of the genus. *Nematology* 16 (9): 1097–1104. http://dx.doi.org/10.1163/15685411-00002837

Peña-Santiago R. & Álvarez-Ortega S. 2014b. Re-description of three species of the genus *Sectonema* Thorne, 1930 (Nematoda: Dorylaimida: Aporcelaimidae) originally studied by E. Altherr. *Zootaxa* 3881: 63–74. <u>http://dx.doi.org/10.11646/zootaxa.3881.1.5</u>

Ronquist F. & Huelsenbeck J. 2003. MrBayes 3: Bayesian phylogenetic inference under mixed models. *Bioinformatics* 19 (12): 1572–1574. <u>http://dx.doi.org/10.1093/bioinformatics/btg180</u>

Siddiqi M.R. 1964. Studies on *Discolaimus* spp. (Nematoda: Dorylaimidae) from India. *Zeitschrift für Zoologische Systematik und Evolutionsforschung* 2: 174–184. <u>http://dx.doi.org/10.1111/j.1439-0469.1964.</u> tb00720.x

Subbotin S.A., Sturhan D., Chizhov V.N., Vovlas N. & Baldwin J.G. 2006. Phylogenetic analysis of Tylenchida Thorne, 1949 as inferred from D2 and D3 expansion fragments of the 28S rRNA gene sequences. *Nematology* 8 (3): 455–474. http://dx.doi.org/10.1163/156854106778493420

Swofford D.L. 2003. *PAUP*: phylogenetic analysis using parsimony (*and other methods)*. Version 4.0b 10. Sunderland, MA, USA, Sinauer Associates.

Tamura K., Stecher G., Peterson D., Filipski A. & Kumar S. 2013. MEGA6: Molecular Evolutionary Genetics Analysis version 6.0. *Molecular Biology and Evolution* 30: 2725–2729. <u>http://dx.doi.org/10.1093/molbev/mst197</u>

Thompson J.D., Gibson T.J., Plewniak F., Jeanmougin F. & Higgins D.G. 1997. The ClustalX windows interface: flexible strategies for multiple sequence alignment aided by quality analysis tools. *Nucleic Acids Research* 25 (24): 4876–4882. <u>http://dx.doi.org/10.1093/nar/25.24.4876</u>

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