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First report of *Hemicriconemoides litchi* associated with *Piper sarmentosum* and revision of the genus *Hemicriconemoides* in Vietnam

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Abstract

The taxonomic status of *Hemicriconemoides litchi*, *H. mangiferae*, and *H. strictathecatus* has been in debates for many years. In this study, a population of *H. litchi* collected from the rhizosphere of *Piper* sarmentosum is characterized using a combination of morphology and molecular data. Our results are in agreement with other authors to maintain the validity of *H. litchi* and provide sequences of *H. litchi* with correct names on GenBank. A revision on the genus *Hemicriconemoides* in Vietnam is also discussed.

Keywords

H. brachyurus, H. cocophilus, H. microdoratus, Hemicriconemoides mangiferae, Sheathoid nematode, Wild betel, Wild pepper.

The sheathold nematodes. *Hemicriconemoides* (Chitwood and Birchfield, 1957), are root-ectoparasitic nematodes that cause damage to various crops (Sikora et al., 2018) with 54 valid species worldwide (Maria et al., 2018). In Vietnam, five Hemicriconemoides species have been reported including H. litchi (Misra and Edward, 1963), H. mangiferae (Siddigi, 1961), H. cocophilus (Loos, 1949; Chitwood and Birchfield, 1957), H. microdoratus (Dasgupta et al., 1969), and H. brachyurus (Loos, 1949; Chitwood and Birchfield, 1957; Dasgupta et al., 1969; Nguyen and Nguyen, 2000). However, to the best of our knowledge, molecular data of Hemicriconemoides spp. in Vietnam are not available, and according to recent debates on the genus Hemicriconemoides, the evaluations for nominal species in the country are needed.

Herein, a population of *H. litchi* associated with *Piper* sarmentosum Roxb. in Vietnam is characterized by morphological and molecular characterizations. Besides, a revision of the genus *Hemicriconemoides* in Vietnam is also provided.

Material and methods

Nematodes were extracted from the rhizosphere samples of *P. sarmentosum* in Vinh Phuc province, Vietnam, using the modified Baermann tray method (Whitehead and Hemming, 1965). For morphological characterization, permanent slides were made (Nguyen et al., 2017). Photographs and measurements were obtained using a Carl Zeiss Axio Lab. A1 light microscope was used equipped with a digital camera.

© 2020 Authors. This is an Open Access article licensed under the Creative Commons CC BY 4.0 license, https://creativecommons.org/licenses/by/4.0/ The 5'-end region of 28S rDNA was amplified using DP391/501 primers (5'-AGCGGAGGAAAAGAAACTAA-3'/5'-TCGGAAGGAACCAGCTACTA-3') and analyzed using Geneious R11 (Nguyen et al., 2019). JB3/JB4 primers (5'-TTTTTTGGGCATCCTGAGGTTTAT-3'/5'-TAA AGAAAGAACATAATGAAAATG-3') were used to amplify *COI* mtDNA gen region (Nguyen et al., 2019). References on *Hemicriconemoides* spp. in Vietnam were collected and evaluated.

Results and discussion

H. litchi Misra and Edward, 1963

Measurements and specimens

After Eroshenko et al. (1985) (population from Vietnam): Details of female are as follows (n=10): L=480 to 520 µm; a=14 to 21; b=4.6 to 5.2; c=13 to 15; V=91 to 93%; stylet=62 to 65 µm; R=120 to 131; Rst=20 to 22; Rex=29 to 32; RV=11 to 13; RVan=5; Ran=6 to 8.

After Nguyen et al. (2020) (population from Vietnam, in this study):

Details of female are as follows (n = 20): $L = 567 \pm 26$ (537-609)µm, $a = 18.3 \pm 1.2$ (16.8-21), $b=5.3\pm0.2$ (5-5.7), $c=28\pm4$ (24-35), $c'=1.1\pm0.1$ $(1-1.3), V=92\pm1$ (91-93) %, VL/VB= 1.6 ± 0.1 (1.5-1.7),lip width = 11.6 ± 0.7 (10.7-12.7) µm, lip height = 5.5 ± 0.6 $(4.7-6.4)\mu m$, metenchium = 54 ± 2 (51-57) μm , telenchium = 11.4 ± 2 (9.2-14) µm, knob = 3.5 ± 0.4 (3-4.1) μ m, stylet=69±3 (65-73) μ m, DGO=4±1 (2.9-5.4) μ m, anterior end to secretory-excretory pore = 136 ± 9 (126-151) μ m, anterior end to nerve ring=89±4 (81-95) μ m, pharynx = 107 ± 3 (104-113) μ m, width at midbody= 31 ± 1 (29-32) µm, width at anus= 18 ± 1 (16.5-20) μ m, tail=21±3 (17-24) μ m, R=120±6 (110-130), $Rst = 18 \pm 0.8$ (17-19), $Roes = 25 \pm 1$ (24-26), $Rex = 32 \pm 2$ (30-34), RV = 10 ± 0.4 (10-11), Rvan = 4.9 ± 0.6 (4-6).

