

# Redescription of *Micrura dellechiajei* (Hubrecht, 1879) (Nemertea, Pilidiophora, Lineidae), a rare Mediterranean species

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*The heteronemertean species Micrura dellechiajei is thus far only known from its type locality in the Gulf of Naples (Italy) and has not been recorded in 120 years. During two oceanographic surveys conducted in Spanish Mediterranean waters, several nemertean specimens were collected, and thorough morphological examination indicated that some of these pertained to the species M. dellechiajei, suggesting that populations may be more widespread than previously thought. Because of the rarity of this species coupled with the fact that its last morphological narrative was given 120 years ago, we here provide a redescription of the species based on the new specimens, complete with illustrations and new data concerning its morphology, and we also place some of the collected specimens in a molecular phylogenetic framework.*

**Keywords:** Heteronemertea, Pilidiophora, *Micrura dellechiajei*, redescription, Mediterranean sea, COI mtDNA, 16S rRNA, phylogeny

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## INTRODUCTION

*Micrura dellechiajei* (Hubrecht, 1879) (Nemertea: Pilidiophora: Lineidae) is an elusive ribbon worm, with a conspicuous lack of information regarding its geographic distribution and general natural history. The species seems to have been first recorded from the general region surrounding Naples, Italy by Delle Chiaje (1829) who attributed his specimens to a species already known at the time, *Cerebratulus bilineatus* (Renier, 1804). However, the formal description of the species was synthesized fully 50 years later by Hubrecht (1879), based on specimens collected from around Capri, an island south of Naples, in the same vicinity as Delle Chiaje's (1829) collection site. These specimens proved to be morphologically compatible with those collected by Delle Chiaje (1829). As opposed to Delle Chiaje (1829), however, Hubrecht (1879) identified morphological apomorphies possessed by the new specimens and, accordingly, erected a new species, *Cerebratulus dellechiajei* Hubrecht, 1879 to accommodate the specimens. Beyond obvious differences in colour patterns between *C. bilineatus* and *C. dellechiajei*, Hubrecht (1879) also noted that, in contrast to *C. bilineatus*, *C. dellechiajei* possesses a short caudal cirrus and numerous eyes – attributes that had come to unify species of the genus *Micrura* Ehrenberg, 1831. As a consequence, Bürger (1895) transferred the species to the genus *Micrura* and provided extensive illustrations of external and internal anatomy of the species. Although Bürger's illustrations

(1895; plate 4, figures 18, 23, 24, 26, 27, 33) are adequate in allowing for identification of specimens belonging to the species, the aged description by Hubrecht (1879) is insufficient in several regards, and thus in dire need of a complete update, using contemporary data. However, since Bürger's (1895) treatment of the species about 120 years ago, no specimens have been recorded and, therefore, no data have been available on which to base a redescription. Numerous gracile heteronemertean species possessing a caudal cirrus and cephalic slits have traditionally been included in the genus *Micrura* but modern molecular phylogenetic studies have revealed an astonishing level of paraphyly within the genus (e.g., Sundberg & Saur, 1998; Schwartz, 2009; Andrade *et al.*, 2012; Kvist *et al.*, 2014). This indicates that the putative synapomorphies previously considered for the genus were wrong, increasing the need for a modern redescription of *M. dellechiajei*, as well as a reconsideration of the entire genus.

During September 2011 and July 2012, numerous nemertean samples were collected as part of an oceanographic campaign (the INDEMARES Project) in the Alborán Sea in southern Spain. This collection effort yielded several specimens of *Micrura dellechiajei*, which had also previously been collected during the 2007/2008 CALMEN oceanographic campaign to the Menorca Channel (Balearic Islands, Spain) – these specimens, however, were poorly preserved and proved refractory to further molecular studies. Importantly, a combination of morphological characters and molecular data seems to be the best approach for inferring species delimitations within Nemertea (Sundberg *et al.*, 2010; Sundberg & Strand, 2010; Strand & Sundberg, 2011; Leasi & Norenburg, 2014; Strand *et al.*, 2014). Therefore, we herein provide a thorough redescription of *Micrura dellechiajei*, based on traditional histological sections as well as external anatomical

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observations, of samples from both the Alborán Sea and the Menorca Channel, approximately 1500 km west of its type locality. We also provide a molecular phylogeny of Heteronemertea to estimate the placement of the species within the class and to ensure the monophyly of the sampled specimens, and we briefly discuss these results in the context of the non-monophyly of most genera within Heteronemertea.

