1	Three new species of nematodes from the syconia of Ficus racemosa in southern India
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Summary – Ficus racemosa with an Indo-Australasian distribution, has so far been recorded to 10 harbour in its fruits, nematode species of the aphelenchoid genera Schistonchus, Ficophagus and 11 Martininema, and species of diplogastrid genera Teratodiplogaster and Pristionchus. The Indian 12 13 species reported so far from *Ficus racemosa* lack comprehensive details on morphology and molecular characterization. In this paper, we describe three new species of nematodes obtained 14 from syconia (enclosed globular infructescence or fruit) of Ficus racemosa found in southern 15 16 India. *Ficophagus glomerata* n. sp. is characterised by small body having b=5.2-9.6, c= 18-23; slightly set-off lip region having well developed cephalic framework; secretory-excretory pore 17 opens near the head; slender stylet with small, rounded/ sloping knobs; ovoid median bulb with 18 relatively posteriorly-placed valve plates; males with sickle-shaped spicules having spatulate or 19 20 hammer-shaped capitulum, represented by an elongate-ovoid condylus and long digitate rostrum 21 and tail conoid with fine, hair-like terminal spike. Teratodiplogaster glomerata n. sp. is 22 characterised by long tubular and narrow stoma with fractal pieces in prostegostom; long rectangular metacorpus; female reproductive system with conspicuous spermatheca and 23 24 amoeboid sperms; males having short, arcuate spicules and keeled gubernaculum; genital papillae in the configuration of P1, P2, C, P3, P4, P5d, (P6d, P7), P8d, Ph and tail conoid with a 25 terminal or subterminal mucro. Pristionchus glomerata n. sp. is characterised by four 26 morphotypes mainly with variations in lip region, stoma, spicules, gubernaculum, and the 27 position of genital papillae. Phylogenetic analyses based upon near-full-length small subunit 28 (SSU) and D2–D3 expansion segments of large subunit (LSU) rRNA genes confirmed affinities 29 with sister species of corresponding genera. 30

Keywords – Fig nematodes, fig wasp, morphology, new species, morphometrics, molecular
 characterization

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35	Nematodes form the most abundant and diverse metazoans in the world and are associated with
36	meso- and macro- faunal arthropods in various associations, ranging from parasitic to mutualistic
37	(Giblin-Davis et al., 2013). These associations have been proposed to lead to speciation and
38	cause increase in nematode diversity and phenotypic plasticity by forming isolated systems
39	(Price, 1980). One such example where such diversification has been observed is the fig-fig
40	wasp-nematode system.
41	The fig-fig wasp system represents a ~75 million-year old obligate relationship (Cruaud et al.,
42	2012) with over 800 known fig species, each associated with a specific pollinator wasp species,
43	although host-switches and pollinator host sharing has been observed in neotropics (Weiblen,
44	2002; Machado et al., 2005). The enclosed fig inflorescence, called a syconium, forms a
45	microcosm containing different species of wasps, bacteria, fungi, mites, and nematodes (Herre et
46	al., 2008). The pollinator wasps are known to be associated with the nematode community
47	comprising of individuals of a single or multiple taxa. The nematodes reported so far from
48	syconia belong to the families, Rhabditidae, Diplogastridae and Aphelenchoididae (Giblin-Davis
49	et al., 2006; Gulcu et al., 2008; Powers et al., 2009; Susoy et al., 2016; Kanzaki et al., 2018)
50	with associations ranging from phoresy to parasitism. Fig nematodes use fig wasps as their
51	vectors in order to move from one fig (microcosm) to another (Krishnan et al., 2010). The
52	nematodes may be transported in the cavity, in intersegmental folds or may cling on to the body
53	surface of the female fig wasp (Giblin-Davis et al., 1995). Parasitodiplogaster,
54	Teratodiplogaster, Pristionchus, Acrostichus, Rhabditolaimus (Diplogastridae);
55	Bursaphelenchus (Aphelenchoididae); Caenorhabditis (Rhabditidae); Schistonchus
56	(Aphelenchoididae); Ficophagus (Aphelenchoididae) and Martininema (Aphelenchoididae) are

57 the genera known to be phoretic on pollinator fig wasps.

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58	HICUS racomosa	the model of	ystem in this study	16.9	monoecious n	lant enectes u	those syconia
50	Ticus rucemosu,	the model s	ystem m uns study	, 15 a	monoccious p	iant species w	mose sycoma,

- 59 harbour a community of nematodes. The nematode species reported from Indian F. racemosa
- 60 include Schistonchus racemosa (Reddy & Rao, 1986), S. osmani (Anand, 2002), S.
- 61 cuculloracemosus (Bajaj & Tomar, 2014), S. flagelloracemosus (Bajaj & Tomar, 2014), S.
- 62 *mucroracemosus* (Bajaj & Tomar, 2014), *Teratodiplogaster racemosus* (Bajaj & Tomar, 2015)
- and *Canalodiplogaster racemosus* (Bajaj & Tomar, 2015); while the species reported from
- 64 Australian F. racemosa include Ficophagus altermacrophylla (Kanzaki et al., 2010) Davies &
- Bartholomaeus, 2015; F. aculeata (Kanzaki et al., 2010) Davies & Bartholomaeus, 2015, F.
- 66 fleckeri (Kanzaki et al., 2010) Davies & Bartholomaeus, 2015 and Teratodiplogaster fignewmani
- 67 (Kanzaki, 2009). The diplogastrid species *Pristionchus racemosa* (Susoy *et al.*, 2016) was
- 68 described from syconia of *F. racemosa* from Vietnam.
- 69 The species reported from India lack critical morphological data for diagnosis and differentiation
- 70 (Davies *et al.*, 2015). All of them also lack molecular characterization which makes future work
- vith re-isolated lines difficult. Thus, there is a need to describe/re-describe the Indian species of
- nematodes associated with fig synconia with modern technologies. The present study deals with
- 73 the morphological characterization of the nematode species isolated from *Ficus racemosa*
- located in southern India along with their molecular characterization to assess their phylogeneticstatus.

76 Materials and Methods

77 Sample collection

- 78 The fig syconia in *Ficus racemosa* pass through five developmental phases: phase A (pre-
- 79 pollination stage, when syconia are at floral bud stage), phase B (pollen-receptive stage, marked
- 80 by entry of pollinators through ostiole, a tiny opening in the syconium), phase C (development

phase of seeds, nematodes and wasps), phase D (wasp dispersal through the opening generated 81 by the male pollinators) and phase E (fruit ripening stage) (Ranganathan et al., 2010). 82 83 Nematodes were collected from the mid-C phase syconia of F. racemosa trees in November 2015 from the campus of the Indian Institute of Science, Bangalore, Karnataka, India at 84 coordinates 13.0219° N, 77.5671° E. 85 86 Isolation and observation Syconia were washed and cut into small pieces using a scalpel. The latex produced by the middle 87 88 layer of the syconium was absorbed using a tissue paper and the pieces of syconia were immersed in sterilized water for 15–20 min in a Petri dish. Nematodes were hand-picked under a 89 90 stereoscopic microscope in sterilized water and later fixed in 4% formalin. The fixed nematodes 91 were then dehydrated using glycerol-alcohol (a mixture of 95 parts of 30% ethanol + 5 parts glycerol) solution (Seinhorst *et al.*, 1959). Nematodes were mounted permanently on slides using 92 93 wax ring technique (De Maeseneer & D' Herde, 1963) for observing under light microscope and conducting confocal imaging using Airyscan LSM 880, Ziess. Drawings were made by tracing 94 the confocal images while measurements were done using Image J 1.46r. 95

96 Scanning Electron Microscopy (SEM)

97 SEM imaging was done to elucidate the surface features of the nematodes. The nematodes were 98 first fixed in 2% glutaraldehyde for 24 h and then post-fixed in 2% osmium tetroxide for 2 h in 99 the dark in a refrigerator. The nematodes were then washed with PBS and transferred into 100 BEEM[®] capsules fitted with nylon mesh of 20 µm pore size to hold the specimens (Bozzola & 101 Russell, 1999). The dehydration was carried out by using a serial gradation of alcohol up to 102 100% and further the nematodes were critical point dried using CO₂. Later, the nematodes were 103 mounted on double-sided carbon tape placed on the stub, sputtered with 10 nm gold for 38 s and

- viewed under SEM JEOL (JSM6510L) at 15 kV at the USIF, Aligarh Muslim University
- 105 (Aligarh).
- 106 *Molecular profiles and phylogeny*
- 107 Nematode DNA was isolated by picking up a single nematode in 5 µl of worm lysis buffer (50
- 108 mM KCl, 10 mM Tris HCL pH 8.3, 2.5 mM MgCl₂, 0.45% NP-40 and 0.45% Tween-20) as per
- the protocol of Williams *et al.* (1992) and keeping it in a water bath at 65°C for 45 min followed
- 110 by 95°C for 15 min. The extracted DNA was stored at -20°C and used as template. Primers
- 111 (Sigma Aldrich) used to amplify D2/D3LSU segment of all three species, were D2A
- 112 (ACAAGTACCGTGGGGAAAGTTG) and D2B (TCGGAAGGAACCAGCTACTA) (Kanzaki
- 113 *et al.*, 2009). Primers used to amplify SSU segments for *F. glomerata* n. sp. were SNF
- 114 (TGGATAACTGTGGTAATTCTAGAGC) (Zeng et al., 2007) and SNR
- 115 (TTACGACTTTTGCCCGGTTC); for *T. glomerata* n. sp. and *P. glomerata* n. sp., these primers
- 116 were SSUF07 (AAAGATTAAGCCATGCATG) and SSU26R
- 117 (CATTCTTGGCAAATGCTTCG) (Hermann et al., 2006, 2007). The PCR products were
- sequenced, and the sequences were deposited in the GenBank database (Accession numbers in
- 119 form of SSU/D2-D3 LSU for *F. glomerata* n. sp. MT903999/ MT903998, for *T. glomerata* n.
- 120 sp. MT904001/ MT904002 and for *P. glomerata* n. sp. MT904000/ MT903997). The sequences
- were aligned by ClustalW and Muscle in MEGA 7 software and were further compared with
- those of other nematode species available at the GenBank database using the BLAST homology
- search program (Zhao et al., 2015 for Ficophagus spp.; Kanzaki et al., 2014 for
- 124 Teratodiplogaster spp. and Susoy et al., 2016 for Pristionchus spp). The best model used to
- generate the Bayesian trees was inferred by using Partition finder ver. 2.1.1. The Akaike-
- 126 supported model, log likelihood (lnL), Akaike information criterion (AIC), the proportion of
- 127 invariable sites, and the gamma distribution shape parameters and substitution rates were also

128	used in the phylogenetic analyses. The inferred model was further used to generate phylogenetic
129	trees using Mr. Bayes software ver. 3.2.6 (Huelsenbeck & Ronquist, 2001) by running the chain
130	for 1,000,000 generations and the 'burnin' was set at 1,000. We used the Markov Chain Monte
131	Carlo (MCMC) method within a Bayesian framework to estimate the posterior probabilities of
132	the phylogenetic trees (Larget & Simon, 1999) using 50% majority rule; the trees were
133	visualized using Fig Tree ver. 1.4.3 (Rambaut, 2006).
134	Results
135	All new species described in this paper have been assigned the specific name "glomerata" since
136	Ficus glomerata is a synonym of Ficus racemosa.
137	Ficophagus glomerata n. sp.
138	(Figs. 1-5)
138 139	(Figs. 1-5) MEASUREMENTS. Table 1, 2.
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139 140 141 142 143	MEASUREMENTS. Table 1, 2. DESCRIPTION <i>Male</i> Body slender, small- to medium-sized, C-shaped when heat-killed, tapering at both ends with maximum width at 2/3 rd of the body length from anterior end. Cuticle with annules < 1 μm wide
139 140 141 142 143 144	MEASUREMENTS. Table 1, 2. DESCRIPTION <i>Male</i> Body slender, small- to medium-sized, C-shaped when heat-killed, tapering at both ends with maximum width at $2/3^{rd}$ of the body length from anterior end. Cuticle with annules < 1 µm wide at mid body. Lateral fields with single thick continuous ridge representing two lines. Lip region
139 140 141 142 143 144 145	MEASUREMENTS. Table 1, 2. DESCRIPTION <i>Male</i> Body slender, small- to medium-sized, C-shaped when heat-killed, tapering at both ends with maximum width at 2/3 rd of the body length from anterior end. Cuticle with annules < 1 μm wide at mid body. Lateral fields with single thick continuous ridge representing two lines. Lip region off-set; cephalic framework moderately sclerotized. Oral aperture surrounded by a sclerotized

separated from isthmus by a constriction; isthmus short, indistinguishable from elongated basal 149 overlap represented by a long dorsal and small ventro-lateral lobe. Secretory-excretory pore 150 151 located in anterior region, 13-16 μ m or ca 1.2-1.5 lip region diam. from anterior end. Pharyngointestinal junction posterior to median bulb in isthmus region. Deirids, hemizonid and phasmids 152 not discernible. Testis usually on left side of intestine; seminal vesicle having amoeboid sperms. 153 154 Spicules slender, strongly arcuate, with hammer-shaped capitulum with a depression comprising 155 an ovoid, spatulate condylus and a long digitate rostrum. Gubernaculum absent. Three pairs of 156 genital papillae in configuration of P1 subventral adcloacal pair and P2 subventral, halfway 157 between cloaca and tail tip and P3 about 3-4 µm anterior to tail terminus. Phasmids located laterally between P1 and P2. Tail conoid, strongly ventrally curved ending into a fine bristle-like 158 spike. Bursa absent. 159