Remarks

The female of *H. litchi* from Vietnam is characterized by having a close sheath fitting; lip region bearing two annuli with first lip annulus protruding outward; frequently curved dorsally stylet with rounded stylet knobs; secretory-excretory pore located approximately seven annuli posterior to the end of pharynx; oval spermatheca full of sperm; and tail with bluntly rounded tail tip (Fig. 1). Morphology and morphometrics of *H. litchi* from Vietnam are in agreement with the description of the type population with small variations that can be seen from other populations of *H. litchi* reported by Van Den Berg et al. (2015). For molecular characterization, the 28S rDNA sequence of *H. litchi* from Vietnam with 938 bp (accession number: MT539313) was 99.8 to 99.9% similar (1-2 bp difference) to sequences of H. litchi from other populations (accession number: KP192481, KF856540, AY780956). The phylogenetic tree based on 28S rDNA sequences showed that the sequence of *H. litchi* from Vietnam was grouped in the same clade (100% PP) with sequences of H. litchi from other populations (Fig. 2). The validation of H. litchi has been revised by many studies (Siddigi, 2000; Geraert, 2010; Chen et al., 2011). Recently, a study of Van Den Berg et al. (2015) agreed with Chen et al. (2011) to maintain the validity of *H. litchi*; however, the sequences of H. litchi provided by the authors still appear as H. strictathecatus on GenBank (accession number: KF856540, KF856541, AY780956, AY780957, AY780958) confusing the identification process, especially for studies using molecular approach only. This study provides the first report of H. lichi associated with P. sarmentosum using an integrated approach of morphology and molecular tools, a 28S rDNA sequence with correct name was also submitted to GenBank. Two COI mtDNA sequences of H. litchi from Vietnam were also obtained and submitted to GenBank (accession number: MT586696, 586697). These are first COI mtDNA sequences of H. litchi on GenBank.

H. mangiferae Siddiqi, 1961

Measurements and specimens

After Eroshenko et al. (1985) (population from Vietnam): Details of female are as follows (n = 10): L = 450 to 560 µm; a = 17 to 22; b = 4 to 5; c = 17 to 24; V = 91 to 94%; stylet=67 to 72 µm; R = 123 to 136; Rst=18 to 20; Roes=26 to 28; Rex=34 to 36; RV=10 to 13; Ran=5 to 8.

After Germani and Anderson (1991) (population from coffee, Vietnam):

Details of female are as follows (n=8): $L=460\pm3$ (400-500) µm; $a=15\pm1.3$ (13.5-16.9); $b=4.7\pm0.1$ (4.5-5); $c=20.8\pm2.58$ (16.5-28.8); $V=92.6\pm0.91$ (91.2-94.1) %, $V'=34\pm4.4$ (28-38); $R=125\pm6.6$ (116-136); Rex=34±3.0 (32-40); Rv=10±1.7 (8-13); Ran=6±2.0 (4-9); Rvan=4±0.6 (3-5); VL/VB=1.42±0.14 (1.22-1.62); stylet=63±1.7 (62-65) µm.

Remarks

Although Chen et al. (2011) differentiated *H. mangiferae* from *H. litchi* using the combination of morphological characterizations and molecular data, Van Den Berg et al. (2015) proposed to transfer



Figure 1: Female of H. litchi from Vietnam. A: Entire body; B: Pharyngeal region; C: Annuli at mid-body; D: Tail region.

species status of *H. mangiferae* provided by Chen to other populations that previously was identified as et al. (2011) to H. strictathecatus. On the contrary, H. mangiferae, including the revision of the population Van Den Berg et al. (2015) also gave the corrections of *H. mangiferae* provided by Germani and Anderson

H. litchi and revision of Hemicriconemoides spp. in Vietnam: Nguyen et al.



0.020

Figure 2: BI phylogenetic tree generated from 28S rDNA sequences (GTR+G+I model, 1×10^6 generations, 20% Burn-in). Bayesian posterior probabilities (in percentage) are given next to each node. Sequences of *H. litchi* from Vietnam are in bold font.