## MATERIALS AND METHODS

### Material examined

During research cruises CALMENo7, CALMENo8 in the Menorca Channel (Balearic Islands, Spain) in July 2007 and June 2008, as well as ALBORÁN-INDEMARES in the Alborán Sea (Spain) in September 2011 and July 2012 (Figure 1, Table 1), numerous nemertean samples were collected by beam trawler and rock dredger. Depths ranged from 57 to 101 m and samples were generally associated with encrusting red algae, organically enriched sediments (shell sand) and communities of sponges and cnidarians. The following redescription of *Micrura dellechiaiei* is based on observations of 25 living specimens (Table 1) and three sets of histological slides, two of these being from the white morphotype (Figure 1A) and one from the olive-colour morphotype (Figure 1C).

### Histology

For the present study, worms were measured and photographed while alive, relaxed in a 7.5% wt/vol solution of MgCl<sub>2</sub> mixed 1:1 with filtered seawater, and fixed in Bouin's (cupric picro-formolacetic) solution. Specimens destined for histology were dehydrated in a graded series of ethyl alcohols, cleared with multiple changes of toluene, and embedded in 56°C mp paraffin wax. Three specimens from Alborán Sea were sectioned and studied for histological features. For this purpose, samples were sectioned at 6 µm, mounted on slides, and stained with Mallory triple stain.

### Molecular analyses

DNA was extracted, amplified, purified and sequenced from three specimens of *Micrura dellechiaiei* (one of them from the white morphotype [Figure 1A] and two of them from the olive-colour morphotype [Figure 1C]) following the exact protocol



Fig. 1. Map showing the collection localities of the sampled specimens of *Micrura dellechiaiei* (Hubrecht, 1879) in the Menorca Channel and the Alborán Sea (stars). The circle denotes the type locality of the species off the coast of Naples, Italy.

described by Kvist *et al.* (2014), but the present study targeted only the mitochondrial loci cytochrome *c* oxidase subunit I (COI) and 16S rRNA. These loci were chosen based on their high rate of nucleotide divergence, i.e. their pertinence for teasing out genetic differences between morphologically similar species (see Discussion), coupled with the fact that one of our specimens has been previously sequenced for six loci and placed in a broader phylogenetic context (see Kvist *et al.*, 2014). Subsequently, the newly generated sequences were conjoined with a set of sequences previously compiled for other palaeonemertean, heteronemertean and hoplonemertean taxa (Table 2) and these were jointly aligned using MAFFT ver. 7 (Katoh & Standley, 2013) employing the L-INS-i strategy and applying default settings. Phylogenetic analyses used both maximum likelihood (ML) and parsimony (MP) methodologies. For ML, RAXML ver. 7.6.3 (Stamatakis, 2006) was applied on the CIPRES Science Gateway platform (Miller *et al.*, 2010) using a partitioned GTR +  $\Gamma$  model of nucleotide evolution, with optimal partition schemes suggested by PartitionFinder ver. 1.1.1 (Lanfear *et al.*, 2012). A heuristic search consisted of 1000 iterations with 25 initial GAMMA rate categories and final optimization with four GAMMA shape categories. For MP, the software TNT (Goloboff *et al.*, 2008) was used with the following settings: 1000 initial addition sequences, five rounds of ratcheting and three rounds of tree fusing, and requiring that the minimum length tree be found a total of 10 times. For both MP and ML, multiparametric bootstrap support values were calculated from 1000 pseudoreplicates with a different starting tree for each iteration. All trees were rooted at the palaeonemertean *Cephalothrix bipunctata* Bürger, 1892 following the phylogenetic hypothesis of Kvist *et al.* (2014).