160 *Female*

Body slender, ventrally curved or C-shaped when heat-killed, tapering at both ends with 161 162 maximum diam. at mid-body. Lateral fields with a single ridge as observed under SEM occupying ~2 µm width; labial sensilla papilliform surrounding oral opening; four cephalic 163 papillae present. Cephalic framework moderately sclerotized, Amphids not discernible under 164 165 LM. Stylet moderately built with well-developed, rounded/ sloping knobs; conus approximately 166 60-65% of stylet length. Procorpus slender ($46.4 \mu m$ - $55.6 \mu m$) long; median bulb well-167 developed, ovoid in shape, with valve plates slightly posterior to middle. Pharyngeal glands and pharyngo-intestinal junction similar to those observed in male. Deirids and hemizonid not seen. 168 Excretory pore located in anterior region, ca 1.2-1.5 lip region diam. from anterior end. 169 170 Reproductive system monodelphic-prodelphic. Ovary large, outstretched or with flexure; reflexed part containing oocytes usually in single tier in the proximal region. Oviduct narrow, 171

spermatheca not differentiated; uterus slightly spacious chamber; post-uterine sac 0.4-0.8 times
vulval body diam.; Vulva post-equatorial; vulval lips protruding in some specimens. Anus a
crescent-shaped slit. Tail ventrally arcuate, four-times anal body diam. long with an obtuse or
mucronate tip.

176 TYPE HABITAT AND LOCALITY

Ficophagus glomerata n. sp. was collected from syconia of *Ficus racemosa* in and around the
campus of Indian Institute of Science, Bangalore, Karnataka, India at coordinates 13.0219° N,
77.5671° E.

180 TYPE SPECIMENS

- 181 One holotype male (*F. glomerata* n. sp./1), eight paratype males (*F. glomerata* n. sp./2-8) and
- 182 nine paratype females (F. glomerata n. sp./1-9) of F. glomerata n. sp. on slides were deposited in
- 183 Indian Institute of Science, Bangalore, Karnataka, India. One female paratype (*F. glomerata* n.
- sp./10) and one male paratype (*F. glomerata* n. sp./9) were deposited in the National Nematode
- 185 Collection, Indian Agricultural Research Institute, New Delhi.
- Etymology: The species name "glomerata" derived from *Ficus glomerata*, is a synonym of *Ficus racemosa*.

188 DIAGNOSIS AND RELATIONSHIP

- 189 The new species F. glomerata n. sp. can be characterised by small body having 'b'=5.2-9.6, 'c'=
- 190 18-23; slightly setoff lip region having well developed cephalic framework; slender stylet with
- 191 small, rounded/ sloping knobs; anteriorly located excretory pore; ovoid median bulb with
- 192 relatively posteriorly-placed valve plates; males with sickle-shaped spicules having hammer-

193	shaped capitulum, represented by an ovoid spatula-shaped condylus and long digitate rostrum
194	and tail conoid with fine, hair-like terminal spike and three pairs of genital papillae.
195	Ficophagus glomerata n. sp. closely resembles F. fleckeri (Davies et al., 2013) Davies &
196	Bartholomaeus, 2015 in most morphometric and morphological characteristics but differs in
197	having less lines (2 vs 4) in lateral fields; greater 'b' value in male (6.7-7.8 vs 4.8-6.4); larger
198	stylet (20-23 μ m vs 15-19 μ m); spicules with an ovoid, spatula-shaped condylus, and rostrum
199	digitate and separate (vs prominent condylus with rostrum merging into ventral arm); male tail
200	with 3-4 μ m hair-like spike (vs 2-3 μ m long conical terminal mucro) and female tail with a
201	narrow blunt mucro or terminus (vs acutely pointed terminus in F. fleckeri).
202	The new species closely resembles F. microcarpus Zeng et al., 2011, in most morphometric and
203	morphological characteristics but differs in relatively smaller 'b' (5.2-9.6 vs 8.5-13.0), 'ć' (1.3-
204	1.9 vs 2.6-3.6) and greater 'c' (18-23 vs 12.3-16.6) values in females; relatively posteriorly
205	located excretory pore (13-16 μ m vs 3.5-5.5 μ m) from anterior end; amphids indiscernible (vs
206	prominent); post-uterine sac longer (more than vulval body diam. vs 0.4-0.8 vulval body diam.);
207	males are relatively larger (604-886 μ m vs 400-486 μ m) with sickle-shaped spicules having an
208	ovoid, spatulate (vs small, rounded) condylus and long, digitate (vs narrowly rounded to short
209	digitate) rostrum and cucullus absent (vs present at distal end of spicules in F. microcarpus).
210	Ficophagus glomerata n. sp. differs from F. virens Bartholomaeus et al., 2009 in having lateral
211	fields with 2 (vs 4) lateral lines; females with relatively smaller 'a' (20-28 vs 28-34) and 'c' (18-
212	23 vs 19-41) values, longer stylet (20-23 μ m vs 14-16 μ m); post-uterine sac longer (more than
213	corresponding vulva body diam. vs 0.6-0.7 vulval body diam.); males having greater 'b' (6.7-7.8
214	vs 3-5) value, sickle-shaped spicules having long, digitate (vs short, rounded to digitate) rostrum
215	and tail with hair-like terminal spike (vs without terminal mucro in F. virens).

216	The new species differs from F. altermacrophylla Lloyd & Davies, 1997 in having lateral fields
217	with 2 (vs 3) lateral lines; larger females (604-886 μ m vs 411-571 μ m) with smaller 'ć' (1.3-1.9
218	vs 2.8-3.8), relatively posteriorly located (13-16 μ m vs 5-9 μ m) excretory pore, longer stylet (20-
219	23 μ m vs 14-18 μ m); and relatively anteriorly located vulva (V= 61.2-76.7 vs 70-95); males
220	having greater body diam. (19 -23 μ m vs 15-19 μ m) and 'b' value (5.2-9.6 vs 3-5); posteriorly
221	located (13-16 μ m vs 3-7 μ m) excretory pore; post-uterine sac longer (more than corresponding
222	vulval body diam. vs 0.4-0.8 vulval body diam.); and smaller (14-17 μ m vs 18-24 μ m), sickle-
223	shaped spicules with an ovoid, spatulate condylus and long, digitate rostrum (vs rose-thorn-
224	shaped spicules with knob-like condylus and short, angular rostrum in S. altermacrophylla).
225	Ficophagus glomerata n. sp. was compared with all the species of Ficophagus reported from
226	India. The earlier reported species, S. racemosa, S. osmani isolated from Ficus racemosa were
227	considered species inquirendae (Davies et al., 2015) along with S. hispida (Kumari & Reddy,
228	1984) isolated from Ficus hispida, due to insufficient information particularly on the position of
229	the excretory pore, spicules and position of the caudal papillae. The present species, however,
230	can be well differentiated from S. racemosa, S. osmani and S. hispida on account of
231	morphometric values including shorter (vs larger) post-uterine sac in females; amoeboid (vs
232	flagellate/ rod-shaped) sperms; smaller (vs longer) spicules and three pairs (vs two pairs) of
233	genital papillae.
234	Bajaj and Tomar (2014) published the description of eight species of Schistonchus isolated from
235	three species of Ficus. Six of them appear to belong to genus Ficophagus largely on the position
236	of excretory pore although the differentiating characters of the species are not clear. Only two
237	species S. flagellobenghalensus (Bajaj & Tomar, 2014), S. mucroracemosus with a posterior

238 excretory pore do not fit into *Ficophagus*. In absence of molecular characterization, the status of

the species could not be confirmed due to considerable morphological fluidity and morphometric

overlap. Nevertheless, to verify the status and distinctness of *F. glomerata* n. sp., comparisons
were made with species described from India.

- 242 The new species differs from *F*. (= *Schistonchus*) *mucrobenghalensus* (Bajaj & Tomar, 2014) n.
- comb. in having relatively smaller 'b' (6.7-7.8 vs 7.2-9.1) and 'c' (12-20 vs 21.9-26.7) values;
- 244 dissimilar larger spicules; gubernaculum absent (*vs* faintly visible); genital papillae three pairs

245 (vs four pairs) and tail with hair-like spike/mucro (vs tail devoid of mucro) in males and smaller

246 'b' value (5.2-9.6 vs 7.0-9.2) in females and tail with short, blunt (vs long, pointed terminal)

247 mucro in both sexes and genital papillae in males three pairs [vs four pairs in F.

248 *mucrobenghalensus*]. A disparity in values of stylet length and excretory pore position was

observed between description (page no. 199) and table (3).

250 The new species differs from *F*. (= *Schistonchus*) antherobenghalensus (Bajaj & Tomar, 2014) n.

comb. in having longer stylet (20-22 μ m vs 15 μ m); dissimilar spicules; and tail with hair-like

spike (vs tail devoid of mucro) in male and tail in females having short, blunt (vs long, pointed

terminal) mucro in *F. antherobenghalensus*. A disparity in position of excretory pore as stated in

description (page no. 200) and that given in table (4) was found.

255 *Ficophagus glomerata* n. sp. differs from *F*. (= *Schistonchus*) *flagelloracemosus* in having

smaller 'b' (6.7-7.8 vs 8.1-11.2), 'c' (12-20 vs 21.1-27.5) and greater 'ć' (1.5-2.7 vs 1.2-1.4)

values in males and smaller 'ć' value (1.3-1.9 vs 3.4-5.2) in females; longer stylet (20-22 μm vs

 14μ m); males with larger (14-17 μ m vs 9 -11 μ m) spicules and gubernaculum absent [vs faintly

259 visible in *F. flagelloracemosus* n. comb.]

260 The new species differs from F. (= Schistonchus) cuculloracemosus n. comb.in having smaller

261 'b' (5.2-9.6 vs 10.9-15.1) value in females; stoma with rounded (vs elongated) knobs; excretory

262	pore relatively	posteriorly	located	(13-15	µm vs 7-12	μm) in males;	spicules	smaller ((14-17	μm

- 263 *vs* 20-22 μm), without (*vs* with cucullus) at distal end in *F. cuculloracemosus* n. comb.
- 264 The morphometric differences with the closely related species of *Ficophagus* reported from
- *Ficus racemosa* have been given in the Table 2.
- 266 Molecular phylogenetic relationship
- 267 Partial SSU and D2-D3 of LSU genes were sequenced for *F. glomerata* n. sp. The relative
- 268 placement of *F. glomerata* n. sp. among the other sequenced *Ficophagus* and *Schistonchus*
- species was analyzed. The Bayesian tree (Fig. 4) for SSU was constructed using Ditylenchus
- 270 *halictus* (Zhao *et al.*, 2015) as an outgroup and suggested that: i) *F. glomerata* n. sp. forms a
- sister species with *S. microcarpus* having 97% posterior probability, ii) the genera
- 272 Aphelenchoides Fischer, 1894, Bursaphelenchus Fuchs, 1937, Laimaphelenchus Fuchs, 1937 or
- 273 Schistonchus are not monophyletic which stands in concordance with earlier studies (Zhao et al.,
- 274 2015). The Bayesian tree inferred from D2-D3 of LSU genes (Fig. 5) using *Aphelenchoides*
- 275 *besseyi* (Zhao *et al.*, 2015) as an outgroup, to analyze the relationships of the species in the genus
- 276 *Schistonchus* suggested that: i) all the sequenced *Schistonchus* species are divided into two
- 277 clades with 100% posterior probability support in accordance with earlier work done on the
- 278 phylogeny of Schistonchus (= Ficophagus) species (Zeng et al., 2011, Zhao et al., 2015); ii) F.
- 279 glomerata n. sp. appeared to be closest to an isolate of *Ficophagus* from *Ficus obliqua* in
- Australia, and forms a clade with 74% posterior probability support; this forms a sister group to
- two other clades, one formed by *F. virens* and another one which includes *F. zealandicus*, *F.*
- altermacrophylla and F. benjamina with 84% posterior probability support. Accession numbers
- in form of SSU/D2-D3 LSU for *Ficophagus glomerata* n. sp. are MT903999/ MT903998.
- 284

285	
286	Teratodiplogaster glomerata n. sp.
287	(Figs. 3, 6-8)
288	MEASUREMENTS. Table 3
289	DESCRIPTION
290	Adult
291	Medium-sized species with thin, slender body, strongly curved in posterior region. Cuticle
292	striated, 0.5-1.0 μ m in thickness. Lateral field with lines indiscernible. Lip region laterally
293	flattened and expanded to form large scoop-like structure. Labial sensilla indiscernible whereas
294	cephalic sensilla club-like. (Fig. 3.A). Stoma long and narrow <i>ca</i> 11-13 times longer than wide.
295	Cheilostom cuticularised, long, funnel-shaped, anterior part slightly wider than posterior part.
296	Gymnostom short, tubular, with thickened cuticularised walls, isotopic and isomorphic.
297	Prostegostom shorter, with three fractal pieces in lateral view; mesostegostom represents
298	considerably long part of stegostom; metastegostom slightly compressed chamber with thickened
299	dorsal wall; metastegostomal armature comprising a short, thorn-like dorsal tooth, a medium-
300	sized triangular right subventral tooth and thin left subventral ridge. Telostegostom sclerotised,
301	narrow, tubular, connecting metastegostom to pharynx. Anterior part of pharynx muscular,
302	longer than posterior part, consisting of a muscular cylindrical (procorpus) of 110-148 μ m length
303	and well developed, elongated, 46-85 μ m long median bulb (metacorpus). Nerve ring

304 surrounding posterior part of isthmus at 178-207 μ m from anterior end. Excretory pore located

below nerve ring at 163 -207 μ m from anterior end.