(1991) to be conspecific with *H. litchi*. Considering the earliest conclusions of Decraemer and Geraert (1992, 1996) and, recently, Van Den Berg et al. (2015), we propose that the population of *H. mangiferae* from Vietnam provided by Eroshenko et al. (1985) should belong to *H. litchi*.

H. cocophilus (Loos, 1949) Chitwood and Birchfield, 1957

Measurements and specimens

After Eroshenko et al. (1985) (population from Vietnam):

Details of female are as follows (n = 10): L = 460 to 500 µm; a = 14 to 17; b = 4.4 to 4.7; c = 13 to 14; V = 89 to 93%; stylet = 49 to 56 µm; R = 101 to 106; Rst = 11 to 12; Rex = 29 to 31; RV = 9 to 10; RVan = 2; Ran = 7 to 8.

After Nguyen (1989) (population from Vietnam, according to Nguyen and Nguyen (2000)):

Details of female are as follows: L=0.37 mm; a=14.6; b=3.6; c=11.1; V=90.4%; stylet=57 μ m.

After Nguyen (1996) (population from Vietnam, according to Nguyen and Nguyen (2000)):

Details of female are as follows (n=5): L=375 to 440 (400.8) µm; a=14.7 to 16.6 (15.3); b=4.3 to 5.0 (4.6); c=15 to 24.5 (19.9); V=91.6 to 95.1 (93.1) %; stylet=49.4 to 50.9 (50.0) µm; VL/VB=0.8 to 1.5 (1.3); R=96 to 107 (103); Rst=14 to 15 (14.6); Roes=23.0; Rex=27 to 28 (27.8); RV=8 to 11 (9.6); RVan=2.0; Ran=6 to 9 (7.6).

Remarks

Morphology of the Vietnamese populations of *H. cocophilus* is highly in accordance with the type population of *H. cocophilus*. Vietnamese population of *H. cocophilus* from Eroshenko et al. (1985) is of equal size, and other populations are of smaller size compared to the type population. These populations can be characterized by the closely fitting sheath; lip region with two annuli, first annulus smaller than second; stylet strong with anchor-shaped knobs; secretory-excretory pore situated 6 to 12 annuli posterior to end of pharynx; vulval flap 1 to 2 annuli long, distinct; anus situated about one annulus posterior to vulva; oval spermatheca with sperms; and tail with a finely rounded tip.

H. microdoratus Dasgupta, Raski & Van Gundy, 1969

Remarks

Nguyen and Nguyen (2000) reported the presence of *H. microdoratus* from a forest in Gia Lai, Vietnam. The female of Vietnamese population of *H. microdoratus* is characterized by having lip region not offset from body contour, bearing two annuli, first annulus larger than second; labial disc slightly elevated, rounded; cuticular annuli 3 to 5 μ m wide at mid-body; stylet knob anchorshaped, 5 to 6 μ m long; dorsal orifice gland located 3 to 5 μ m posterior to stylet base; secretory-excretory pore located 97 to 105 μ m from anterior end; vulva located 11 annuli anterior to tail end, vulvar sheath present; anus located 1 annulus posterior to vulva; and tail tapering to a rounded tail tip.

H. brachyurus (Loos, 1949) Chitwood and Birchfield, 1957

Measurements and specimens

After Germani and Anderson (1991) (population from pepper, Vietnam):

Details of female are as follows (n=9): $L=440\pm4$ (360-460) µm; $a=11\pm0.6$ (9.8-11.6); $b=4.7\pm0.3$ (4.3-5.3); $c=17.3\pm4.06$ (13.7-26.8); $V=92.4\pm1.42$ (90.8-95.5) %; $V'=33.2\pm5.2$ (20-38) µm; $R=97\pm5.4$ (93-105); RSt=13\pm0.9 (12-14); ROe=20\pm1.4 (19-23); Rhem=24±1.5 (21-25); Rex=29±1.6 (27-31); RV=8±0.7 (7-9); Ran=7±1.0 (6-8); Rvan=1; VL/ VB=1.18±0.19 (0.71-1.33); stylet=53±3.5 (47-59) µm.

Remarks

All morphometrics of the Vietnamese population of *H. brachyurus* provided by Germani and Anderson (1991) are totally in agreement with the description of Loos (1949).

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