Table 1. List of localities of the sampled specimens of *Micrura dellechiaiei* in the Menorca Channel (Balearic Islands) and Alborán Sea (South Spain).

No. of specimens	Date	Depth (m)	GPS coordinates	Cruise	Sample code	Sampling method
2	03/07/2007	61	39° 46' 00" N 03° 34' 00" E	Calmeno7	Lance 6	Trawling
2	04/07/2007	57	39° 57' 30" N 03° 45' 15" E	Calmeno7	Lance 7	Trawling
4	30/06/2008	60	40° 00' 00" N 03° 43' 60" E	Calmeno8	DR0044	Rock dredger
3	23/09/2011	96	35° 52' 43" N 03° 04' 40" W	Alborán-Indemares	BV14	Beam trawling
1	23/09/2011	96	35° 52' 40" N 03° 04' 39" W	Alborán-Indemares	BV15	Beam trawling
1	23/09/2011	92	35° 53' 06" N 03° 04' 44" W	Alborán-Indemares	BV16	Beam trawling
7	24/09/2011	101	36° 00' 24" N 02° 55' 19" W	Alborán-Indemares	BV21	Beam trawling
2	08/09/2011	80	35° 53' 24" N 03° 04' 40" W	Alborán-Indemares	DR01	Rock dredger
1	24/09/2011	48	35° 57' 39" N 02° 58' 51" W	Alborán-Indemares	DR20	Rock dredger
2	21/07/2012	81	36° 00' 28" N 02° 53' 16" W	Alborán-Indemares	DR40	Rock dredger

**Table 2.** List of specimens included in the phylogenetic analyses, with MCZ voucher numbers (more information on each specimen is accessible through <http://mczbase.mcz.harvard.edu/>), and GenBank accession numbers. Specimens newly sequenced for the present study are denoted in bold font.

Specimen ID	MCZ voucher	COI	16S
<i>Cephalothrix bipunctata</i>	IZ-133009	KF935501	KF935447
<i>Cephalothrix filiformis</i>	IZ-135328	HQ848617	JF277593
<i>Cephalothrix filiformis</i>	IZ-135327	HQ848616	JF277594
<i>Cephalothrix hongkongiensis</i>	IZ-134153	HQ848615	JF277590
<i>Cephalothrix hongkongiensis</i>	IZ-134153	HQ848614	JF277591
<i>Cephalothrix rufifrons</i>	IZ-135330	HQ848604	JF277592
Interstitial cephalotrichid	DNA 106139	HQ848618	JF277589
<i>Tetrastemma vittigera</i>	IZ-25171	KF935540	KF935491
<i>Tetrastemma matidae</i> sp.	IZ-132537	KF935542	KF935493
<i>Tetrastemma</i> sp.	IZ-132742	KF935541	KF935492
<i>Hubrechtella ijimai</i>	IZ-135342	KF935520	KF935470
<i>Hubrechtella dubia</i>	IZ-134224	HQ848631	JF277630
<i>Baseodiscus</i> cf. <i>delineatus</i>	IZ-133729	KF935502	KF935448
<i>Baseodiscus</i> sp.	IZ-135319	JF277568	HQ848588
<i>Baseodiscus mexicanus</i>	IZ-135321	KF935503	KF935449
<i>Baseodiscus unicolor</i>	IZ-135323	KF935505	KF935451
<i>Baseodiscus</i> sp.	IZ-135324	HQ848589	JF277569
<i>Micrura</i> sp.	IZ-133724	KF935509	KF935456
<i>Micrura</i> sp.	IZ-132532	KF935510	KF935457
<i>Micrura</i> sp.	IZ-132529	KF935511	KF935458
<i>Micrura verrilli</i>	IZ-134451	KF935508	KF935455
<i>Micrura fasciolata</i>	IZ-133719	HQ848578	JF277586
<i>Micrura fasciolata</i>	IZ-135347	HQ848577	JF277585
<i>Micrura ignea</i>	IZ-135349	KF935507	KF935454
<i>Micrura ignea</i>	IZ-133720	KF935506	KF935453
<i>Micrura ignea</i>	IZ-133721	HQ848587	JF277588
<i>Micrura rubramaculosa</i>	IZ-132531	KF935513	KF935460
<i>Micrura dellechiajei</i>	IZ-132745	KF935514	KF935461
<b><i>Micrura dellechiajei</i></b>	<b>IZ-132745</b>	<b>KP893673</b>	<b>KP893675</b>
<b><i>Micrura dellechiajei</i></b>	<b>IZ-132745</b>	<b>KP893674</b>	<b>KP893676</b>
<i>Micrura purpurea</i>	IZ-133723	HQ848586	JF277577
<i>Micrura chlorapardalis</i>	IZ-132530	KF935512	KF935459
<i>Zygeupolia rubens</i>	IZ-133731	HQ848585	JF277574
Freshwater heteronemertean	DNA 106130	HQ848584	JF277587
<i>Lineus</i> sp.	IZ-132744	KF935518	KF935468
<i>Lineus viridis</i>	IZ-135346	HQ848579	JF277582
<i>Lineus torquatus</i>	IZ-134170	HQ848574	JF277572
<i>Lineus acutifrons</i>	IZ-135343	GU590937	JF277573
<i>Lineus bilineatus</i>	IZ-135345	DQ280014	JF277571
<i>Notospermus</i> sp.	IZ-134234	KF935516	KF935465
<i>Notospermus</i> sp.	IZ-135528	KF935515	KF935463
<i>Cerebratulus lacteus</i>	IZ-134506	HQ848576	JF277575
<i>Cerebratulus leucopsis</i>	IZ-135331	KF935517	KF935467
<i>Cerebratulus marginatus</i>	IZ-134542	HQ848575	JF277576
<i>Ramphogordius sanguineus</i>	DNA 103903	HQ848580	JF277583
<i>Ramphogordius lacteus</i>	IZ-135373	KF935519	KF935469
<i>Ramphogordius lacteus</i>	IZ-135372	HQ848583	JF277584
<i>Riseriellus occultus</i>	IZ-135375	HQ848581	JF277581
<i>Riseriellus occultus</i>	IZ-135376	HQ848582	JF277580
<i>Riseriellus occultus</i>	IZ-135377	HQ848633	JF277579