306 *Male*

307 Body strongly arcuate in tail region. Testis single and outstretched; spermatocytes small, arranged irregularly in single or double rows at distal end. Spicule short, thick, slightly arcuate to 308 strongly arcuate consisting of small and a reniform capitulum. Gubernaculum slender, bow-309 shaped to prominently keeled; weakly arcuate, tapering to blunt proximally, broad and keeled in 310 the middle and finely attenuated at distal end. Eight pairs of genital papillae present with 311 312 configuration of P1 (3.5-4 cloacal body diam. anterior to cloacal opening), P2 (1.5 cloacal body diam. anterior to cloacal opening) and P3 (just adjacent to cloacal lip), P4/P5 (located at one and 313 a half cloacal body diam. posterior to spicules or about 1/3rd of distance from spicules to tail tip, 314 315 P6-7 form a group with a rudimentary bursal membrane around, P8d (located subdorsally near tail tip). Phasmids located after P8d. Tail broad and strongly arcuate ventrally. Tail tip rounded 316 with a small mucro present terminally or subterminally. 317

318 *Female*

Reproductive system didelphic, amphidelphic; anterior ovary on the right and posterior on left of 319 320 the intestine. Both genital branches equally developed. Each ovary reflexed at its total length, 321 oocytes arranged in several rows in distal half and shift to a single file proximally. Oocytes show granular texture; oviducts slender with posterior part serving as spermatheca, containing 322 amoeboid sperms; uterus with thick walls, occasionally containing one or two egg(s). Vagina 323 perpendicular to body axis; four small vaginal glands present, observed in lateral view; vulval 324 325 lips protruding, without vulval flap. Anus a dome-shaped slit, not protuberant. Tail broad, long, weakly tapering to a rounded terminus with a small blunt mucron. 326

327 TYPE HABITAT AND LOCALITY

328	Teratodiplogaster glomerata n. sp. was collected from Ficus racemosa host trees situated in and
329	around the campus of Indian Institute of Science, Bangalore, Karnataka, India at coordinates
330	13.0219° N, 77.5671° E.

331 TYPE SPECIMENS

One holotype male (*T. glomerata* n. sp./1), eight paratype males (*T. glomerata* n. sp./2-8) and

nine paratype females of *T. glomerata* n. sp. on slides were deposited in Indian Institute of

Science, Bangalore, Karnataka, India. One female paratype (*T. glomerata* n. sp./10) and one

male paratype (*T. glomerata* n. sp./9) were deposited in the National Nematode Collection,

336 Indian Agricultural Research Institute, New Delhi.

337

Etymology: The species name "glomerata" derived from *Ficus glomerata*, is a synonym of *Ficus racemosa*.

340 DIAGNOSIS AND RELATIONSHIP

Teratodiplogaster glomerata n. sp. is characterized by the presence of relatively narrow and 341 342 elevated, laterally compressed lip region with fused lips, long tubular and narrow stoma with fractal pieces in prostegostom; long rectangular metacorpus; almost cylindroid to slightly 343 expanded basal bulb; female reproductive system with conspicuous spermatheca and amoeboid 344 sperms; males having short, arcuate spicules with tapering distal ends and curved and keeled 345 gubernaculum; two pairs of precloacal genital papillae (P1, P2, C, P3, P4, P5d, (P6, P7), P8d, 346 Ph) and tail conoid with a terminal or subterminal mucro. Comparison has been made with 347 three species of Teratodiplogaster reported so far from Ficus syconia viz., T. fignewmani 348 (Kanzaki et al., 2009) isolated form Ficus racemosa (Australia), T. martini isolated from Ficus 349

- *sycomorus* (Africa) (Kanzaki *et al.*, 2012) and *T. variegatae* isolated from *Ficus variegata*(Japan) (Kanzaki *et al.*, 2014).
- 352 *Teratodiplogaster glomerata* n. sp. differs from *T. fignewmani* in having smaller males (1195-
- $1345 \ \mu m \ vs \ 1850-2700 \ \mu m$) and females (1388-1657 \ $\mu m \ vs \ 2160-3895 \ \mu m$); females with
- smaller 'b' value (7.3-8.2 *vs* 9.5-13.5); stoma with (*vs* without) fractal dot-like arcade syncytia;
- amoeboid (vs lemon-shaped) spermatids; males having smaller 'c' (10.6-12.8 vs 16.9-26.6) and
- 356 greater 'ć' (6.2-8.5 *vs* 3.4-5.7) values; shorter (17-20 *vs* 25-32 μm), arcuate (*vs* stout, fusiform to
- bow-shaped) spicules; keeled (vs trough-shaped) gubernaculum and two pairs [vs three pairs of
- 358 precloacal papillae in *T. fignewmani apud* Kanzaki *et al.*(2009)].
- 359 The new species differs from *T. martini* Kanzaki *et al.*, 2012 in having smaller males (1195-1345
- $vs 1825-2650 \,\mu\text{m}$) and females (1388-1657 vs 1750-3100 μm); females with smaller 'c' (8.3-
- 361 13.0 *vs* 15.9-26.3) and greater 'ć' (7.7-13.5 *vs* 4.8-7.3) values; relatively smaller metacorpus;
- 362 males having smaller 'c' (10.6-12.8 *vs* 18.1-25.0) and greater 'ć' (6.2 -8.5 *vs* 3.5-4.9) values;
- relatively shorter (17-20 vs 18-28 μm) spicules and keeled (vs L-shaped) gubernaculum and
- 364 genital papillae having two precloacal pairs (vs three precloacal pairs) and with dissimilar
- 365 configuration [P1, P2, C, P3, P4, P5d, (P6, P7) P8d, Ph vs P1, P2, P3, C, P4d, P5 (P6, P7), P8d,
- 366 Ph in *T. martini apud* Kanzaki *et al.* (2012)].
- 367 The new species differs from *T. variegatae* Kanzaki *et al.*, 2014 in having larger females (1388-
- $1657 \,\mu\text{m} vs \, 805{-}1173 \,\mu\text{m}$) having greater 'a' (68.8-79.7 vs 26.1-45.6), 'b' (7.3-8.2 vs 3.8-4.9)
- and 'ć' (7.7-13.5 vs 3.2-3.8) values; smaller 'c' (8.3-13.0 vs 17.0-19.9); lip region elevated (vs
- 370 flattened and low); metacorpus demarcated (vs indistinguishable); males having greater 'a'
- 371 (62.0-69.6 vs 29.9-47.4), 'b' (5.9-6.3 vs 3.8-5.4) and 'c' (6.2 -8.5 vs 3.5-4.9) values; smaller 'c'
- (10.6-12.8 vs 18.1-25.0); relatively shorter $(17-20 \mu m vs 17-30 \mu m)$ spicules with simple, arcuate

373	vs keel-like dorsal	part): gen	tal papillae	with dissimilar	configuration	(P1. P2.	C. P3	. P4.	P5d.

374 (P6, P7) P8d, Ph vs P1, P2, P3, C, (P4, P5d), (P6, P7), P8d, Ph and tail terminus with simple [vs

star-shaped mucro in *T. variegatae apud* Kanzaki *et al.*(2014)].

- The new species differs from *T. racemosus* Bajaj & Tomar (2015) in having smaller males
- 377 (1195-1345 μ m vs 1412-2184 μ m) having relatively smaller stoma (19-21 μ m vs 20-25 μ m);
- smaller (17-20 vs 19-25 μm) spicules; proximally attenuated (vs blunt) gubernaculum; genital
- papillae eight pairs (vs seven pairs) with dissimilar configuration (P1, P2, C, P3, P4, P5d, (P6,

280 P7) P8d, Ph *vs* P1, P2, C, P3, P4, Ph, (P5, P6), P7d; relatively smaller females (1388-1657 μm *vs*

- 381 1587-2444 μ m) with smaller 'c' value (10.6-12.8 vs 12.9-17.8); smaller (108-189 μ m vs 177-298
- μm) tail, and tail usually with subterminal (vs terminal mucro in T. racemosus apud Bajaj &

383 Tomar, 2015).

384 The new species differs from *Ceratosolenus* (=*Rhabditolaimus*) *racemosa* (Anand, 2005) having

presence of genital papillae (*vs* absence) and bow-shaped gubernaculum (*vs* rod-shaped). The

species reported from India that include species belonging to both genus *Rhabditolaimus* and

387 *Teratodiplogaster* have insufficient description, poor illustration and lack molecular

388 characterization hence making it difficult to validate these as separate species. The morphometric

differences within all the *Teratodiplogaster* species known so far, have been shown in Table 1.

390

391 Molecular phylogenetic relationship

Partial SSU and D2-D3 of LSU genes were sequenced for *T. glomerata* n. sp. The relative

- 393 placement of *T. glomerata* n. sp. among the other sequenced *Teratodiplogaster* and
- 394 Paradiplogaster species was analyzed. The Bayesian tree (Fig. 8) constructed using Koerneria

395	luziae (Kanzaki et al., 2014) as an outgroup suggested that: i) Parasitodiplogaster forms a
396	monophyletic clade in relation to the Teratodiplogaster clade, ii) In the Teratodiplogaster clade,
397	the Teratodiplogaster species collected from Ficus racemosa shows a monophyletic relation
398	with T. variegatae, and iii) T. glomerata n. sp. forms a sister species of T. fignewmani and also
399	shows a monophyletic relationship. Accession numbers in form of SSU/D2-D3 LSU for T.
400	<i>glomerata</i> n. sp. are MT904001/ MT904002.
401	Pristionchus glomerata n. sp.
402	(Figs. 9-15)
403	MEASUREMENTS. Table 4
404	DESCRIPTION
405	Adult
406	Body stout, length ranging from 1-1.5 mm. Cuticle thick, transversely annulated without
407	punctations or longitudinal striations. Lateral field showing presence of a single ridge. Stomal
408	morphology variable in five morphotypes of the species. Anterior part of pharynx (= pro- and
409	metacorpus) 1.5 times as long as posterior part (isthmus and basal bulb). Procorpus muscular,
410	stout, 105.8 \pm 7.3 μm long; metacorpus very muscular representing rectangular or ovoid median
411	bulb of $43.8\pm5.6\mu m$ dimension; is thmus narrow and not muscular; basal bulb small, glandular.
412	Pharyngo-intestinal junction conspicuous. Nerve ring encircling middle or anterior to middle
413	region of isthmus. Excretory pore faintly visible, with position ranging from anterior level of
414	basal bulb to pharyngo-intestinal junction. Hemizonid and deirids obscure. Five different morphs
415	could be observed out of which the pharynx of two morphotypes (α , γ) possessed "fish-bone" or
416	zipper-like lumen; pharyngeal lumen of morph ε is smooth. Pharynx comprising an anterior

417 corpus continuing into a swollen metacorpus which is followed by a narrow isthmus terminating418 into an elongate expanded basal part.

419 *Male* (general morphology comprising γ , δ , ε)

Body straight to ventrally arcuate except strongly ventrally curved posterior region. Testis single, 420 ventrally located at 1/3rd of genital branch from anterior end. Spermatocytes arranged in 2–5 421 422 rows followed by amoeboid spermatids arranged in multiple rows in genital tract. Vas deferens not clearly separated from other parts of gonad. Spicules free, bow-shaped having a wide, 423 424 bilobed capitulum and pointed distal end. Gubernaculum conspicuous, flared anteriorly with ventral and dorsal walls separated at an angle of 45° with a prominent, curved dorsal wall. Tail 425 426 conical, tapering down abruptly, with long spike. Nine pairs of genital papillae arranged in 427 configuration: P1, P2d, P3, C/P4, P5, Ph (P6, P7, P8), P9d. P1 located at 1.5 cloacal body diam. anterior to cloaca, P2 was located at 1 cloacal body diam. anterior to cloaca, P3 present above the 428 cloaca, P4 just below the cloacal opening, P5 small 4–5 µm anterior to P6 that form a group with 429 430 P7 and P8, P9 posterior to P8 directed dorsally. Phasmid located between P5 and P6. Tail spike 431 about 2.5-3.0 cloacal body diam. long.

432 *Female* (morph α)

Body slightly arcuate when heat killed. Gonad didelphic, ovaries reversed. Anterior gonad right
of intestine while posterior one on left side of intestine. Oocytes mostly arranged in multiple
rows. Receptaculum seminis not observed; posterior part of oviduct serving as spermatheca
holding small amoeboid sperm cells. Uterus spacious with 2–4 intra-uterine eggs. Vagina
perpendicular to body surface; vulva elliptical slit-like with protruding lips. Posterior anal lip
prolapsed. Tail long, conical to filiform. Tail spike about 2.5–3 cloacal body diam. long.

439

440 TYPE HABITAT AND LOCALITY

The type specimen was collected from *Ficus racemosa* host trees situated in and around the
campus of Indian Institute of Science, Bangalore, Karnataka, India (GPS: (GPS: 13.0219° N,
77.5671° E).

444 Morphs

Of the five morphotypes found, α morphs are represented by females while γ , δ , ε are represented by males. The morphs mainly show phenoplastic traits in the anterior region especially the labial region, stoma and pharyngeal lumen. However, the general characters of females remain the same while males also show similar spicules and gubernaculum morphologies. Morphological illustrations and photographs are shown in Figs. 9 (A), 10-15.

450

Morph α (Females): Lip region wide, slightly offset with six equal-sized, rounded lips; 451 452 cheilostom equal or slightly longer than gymnostom, absence of tightly packed cheilostomal 453 rugae, cheilostomal flap weak; gymnostom with prominent serrated anterior margin; promesostegostom without lobes; metastegostom with claw-like dorsal tooth, curved right 454 subventral tooth, and coarsely serrated left subventral ridge overlapping $2/3^{rd}$ of the dorsal tooth; 455 telostegostom long, sclerotized with straight dorsal and wide tapering subventral walls. 456 457 Pharyngeal lumen zipper-like. Conspicuously larger phasmids present in anterior half of the tail, 458 not found in any other morph.

459

460	Morph γ (Male): Lip region narrow; lips distinct; six equal-sized, amalgamated, stoma funnel-
461	shaped, wider anteriorly and tapering at base; cheilostom thick; gymnostom relatively thin,
462	smooth, narrower; metastegostom with a large flap-like dorsal tooth, a triangular right
463	subventral tooth and left subventral ridge; stegostom simple, smooth, continuing into zipper-like
464	(fish bone-like) pharyngeal lumen.
465	
466	Morph δ (Male): Lip region with six equal-sized, amalgamated lips; cheilostom thick with
467	sloping walls, arched; gymnostom half the length of the cheliostom, thickened, arched;
468	metastegostom with a triangular dorsal tooth and right subventral, dagger-like tooth and a left
469	subventral plate; telostegostom shallow, short, weakly sclerotized.
470	
471	Morph ϵ (Male): Anterior labial region with umbrella-like flap having six labial projection; left
472	side possessing a slit with a round opening at base, the flap in few specimens covers the
473	gymnostom and cheilostom of one side and with six ribs consisting of labial papillae within the
474	cuticle; cheilostom smooth, anterior margin marked by per- and interradial notches; cheilostom
475	and gymnostom equal in length and project outward, metastegostom simple, smooth, except for a
476	flat, thin, claw-like dorsal tooth, ventral tooth absent; telostegostom weakly sclerotized,
477	indistinct.
478	TYPE HABITAT AND LOCALITY

479 *Pristionchus glomerata* n. sp. was collected from *Ficus racemosa* host trees situated in and
480 around the campus of Indian Institute of Science, Bangalore, Karnataka, India at coordinates
481 13.0219° N, 77.5671° E.

482 TYPE SPECIMENS

483	One holotype male ε morph (<i>P. glomerata</i> n. sp./1), eight paratype males (<i>P. glomerata</i> n. sp./2-
484	8) and nine paratype females (P. glomerata n. sp./1-9) of P. glomerata n. sp. on slides were
485	deposited in Indian Institute of Science, Bangalore, Karnataka, India. Holotype male (ϵ
486	morphotype) (P. glomerata n. sp./9) was deposited in the National Nematode Collection, Indian
487	Agricultural Research Institute, New Delhi.

488

489 Etymology: The species name "glomerata" derived from *Ficus glomerata*, is a synonym of *Ficus*490 *racemosa*.

491 DIAGNOSIS AND RELATIONSHIP

492 Pristionchus glomerata n. sp. is characterized by its phoretic relationship with Ceratosolen 493 *fusciceps*, the pollinator wasp of the *Ficus racemosa*, and presented four different morphotypes *viz.*, α , γ , δ and ε types. The morphs mainly show variations in labial region ranging from 494 495 rounded, truncate to umbrella-shaped, stoma from barrel-shaped to cuboidal or elongated, dorsal tooth being claw-shaped to triangular, dagger-like and pharyngeal lumen simple to zipper-like; 496 497 female reproductive system didelphic, ovaries reversed; oocytes mostly arranged in multiple 498 rows; proximally dilated oviduct holding amoeboid sperm cells; 2–4 intra-uterine eggs 499 occasionally present; vulva elliptical slit-like with protruding lips; males having a bow-shaped 500 spicules with a wide, bilobed capitulum; gubernaculum conspicuous, with 45° bent, curved 501 dorsal wall. Tail conical, tapering down abruptly, with long spike. Nine pairs of genital papillae 502 arranged in configuration P1, P2/ P2d, P3, C/P4, P5, Ph, (P6, P7, P8), P9d. The morphotype a possessed exceptionally large phasmids. 503

The new species is distinguished from all other Diplogastridae, except *P. borbonicus* Susoy *et al.*, 2016, *P. sycomori* Susoy *et al.*, 2016 and *P. racemosae* Susoy *et al.*, 2016 by the presence of

506 morphs of laterally symmetrical and laterally asymmetrical stomatal structures. The diagnostic characteristic features of this species are the gross morphological differences in the morphs 507 508 compared with those of the species reported from fig species. Among the fig-associated Pristionchus species, P. glomerata n. sp. comes close to P. racemosae in having the basic 509 structural similarity in the stomal components of four morphs; however, relatively cylindrical (vs 510 511 globular) buccal cavity in γ and δ morphotypes and relatively rectangular and less expanded 512 (funnel-shaped and well expanded) lip region in ε morphotype in *P. racemosa*e and lacks β 513 morph.

514 P. glomerata n. sp. differs from P. sycomori in having lips moderately developed (vs large), equal-sized (vs lateral lips higher and wider than subventral and subdorsal lips); labial region 515 with ring of conspicuous, thin cuticular cheilostomal filaments); cheilostom (equal or smaller vs 516 517 twice the length of gymnostom) with less developed cheilostomal rugae (vs thick, tightly packed 518 rugae); gymnostom without (vs with) heavy punctations, lip region slightly narrower (vs 519 conspicuously wider) than the adjoining body; anterior labial margin smooth (vs projected, 520 "wave"-like) in α morphotype and larger, relatively expanded and umbrella-like (vs less expanded, barrel-shaped) lip region in ε morphotype in *P. sycomori apud* Susoy *et al.* (2016) 521 522 (Fig. 15).

Pristionchus glomerata n. sp. differs from *P. borbonicus* in having lips moderately developed (*vs* large), equal-sized (*vs* lateral lips higher and wider than subventral and subdorsal lips); labial region with ring of conspicuous, thin cuticular cheilostomal filaments); cheilostom (equal or smaller *vs* twice the length of gymnostom) with less developed cheilostomal rugae (*vs* thick, tightly packed rugae); gymnostom without (*vs* with) heavy punctations, lip region slightly narrower (*vs* conspicuously wider) than the adjoining body; anterior labial margin smooth (*vs*

projected, "wave"-like) in α morphotype and larger, relatively expanded and umbrella-like (*vs*less expanded, barrel-shaped) lip region in ε morph in *P. sycomori apud* Susoy *et al.* (2016). *Pristionchus glomerata* n. sp. differs from *Canalodiplogaster racemosus* (Bajaj & Tomar 2015)
in having 9 pairs of genital papillae (*vs* 7 pairs), conspicuous gubernaculum (*vs* large pouch
shaped) and spicule monomorphic (*vs* dimorphic).

534 REMARKS

Bajaj and Tomar (2015) reported a new genus Canalodiplogaster with C. racemosus as its 535 536 species from the fig F. racemosa along with three other new genera. The description and 537 illustrations of the former indicated it to be a species of *Pristionchus* with three morphotypes. 538 The authors have identified the morphs as stenostomous and eurystomous individuals. The 539 present species *P. glomerata* n. sp. with five morphs can be distinctly differentiated from the species (= C. racemosus) in having γ morph with funnel-shaped (vs barrel-shaped) buccal cavity; 540 smaller (20-22 μ m vs 29-43 μ m) spicules and morphotype ε with umbrella-like lip region 541 represented by only males (vs females in the species (=C. racemosus) apud Bajaj and Tomar 542 (2015). The status of the latter species can be verified by revisiting the species for the molecular 543 data and for detailed information about all the morphotypes associated with the species. 544

545

546 Molecular phylogenetic relationship

547 Partial SSU and D2-D3 of LSU genes were sequenced for *P. glomerata* n. sp. The relative

548 placement of *P. glomerata* n. sp. among the other known sequenced *Pristionchus* species was

analyzed. The Bayesian tree (Fig. 16) constructed using Koerneria sp. (Kanzaki et al., 2014),

550 Parapristionchus giblindavisi and Micoletzkya masseyi (Susoy et al., 2016) as an outgroup,

suggested that: i) The fig-associated *Pristionchus* form a polyphyletic clade in relation to *P*.

552 *bucculentus, P. elegans* and *P. fissidentatus*, ii) In the fig-associated *Pristionchus* clade,

553 Pristionchus collected from Ficus racemosa shows a monophyletic relation to other fig-

associated *Pristionchus* species, and iii) *P. glomerata* n. sp. is a sister species of *P. racemosae*

and also shows a monophyletic relationship. Accession numbers in form of SSU/D2-D3 LSU for

556 *Pristionchus glomerata* n. sp. are MT904000/ MT903997.

557

558 Discussion

559 Nematodes have long been associated with the fig syconium as a substratum for their growth and reproduction whereas the pollinators associated with figs are used as vehicles for their transport 560 561 from one syconium to the other (Krishnan et al., 2010). These nematodes in association with the 562 fig syconium have shown millions of years of co-evolution, co-speciation and co-diversification (Herre, 1993; Davies et al., 2015). Such associations might range from being commensal to 563 parasitic in nature (Giblin-Davis et al., 2013). The fig-associated nematodes of families 564 Aphelenchoididae and Diplogastridae have shown independent phylogenetic radiation (Kanzaki 565 et al., 2009; Davies et al., 2015). The former family constitutes plant-parasitic nematodes that 566 567 feed on the anthers and the epidermis of the female florets of the fig (DeCrappeo & Giblin-Davis, 2001; Vovlas et al., 1992; 1996; 1998; Giblin-Davis et al., 1995, 2006; Center et al., 568 1999) whereas the latter family includes nematodes which can be fungal feeders, bacteriovores 569 570 or insect parasites (Susoy et al., 2016). *Ficus racemosa*, the fig plant under study in this paper, is a monoecious species of subgenus 571 572 Sycomorus commonly found in Bangalore, South India. A single pollinator wasp species

573 *Ceratosolen fusciceps* and six non-pollinator fig wasp species (Ghara & Borges, 2010) have been

574 found associated with this fig species, of which the former serves a reliable vehicle for nematode

575	transport (Krishnan et al., 2010, Gupta & Borges, 2019). So far, the nematode species reported
576	from Ficus racemosa belong to genera Schistonchus, Ficophagus, Teratodiplogaster and
577	Pristionchus (=Canalodiplogaster) (Reddy & Rao, 1986; Anand, 2002; Anand, 2005; Bajaj &
578	Tomar, 2014, 2015). However, in the present study we could find only three species associated
579	with Ficus racemosa, each a representative of Ficophagus, Teratodiplogaster and Pristionchus.
580	Coincidentally, all the species described from India lack molecular characterization and are
581	differentiated mainly on minor morphological characters without any report of phenotypic
582	plasticity thus causing difficulty in ascertaining their correct status in comparison with other
583	species (Anand, 2002; Anand, 2005; Reddy & Rao, 1986; Davies et al., 2013; 2015).
584	Ficophagus glomerata n. sp. forms a sister species with F. microcarpus isolated from Ficus
585	microcarpa in China under the phylogenetic analysis done using partial SSU region and
586	Ficophagus sp. isolated from Ficus obliqua using the D2–D3 segment of LSU. It was found to
587	be distantly related to the species found in Australia suggesting that the species might have
588	diverged out separately either because of geographical range variation or due to host shift. The
589	phylogenetic trees stand in concordance with the earlier study where it was proposed that the
590	genera Schistonchus and Ficophagus represented two evolutionary lines of the polyphyletic
591	clade (Davies et al., 2010).
592	The diplogastrid species, T. glomerata n. sp. shows affinities with T. fignewmani and T.
593	racemosus. The presence of T. glomerata n. sp. in Ficus racemosa syconia suggests that a

speciation event might have occurred due to large geographical ranges and genetic isolation of F.

- 595 *racemosa* in south India compared to the south-east Asian populations (Bain *et al.*, 2016). The
- 596 phylogenetic analysis done using SSU and LSU shows that *Teratodiplogaster* forms a
- 597 monophyletic clade with respect to the *Pristionchus* clade. The new species *T. glomerata* n. sp.
- 598 forms a sister species to *T. fignewmani*. The data still stands in concordance with the earlier

study suggesting the *Parasitodiplogaster* Poinar, 1979, and *Teratodiplogaster* Kanzaki *et al.*2009 clades to be monophyletic and separated into five groups, i.e. *Parasitodiplogaster australis*Bartholomaeus, *et al.* 2009, *Parasitodiplogaster sycophilon* Poinar 1979, *Parasitodiplogaster maxinema* Poinar & Herre, 1991, *Parasitodiplogaster laevigata* Giblin-Davis *et al.* 2006 and *Parasitodiplogaster citrinema* Poinar & Herre, 1991, and the *Teratodiplogaster* Kanzaki *et al.*2009 clade (Wöhr *et al.*, 2014).

Pristionchus, another genus of Diplogastridae, is known to be mainly associated with beetles but 605 606 recently has been reported from Ficus species (Herrmann et al., 2006, 2007; Kanzaki et al., 607 2011, 2012a, 2012b). The species found so far are *P. borbonicus* collected at Grand Étang, La Réunion Island, from F. mauritiana; P. sycomori collected in South Africa from F. sycomorus; 608 P. racemosae collected in Vietnam from F. racemosa (Susoy et al., 2016) and a species of 609 Pristionchus reported (Bajaj & Tomar, 2015) from India with the name Canalodiplogaster 610 611 racemosus. According to the phylogenetic analysis, *Pristionchus glomerata* n. sp. forms a 612 closely related sister species to *P. racemosae* where the branch length increase in the arm of Pristionchus glomerata n. sp. depicts speciation with recent divergence. The species present in 613 *Ficus racemosa* forms a monophyletic clade to the other *Pristionchus* species which are 614 615 associated with figs. The close association is suggestive of speciation which might have occurred due to large geographical range variations. The occurrence of five morphotypes is an ideal case 616 617 of character displacement for the efficient sharing of resources in the microhabitat of syconium populated by a good number of species. The morphs (δ , ε) with simple, tube-like buccal cavities 618 619 devoid of effective armature seem to compete for a temporary bacterial food source whereas the predatory morphs (α , γ) emerge next coinciding with the proliferation of wasp-transmitted 620 nematodes in the syconium. Development of greater number of morphs might be due to the high 621 climatic and seasonal variation which is observed in southern India (Gadgil & Joshi, 1983; 622

Gunnel 1997; Peel *et al.*, 2007; Shenoy & Borges, 2010; Chanam *et al.*, 2014). We observed sex specific stomatal morphs (β = adult female only) versus (γ , δ , ε = adult males only) which requires further validation through a study similar to that reported in Susoy *et al.* (2016) with *Pristionchus sycomori* where different early and late interfloral (phase C) figs were examined and large numbers of *Pristionchus* were sexed and assessed for stomatal morphotype. This will be attempted in the future.

The phylogenetic analyses of these nematodes using LSU and SSU of rRNA sequences have 629 630 helped in the better understanding of nematode diversification. There are instances in which a single *Ficus* species is known to be associated with several nematode species belonging to the 631 same genus, for example, several species of Schistonchus associated with Ficus racemosa in 632 Australia (Davies *et al.*, 2010), which is suggestive of opportunities for host switching for such 633 cosmopolitan fig species (Zhang et al., 2006). Future studies on nematode taxonomy related to 634 *Ficus* species should involve sequencing of at least two genes which might help to better resolve 635 636 the phylogenetic trees of these nematode species. Special emphasis should be given to comparative genomic analyses using the new species and its close relatives which would yield 637 interesting information concerning the genes involved in different life history traits, switching 638 639 feeding habitats and the development of plant and animal parasitism. Further investigation on *Pristionchus* species associated with *Ficus* species might help us to conduct polymorphism 640 641 studies across the several geographical landscapes to understand the relationships between form 642 and function and how evolution may have proceeded from a common ancestor. This might even lead to better understanding of life history adaptations used by these nematode species. This 643 644 study also paves the way for further research on fig nematodes related to their biodiversity, 645 distribution, evolutionary history, host/carrier relationships with pollinators associated with the 646 *Ficus* systems and co-speciation between fig and fig associated nematodes.

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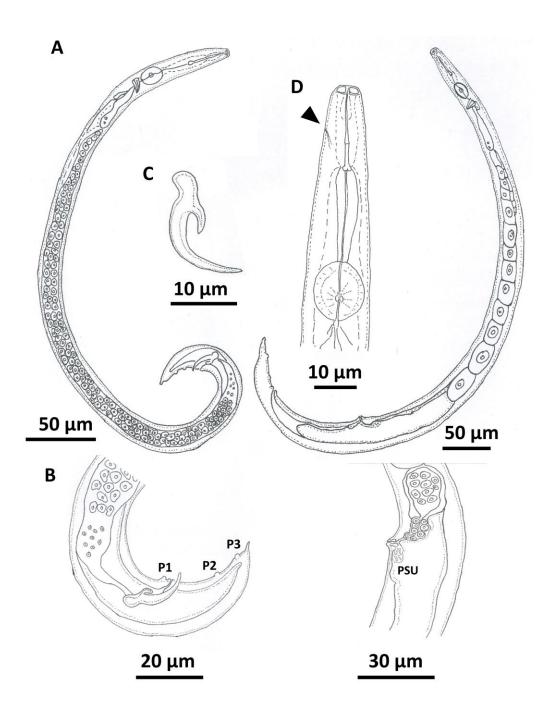


Fig. 1. *Ficophagus glomerata* n. sp. lateral view Male A: Habitus, B: Tail region, C: Spicule, D: Anterior region (similar in case of female); Female E: Habitus and F: Vulval region.

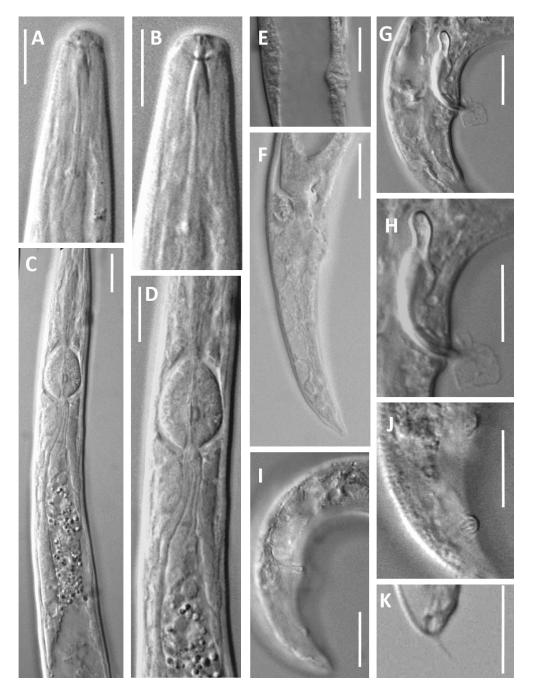


Fig. 2. Photomicrographs of *Ficophagus glomerata* n. sp., anterior end of a male showing a stylet and basal knob (A - B) and a distinct median bulb and pharyngeal glands (C - D). Vulval region of female showing vulval opening (E), and tail region of the male showing anal opening (F), spicule (G - H), genital papillae (I - J) and spicate tip (K). Scale bar: (A - K) 10 µm.

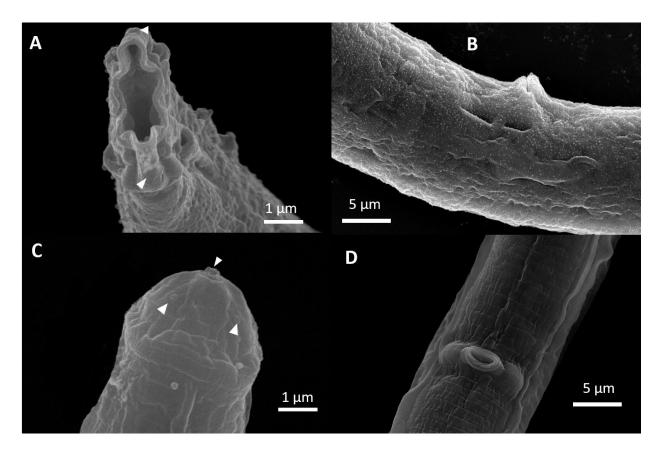


Fig. 3. SEM observations: A: Anterior end; B: Vulval region of female of *Teratodiplogaster glomerata* n. sp.; C: Anterior end; D: Vulval region of female *Ficophagus glomerata* n. sp. (arrows indicate points to the amphidal pore and cephalic sensilla).

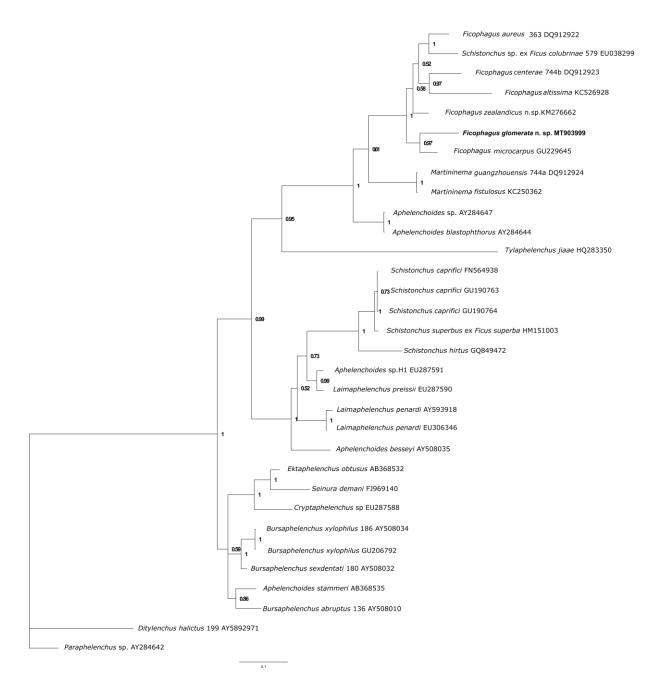


Fig. 4. The Bayesian tree inferred from the 18S gene for *Ficophagus glomerata* n. sp. under the GTR + I + G model (lnL = 5957.9728; freqA = 0.2547; freqC = 0.1966; freqG = 0.2654; freqT = 0.2833; R (a) = 0.0959; R (b) = 0.2182; R (c) = 0.1305; R (d) = 0.0852; R (e) = 0.4025; R (f) = 0.0676; Pinva = 0.27; Shape = 0.593). The accession numbers of the compared sequences are indicated in the form: SSU. Posterior probability values exceeding 50% are given on appropriate clades.

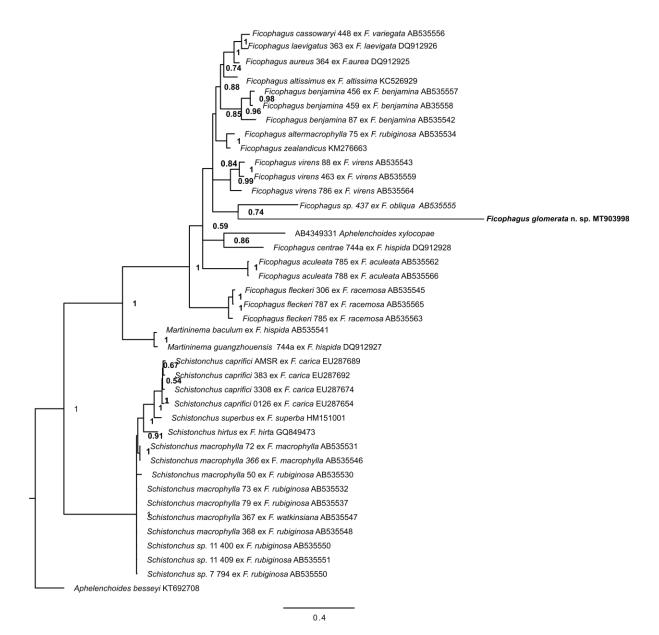


Fig. 5. The Bayesian tree inferred from the 28S gene for *Ficophagus glomerata* n. sp. under the GTR + I + G model (lnL = 9439.3496; freqA = 0.2437; freqC = 0.1807; freqG = 0.2957; freqT = 0.28; R (a) = 0.0592; R (b) = 0.2708; R (c) = 0.1709; R (d) = 0.0879; R (e) = 0.397; R (f) = 0.0772; Pinva = 0.0117; Shape = 0.7021). The accession numbers of the compared sequences are indicated in the form: D2-D3 LSU. Posterior probability values exceeding 50% are given on appropriate clades.

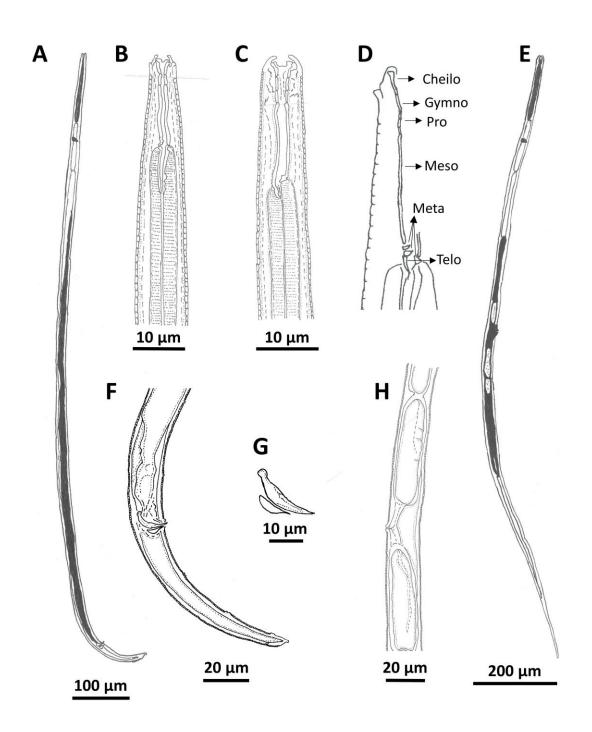


Fig. 6. *Teratodiplogatser glomerata* n. sp. lateral view (A) Habitus of male, (B) Anterior region of male, (C) Anterior region of female, (D) Schematic representation of lateral view of stomatal morphology, (E) Habitus of male, (F) Tail region of male, (G) Spicule and gubernaculum and (H) Vulval region of female.

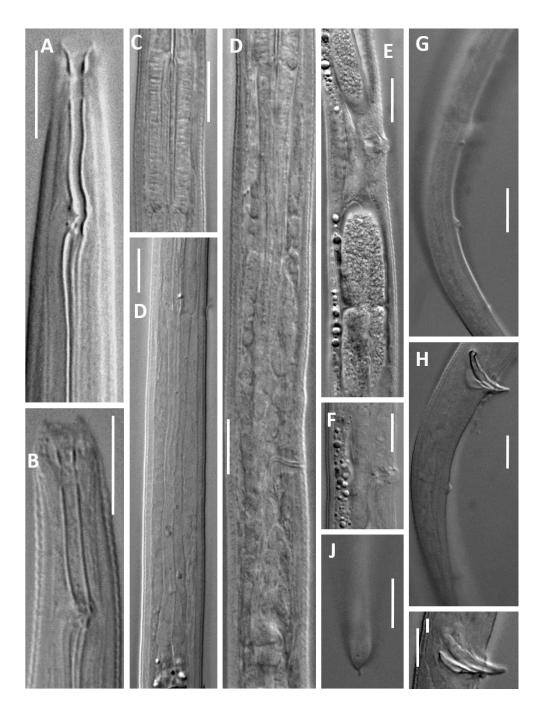


Fig. 7. Photomicrographs of *Teratodiplogaster glomerata* n. sp. (A, B) Head morphology of a male and female respectively. Anterior and posterior pharynx (C – D), (E) Female gonadal region showing vulva region showing the vulval opening and the eggs, (F) vulval opening, (G – H) Tail region of a male at different focal planes showing genital papillae, (I) the male spicule, and (J) the mucronated tip. Scale bar: $(A – J) 10 \mu m$.

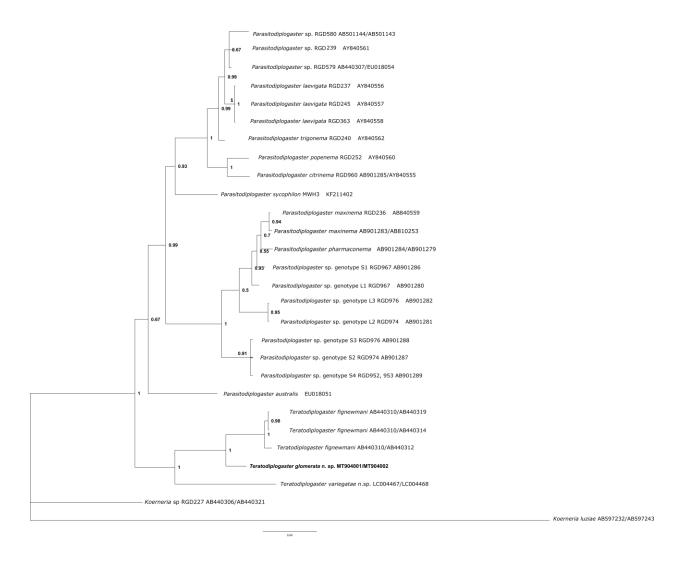


Fig. 8. The Bayesian tree inferred from the 18S and 28S gene for *Teratodiplogaster glomerata* n. sp. under the GTR + I + G model (lnL = 6263.9613; freqA = 0.1647; freqC = 0.2377; freqG = 0.336; freqT = 0.2617; R (a) = 0.0565; R (b) = 0.2121; R (c) = 0.1209; R (d) = 0.0504; R (e) = 0.4818; R (f) = 0.0783; Pinva = 0.251; Shape = 0.915). The accession numbers of the compared sequences are indicated in the form: SSU/D2-D3 LSU. Posterior probability values exceeding 50% are given on appropriate clades.

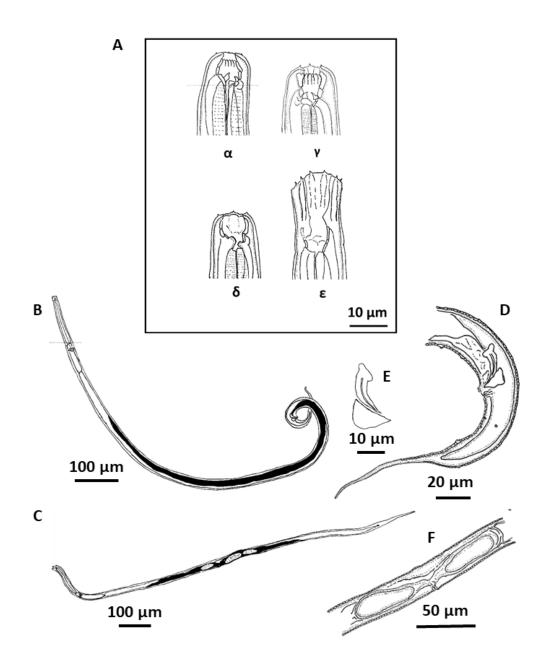


Fig. 9. *Pritionchus glomerata* n. sp. lateral view (A) Different morphs anterior structure, (B) Habitus of male, (C) Habitus of female, (D) Tail region of male, (E) Spicule and (F) Vulval region of female.



Fig. 10. Photomicrographs of α morph (adult females only) of *Pristionchus glomerata* n. sp. (A) Head morphology, (B) Anterior and posterior pharynx, (C) different focal plane, arrows show spicule in A and capitulum in B, (C) Gonadal region of female showing vulval opening and eggs, (D) Anal opening and tail region (E – F). Scale bar: (A – F) 10 µm.

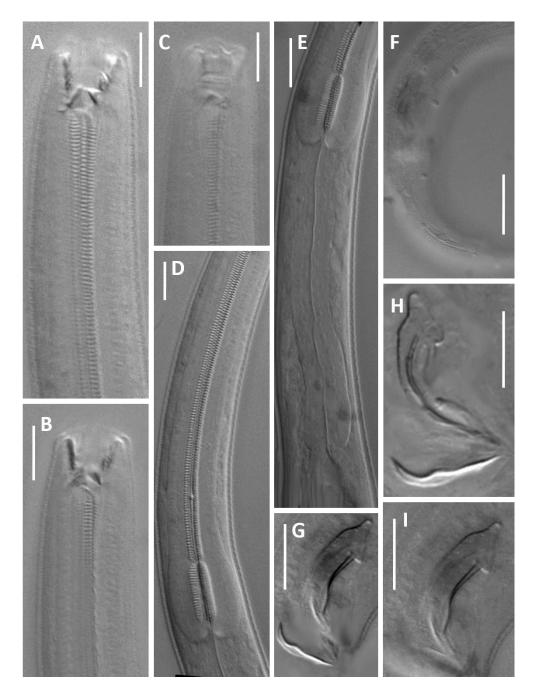


Fig. 11. Photomicrographs of γ morph (adult males only) of *Pristionchus glomerata* n. sp. (A – C) Head morphology at different focal planes, (D) Anterior pharynx, (E) Posterior pharynx, (F) Genital papillae, (G – I) male spicule and capitulum at different focal planes. Scale bar: (A – I) 10 μ m.

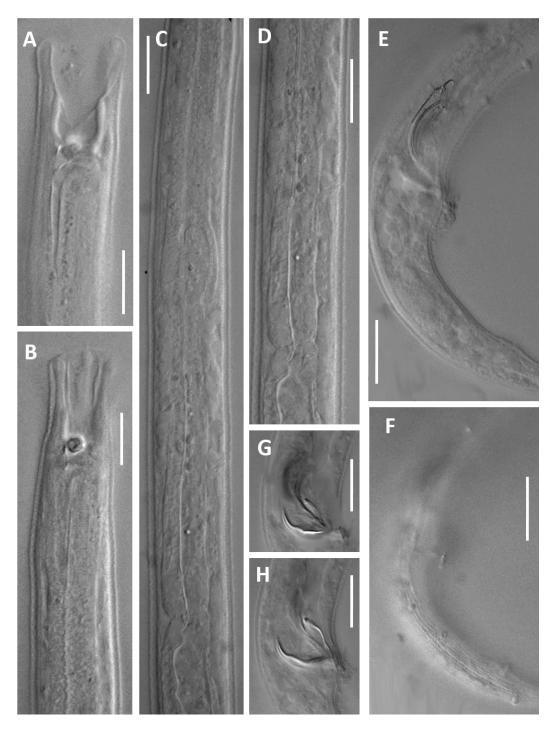


Fig. 12. Photomicrographs of ε morph (adult males only) of *Pristionchus glomerata* n. sp. (A – B) Head morphology at different focal planes, (C – D) Anterior pharynx and posterior pharynx, (E – F) Tail region at different focal planes showing genital papillae, (G – H) male spicule and capitulum at different focal planes. Scale bar: (A – H) 10 µm.

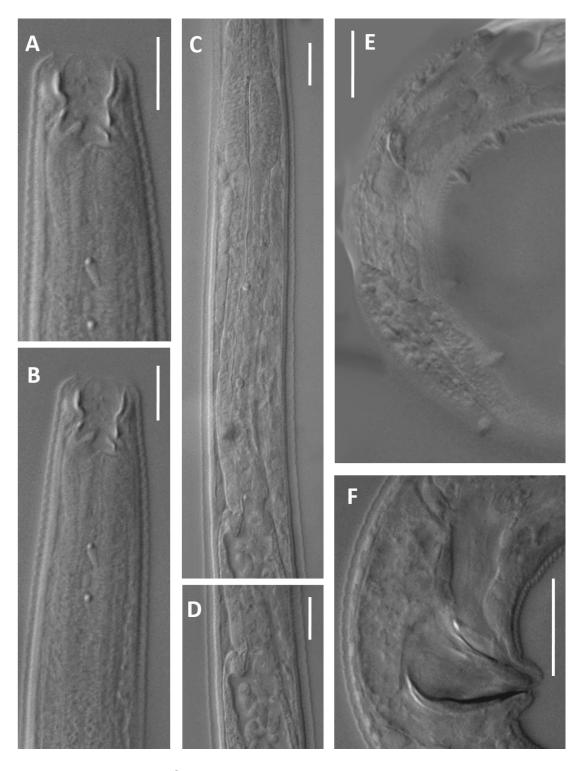


Fig. 13. Photomicrographs of δ morph (adult males only) of *Pristionchus glomerata* n. sp. (A – B) Head morphology at different focal planes, (C – D) Anterior and posterior pharynx, (E) Genital papillae, (F) male spicule and capitulum. Scale bar: (A – F) 10 µm.

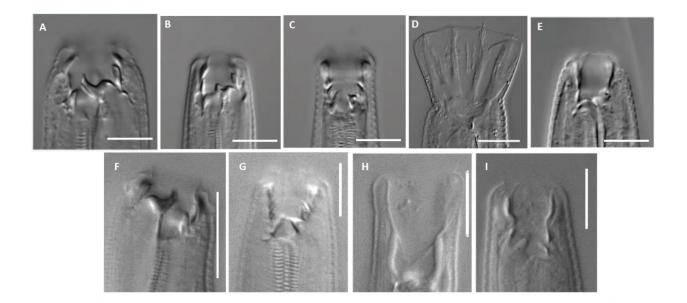


Fig. 14. Discrete comparable adult morphs present in *P. racemosae* (A–E)[taken from Susoy *et al.* 2016) and *P. glomerata* n. sp. (F–I). The species morphs of *P. racemosae* (from left to right) : α , β , γ , ε and δ and for *P. glomerata* n. sp. (from left to right): α , γ , ε and δ Scale bar: (D) 20 μ m, (A–C, E–I) 10 μ m.

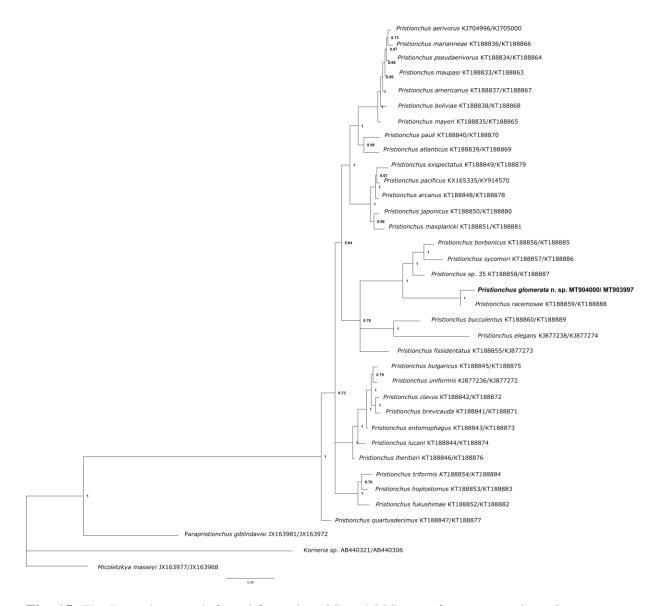


Fig. 15. The Bayesian tree inferred from the 18S and 28S gene for *Pristionchus glomerata* n. sp. under the GTR + I + G model (lnL = 8781.2399; freqA = 0.2345; freqC = 0.1834; freqG = 0.3026; freqT = 0.2795; R (a) = 0.0543; R (b) = 0.2938; R (c) = 0.1071; R (d) = 0.0836; R (e) = 0.3843; R (f) = 0.0769; Pinva = 0.168; Shape = 0.852). The accession numbers of the compared sequences are indicated in the form: SSU/D2-D3 LSU. Posterior probability values exceeding 50% are given on appropriate clades.

Table 1. Morphometric characteristics of Ficophagus glomerata n. sp. All measurements are in µm and in

the form: mean \pm s.d. (range).

	Ficophagus. glomerata n. sp.									
	М	ale	Female							
Character	Holotype	Paratypes	Paratypes							
1		9	10							
L	525	567 ± 58 (469-656)	761 ± 81 (604-886)							
a	25	26.5 ± 2.4 (23.4-30.9)	24.0 ± 2.3 (20-28)							
b	7.3	7.4 ± 0.3 (6.7-7.8)	7.6 ± 1.4 (5.2-9.6)							
2	17.2	16.6 ± 2.6 (12-20)	20.7 ± 1.6 (18-23)							
с'	1.8	2.0 ± 0.4 (1.5-2.7)	1.6 ± 0.2 (1.3-1.9)							
T or V (%)	76.9	72 ± 3.3 (65-77)	$\begin{array}{c} 68.6 \pm 4.9 \\ (61.2 \text{-} 76.7) \end{array}$							
Max. body diam.	20.9	21.3 ± 1.0 (19-23)	31.4 ± 0.7 (30.2-32.1)							
Stylet length	21.2	21.3 ± 0.6 (20-22)	22.0±0.9 (20-23)							
Median bulb (length)	16.6	16.2 ± 0.6 (14-17)	14.3 ± 0.9 (12.4 -15.6)							
Median bulb (width)	10.6	$\begin{array}{c} 10.6 \pm 0.3 \\ (10\text{-}11) \end{array}$	10.4 ± 0.5 (10-11)							
Ant.enf to SE pore	14	14 ± 0.7 (13-15)	14.5 ± 0.8 (13-16)							
Ant. end to base of metacorpus	55	59 ± 5.5 (51-68)	64.6 ± 2.3 (62-69)							
Ant. end to base of pharyngeal gland	114	80.4 ± 12 (68-114)	101.5 ± 15 (78-114)							
Nerve ring from anterior end	68	70.8 ± 6 (62-82)	77.9 ± 6.1 (71-87.4)							
Post uterine sac	-	_	6.0 ± 0.3 (5.4-6.3)							
Tail length	35.1	33.5 ± 6.6 (23.6-42)	38.9 ± 2.1 (34.5-42.3)							
Spicule length (chord)	16.6	16.8 ± 1 (14-17)								

Table 2. Comparative morphometric data of *Ficophagus glomerata* n. sp., other congeners and two *Schistonchus* sp. described from India, associated with *Ficus racemosa*. All measurements are in µm and in the form mean ± s.d. (range).

Character	S. rac	emosa	<i>S. os</i>	smani	F. ac	uleate	F. fl	eckeri	F. alterm	acrophylla	F. cucullo	oracemosus	F. flagello	racemosus	F. glome	rata n. sp. *
	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female
	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes
n	35	35	30	30	15	14	22	27	12	10	11	14	12	15	9	10
L	653 (500-660)	792 (600-800)	(700-840)	(718-910)	450 ± 33 (396-523)	619 ± 35 (548-671)	547 ± 32 (498-609)	727 ± 78 (572-889)	560 ± 44 (445-618)	500 ± 49 (411-571)	523.6±56.1 (441-609)	785.7±57.5 (707-906)	379.8±28.6 (342-446)	475.9±18.3 (442-504)	567 ± 58 (469-656)	761 ± 81 (604-886)
a	26 (24-26)	117.2 (14-21)	(26-30)	(27-34)	27 ± 2.8 (23-33)	18 ± 2.2 (14-23)	30 ± 1.5 (27-33)	23 ± 3.9 (15-35)	28 ± 2 (25-31)	22 ± 3 (17-29)	26.6±1.57 (24.5-29.3)	22.1±2.81 (17.6-27.8)	27.3±3.3 (22.8-32.9)	26.9±1.52 (24.3-30.4)	26.5 ± 2.4 (23.4-30.9)	$24.0 \pm 2.3 \\ (20-28)$
b	5.4 (5.2-6.2)	(5.2-6.2)	(10-13)	(11.5- 13.5)	3.3 ± 0.2 (2.6-4.0)	$\begin{array}{c} 4.9 \pm 0.2 \\ (4.7 \text{-} 5.1) \end{array}$	$5.5 \pm 0.5 \\ (4.8-6.4)$	7.2 ± 0.3 (7.0-7.4)	4 ± 0.6 (3-5)	5 ± 1.2 (3-8)	7.9±0.88 (6.0-9.5)	12.3±1.12 (10.9-15.1)	9.1±0.84 (8.1-11.2)	9±0.54 (8.0-10.1)	7.4 ± 0.3 (6.7-7.8)	7.6 ± 1.4 (5.2-9.6)
с	28 (22-28)	21.6 (18-22)	(12-15)	(14.8- 19.6)	12.2 ± 1.0 (11-14)	19.3 ± 2.0 (17-23)	18 ± 1.5 (16-21)	21 ± 3.4 (12-29)	15 ± 1 (13-16)	17 ± 2 (14-21)	17.6±1.3 (16-20.5)	16.9±2.36 (14.5-21.1)	24.3±2.1 (21.1-27.5)	13.9±1.1 (12.8-16.8)	16.6 ± 2.6 (12-20)	$20.7 \pm 1.6 \\ (18-23)$
c'	-	-	-	-	$\begin{array}{c} 2.3 \pm 0.2 \\ (1.8 \text{-} 2.8) \end{array}$	2.4 ± 0.4 (2.0-3.1)	$\begin{array}{c} 1.9 \pm 0.2 \\ (1.6 \text{-} 2.4) \end{array}$	2.6 ± 0.7 (1.6-4.6)	2 ± 0.2 (1.7-2.4)	3 ± 0.3 (2.8-3.8)	1.8±0.14 (1.7-1.9)	3.4±0.5 (2.7-4.1)	1.3±0.09 (1.2-1.4)	4.5±0.48 (3.4-5.2)	2.0 ± 0.4 (1.5-2.7)	1.6 ± 0.2 (1.3-1.9)
T or V (%)	31 (30-48)	60 (60-76)	-	(78-84)	55 ± 1 (48-57)	75 ± 1.2 (73-77)	41 ± 13.4 (22-80)	76 ± 4.3 (67-86)	330 ± 49 (273-401)	85 ± 9 (70-95)	-	73.2±2.0 (70.1-77.6)	-	76.2±0.96 (74.7±78.2)	72 ± 3.3 (65-77)	68.6 ± 4.9 (61.2-76.7)
Max. body diam.	-	-	(25-28.5)	(21-27)	17 ± 1 (15-19)	34 ± 3 (29-40)	18 ± 1.5 (15-19)	33 ± 7 (19-54)	18 ± 2 (15-19)	23 ± 2 (19-25)	-	-	-	-	21.3 ± 1.0 (19-23)	$\begin{array}{c} 31.4 \pm 0.7 \\ (30.2 \text{-} 32.1) \end{array}$
Stylet length	-	23 (21-24)	(19-21)	(21-27)	15 ± 2 (13-18)	14 ±1.5 (13-17)	17 ± 0.8 (15-17)	18 ± 1.5 (15-19)	22 ± 3 (16-25)	21 ± 2 (17-26)	22 (22)	22 (22)	14 (14)	14 (14)	21.3 ± 0.6 (20-22)	22.0±0.9 (20-23)
Median bulb (length)	17 (13-17)	17 (13-17)	(10-12)	(12-13)	-	-	-	-	-	-	-	-	-	-	16.2 ± 0.6 (14-17)	14.3 ± 0.9 (12.4 -15.6)
Median bulb (width)	13 (10-13)	15 (12-15)	(8-9)	(8-9)	-	-	-	-	-	-	-	-	-	-	$\begin{array}{c} 10.6 \pm 0.3 \\ (10\text{-}11) \end{array}$	10.4 ± 0.5 (10-11)
Ant. end to Ex pore	-	-	-	-	6 ± 2.0 (4-9)	6 ± 1.1 (5-8)	10 ± 1.4 (8-11)	8.0 ± 0.6 (7-8)	6 ± 1 (3-7)	6 ± 2 (5-9)	10±1.89 (7-12)	10.1±1.8 (7-14)	16.7±2.0 (12-19)	11.4±0.44 (11-13)	14 ± 0.7 (13-15)	14.5 ± 0.8 (13-16)

Ant. End to base metacorpus	-	-	-	-	53 ± 3 (49-58)	54 ±7 (47-77)	66 ± 3.7 (60-73)	58 ± 9.5 (41-87)	65 ± 7 (60-71)	69 ± 6 (55-77)	-	-	-	-	59 ± 5.5 (51-68)	64.6 ± 2.3 (62-69)
Ant. End to base pharyngeal gland	-	-	-	-	135 ± 7 (123-154)	127 ± 5 (117-135)	100 ± 6 (87-110)	119 ± 11 (112-127)	130 ± 20 (112-163)	10 ± 20 (114-168)	-	-	-	-	80.4 ± 12 (68-114)	101.5 ± 15 (78-114)
Nerve ring from anterior end	-	-	-	-	60 ± 3 (53-65)	57 ± 4 (52-63)	72 ± 4.5 (65-78)	66 ± 3 (61-70)	72 ± 8 (60-85)	76 ± 8 (60-83)	-	-	-	-	70.8 ± 6 (62-82)	77.9 ± 6.1 (71-87.4)
Post uterine sac	-	-	-	-	-	6 ± 1.5 (4-9)	-	18 ± 8 (6-39)	-	14 ± 3 (8-20)	-	-	-	-	-	6.0 ± 0.3 (5.4-6.3)
Tail length	-	-	-	-	37 ± 4 (31-43)	32 ± 3 (28-36)	30 ± 3 (26-38)	37 ± 8 (24-62)	34±2 (28-37)	29 ± 4 (23 -34)	29.7±2.1 (25-32)	47.3±7.58 (35-62)	15.7±0.86 (14-16)	34.3±2.58 (30-38)	33.5 ± 6.6 (23.6-42)	38.9 ± 2.1 (34.5-42.3
Spicule length (chord)	-	-	(37-39.5)	-	16 ± 1 (15-17)	-	17 ±0.9 (16-18)	-	21 ± 2 (18-24)	-	21.4±0.67 (20-22)	-	10.1±0.5 (9-11)	-	16.8 ± 1 (14-17)	-

4.6 ± 2.3 (62-69))1.5 ± 15 78-114) 7.9 ± 6.1 71-87.4) .0 ± 0.3 5.4-6.3) 3.9 ± 2.1 4.5-42.3)

Table 3. Comparative morphometric data of *Teratodiplogaster glomerata* n. sp. and other congeners associated with *Ficus racemosa*. All measurements are in μ m and in the form: mean \pm s.d. (range).

Character	T. var	T. variegatae		T. fignewmani		T. martini		T. racemosus			T. glomerata n. sp.		
	Male	Female	Male	Female	Male	Female	Male	Female	1	Male	Female		
	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Paratypes	Holotype	Paratypes	Paratypes		
	9	10	10	11	10	13	9	13		9	10		
	1019 ± 117 (854-1283)	965 ± 120 (805-1173)	2119 ± 311 (1850-2700)	2705 ± 636 (2160-3895)	2171 ± 267 (1825-2650)	2253 ± 325 (1750-3100)	1771.1 ± 295.6 (1412-2184)	2030.9 ± 236.2 (1587-2444)	1329	1271 ± 48 (1195-1345)	1506 ± 83 (1388-1657)		
	40.9 ± 4.6 (29.9-47.4)	30.7 ± 5.8 (26.1-45.6)	$\begin{array}{c} 67.8 \pm 6.6 \\ (52.9\text{-}75.9) \end{array}$	$\begin{array}{c} 79.9 \pm 10.6 \\ (59.0 \text{-} 92.0) \end{array}$	62.6 ± 4.6 (52.6-70.0)	67.1 ± 6.9 (54.5-79.5)	77.9 ± 10 (66.5-92.8)	81.4 ± 4.4 (74.7-89.1)	63.3	64.7 ± 2.9 (62.0-69.6)	72.4 ± 3.4 (68.8-79.7)		
	4.4 ± 0.4 (3.8-5.4)	4.2 ± 0.3 (3.8-4.9)	9.1 ± 1.3 (7.8-11.7)	11.0 ± 1.5 (9.5-13.5)	8.2 ± 0.5 (7.7-9.3)	8.1 ± 0.7 (6.7-9.3)	8.2 ± 1.1 (6.5-9.4)	8.9 ± 0.5 (7.7-9.8)	6.2	6.1 ± 0.1 (5.9-6.3)	7.7 ± 0.3 (7.3-8.2)		
	16.9 ± 1.4 (15.0-19.7)	$18.2 \pm 1.2 \\ (17.0-19.9)$	19.4 ± 2.6 (16.9-26.0)	13.0 ± 1.9 (10.2-16.4)	21.6 ± 2.4 (18.1-25.0)	$19.4 \pm 2.9 \\ (15.9-26.3)$	$\begin{array}{c} 14.9 \pm 1.6 \\ (12.9\text{-}17.8) \end{array}$	8.3 ± 0.5 (7.3-11.2)	12	11.4 ± 0.8 (10.6-12.8)	10.2 ± 1.6 (8.3-13.0)		
	3.1 ± 0.1 (3.0-3.4)	3.5 ± 0.2 (3.2-3.8)	5.1 ± 0.6 (3.4-5.7)	9.3 ± 1.8 (6.8-11.9)	4.2 ± 0.4 (3.5-4.9)	5.8 ± 0.6 (4.8-7.3)	7 ± 0.51 (6.2-7.9)	15.7 ± 1.8 (11.8-19)	7.1	7 ± 0.6 (6.2-8.5)	$\begin{array}{c} 10.9 \pm 1.9 \\ (7.7\text{-}13.5) \end{array}$		
v	40.2 ± 6.2 (29.8-51.3)	$59.4 \pm 1.0 \\ (58.3-61.1)$	61.9 ± 9.2 (51.9-83.7)	50.0 ± 1.5 (46.2-52.2)	69.1 ± 6.8 (58.5-80.6)	50.5 ± 1.7 (47.8-53.3)	-	-	64	58.5 ± 5.3 (51.4-65.6)	49.7 ± 2.3 (46.5-55.8)		
ax. body m.	25 ± 4.1 (21-35)	32 ± 3.4 (24-35)	32 ± 8.2 (26-51)	35 ± 13.0 (25-66)	35 ± 5.5 (29-45)	34 ± 5 (25-44)	-	-	21	20 ± 0.9 (18-21)	21 ± 0.8 (19-22)		
oma diam.	4.4 ± 0.6 (3.4-5.3)	4.2 ± 0.6 (3.4-4.8)	-	-	5.0 ± 0.8 (4.0-6.0)	5.0 ± 0.9 (4.0-7.0)	-	-	-	-	-		
oma length	22 ± 1.2 (20-23)	22 ± 1.1 (20-24)	23 ± 1.4 (20-25)	23 ± 2.4 (20-28)	26 ± 3.4 (21-30)	27 ± 5.7 (15-34)	22.5 ± 1.6 (20-25)	22.9 ± 1.5 (21-26)	21	$20.1 \pm 0.8 \\ (19-21)$	21.9 ± 1.1 (20-23)		
rior ynx h	130 ± 6.8 (120-142)	125 ± 7.3 (115-138)	126 ± 8.0 (112-134)	139 ± 24 (102-178)	168 ± 28 (10-240)	174 ± 17 (150-200)	131.1 ± 12.8 (110-150)	138.7 ± 12.1 (119-160)	139	132 ± 10 (110-146)	135.5 ± 5.4 (129-148)		
sterior arynx gth	74 ± 4.8 (63-80)	76 ± 5.1 (70-85)	81 ± 7.3 (68-92)	82 ± 14 (43-98)	73 ± 10 (50-85)	75 ± 17 (50-110)	78.9 ± 7.5 (71-90)	83.2 ± 4 (77-89)	54	56 ± 6 (47-69)	57.7 ± 11.1 (46-85)		

Ratio of anterior and posterior pharynx	1.8 ± 0.1 (1.6-2.0)	1.7 ± 0.1 (1.5-1.7)	1.6 ± 0.1 (1.4-1.7)	1.7 ± 0.3 (1.4-2.4)	-		1.7±0.1 (1.5- 1.8)	1.7 ± 0.1 (1.5-1.8)	2.57	$2.26 \pm 0.3 \\ (1.6-2.7)$	$2.42 \pm 0.4 \\ (1.6-2.9)$
Nerve ring from anterior end	177 ± 10 (162-189)	172 ± 10 (156-188)	188 ± 13 (177-214)	202 ± 33 (143-252)	-	-	-	-	186	178 ± 8 (157-186)	187 ± 8 (178-203)
Excretory pore from anterior end	190 ± 12 (167-210)	186 ± 13 (170-206)	217 ± 19 (202-261)	235 ± 49 (166-321)	-	228 ± 17 (206-252)	196.4 ± 14.1 (177-215)	198.5 ± 19.6 (172-224)	207	196 ± 7 (183-207)	180 ± 10 (163-196)
Testis length	408 ± 71 (327-561)	-	1334 ± 409 (1050-2260)	-	1493 ± 168 (1225-1750)	-	-	-	851	746 ± 86 (614-866)	362 ± 45 (312-423)
Anterior ovary length	-	76 ± 11 (58-93)	-	294 ± 174 (150-603)	-	233 ± 44 (170-325)	-	-	-	-	360 ± 46 (308-426)
Posterior ovary length	-	85 ± 17 (65-115)	-	268 ± 141 (160-560)	-	230 ± 54 (130-350)	-	-	-	-	362 ± 45 (312-423)
Anal or cloacal body diam.	19 ± 1.8 (16-21)	15 ± 1.3 (13.5-17.5)	22 ± 3.3 (19-30)	23 ± 6.4 (17-35)	24 ± 2.8 (21-29)	20 ± 2.9 (17-28)	-	-	14	16 ± 1.8 (13-19)	13.7 ± 0.4 (13-14)
Tail length	60 ± 4.5 (51-65)	53 ± 4.7 (46-59)	110 ± 11 (98-136)	207 ± 27 (163-256)	101 ± 8.7 (88-116)	118 ± 13 (92-138)	118 ± 12.7 (95-137)	$245.8 \pm 41.1 \\ (177-298)$	99	110 ± 6 (99-118)	151 ± 28 (108-189)
Spicule (curved)	27 ± 2.4 (24-30)	-	27 ± 2.1 (25-32)	-	25 ± 3.6 (18-28)	-	20.7 ± 2.1 (19-25)	-	19	19 ± 1.0 (17-20)	-
Spicule (chord)	23 ± 2.0 (20-26)	-	24 ± 2.0 (23-29)	-	24 ± 3.8 (17-30)	-	-	-	18	18 ± 1.0 (16-19)	-
Gubernaculum length	11 ± 0.7 (10-12)	-	15.0 ± 1.0 (13-16)	-	-	-	11.7 ± 1.3 (10-14)	-	10	12 ± 1 (10-14)	-

 2.42 ± 0.4 (1.6-2.9) 187 ± 8 (178-203) 180 ± 10 (163-196) 362 ± 45 (312-423) 360 ± 46 (308-426) 362 ± 45 (312-423) 13.7 ± 0.4 (13-14)

Table 4. Morphometrics of Pristionchus glomerata n. sp. collected from Ficus racemosa of Southern

	Pristionchus. glomerata n. sp.									
		Male	Female							
Character	Holotype (ε morph)	Paratypes	Paratypes							
n		9	10							
L	1253.2	$\frac{1185.6 \pm 122.6}{(1023.8-1411.8)}$	$\begin{array}{r} 1067.7 \pm \ 257.6 \\ (693.4 \text{-} 1539.5) \end{array}$							
a	57.2	53.3 ± 6.7 (31.1-66.8)	51.8 ± 10.7 (37.7-68.2)							
b	6.7	6.5 ± 0.6 (5.6-7.6)	5.9 ± 1.2 (4.3-7.9)							
с	8.8	9.3 ± 1.0 (8.2-11.5)	7.5 ± 0.9 (5.8-9)							
c'	7.6	7.1 ±0.5 (6.5-7.8)	9.2± 1.9 (7.1 -12)							
T or V	62.3	61.2 ± 4.8 (54-68.8)	51.3 ± 5.6 (37.9-58.7)							
Max. body diam.	21.9	21.6 ± 1.5 (10.1-24.1)	20.9 ± 2.9 (16.6-24.6)							
Stoma length	7.6	12.5 ±6.2 (6.3-18.4)	7.7 ± 2.9 (5.1 - 14.9)							
Anterior pharynx length	110.9	109.8 ± 4.3 (104.2-119.4)	101.5 ± 13.0 (90.3-128.2)							
Posterior pharynx length	74	73.7 ± 4.2 (68.1-82.3)	$75.6 \pm 10.8 \\ (59.2-87.1)$							
Ratio of anterior and posterior pharynx	1.5	1.46 ± 0.04 (1.4-1.5)	$\begin{array}{c} (3.12 + 0.17) \\ 1.36 \pm 0.15 \\ (1.1 - 1.5) \end{array}$							
Neck length (head to the base of pharynx)	192.5	196.0 ± 9.7 (185.5-217.5)	184.85 ± 21.4 (158.6-230.2)							
Excretory pore from anterior end	53.4	59.4 ± 6.9 (52.4-75.4)	51.9 ± 5.7 (46.4-63.7)							
Anterior ovary length	-	-	$(40.\pm0.7)$ 226.3 ± 44.9 (178.8-314.6)							
Posterior ovary length	-	-	268.3 ± 55.9 (325.1-187.2)							
Testis length	780.3	728 ± 112.9 (562.8-910.5)								
Vulva to anus distance	-	-	392.6 ± 170.5 (783.7 -196.6)							
Anal or cloacal body diam.	18.7	17.9 ± 1.4	15.4 ± 1.8							
Tail length	142.4	(16.1-21.4) 126.9 ± 9.3 (113.4, 142.4)	(12.2-16.8) 139.9 ± 23.7 (117.0, 172.6)							
Spicule (curved)	21.2	(113.4-142.4) 21.42 ± 0.5 (20.2, 22.4)	(117.0-172.6)							
Spicule (chord)	17.2	(20.2-22.4) 17.3 ± 0.1 (17.1 ± 17.6)	-							
Gubernaculum length	16.5	(17.1-17.6) 16.4 ± 0.1 (16.2-16.8)	-							