## TAXONOMY

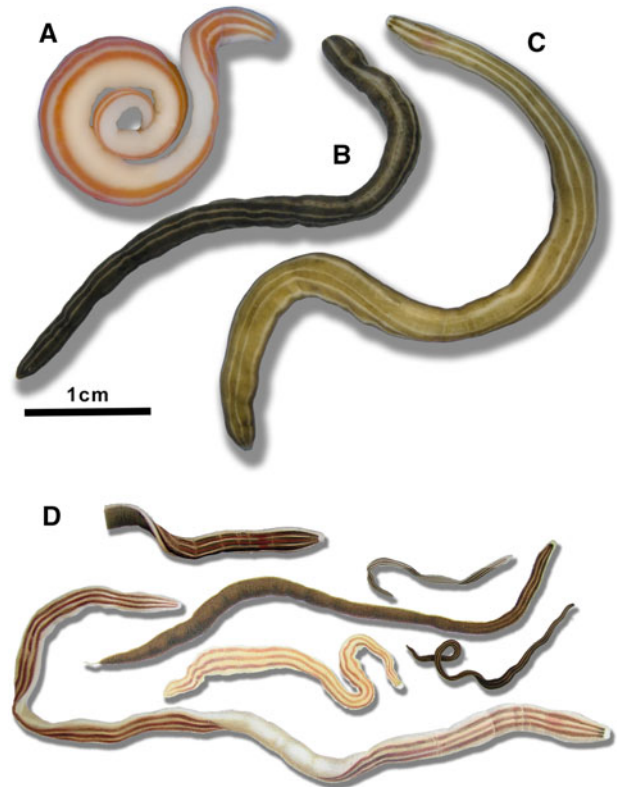
### SYSTEMATICS

Class ANOPLA Schultze, 1851  
 Subclass heteronemertea  
 Genus *Micrura* Ehrenberg, 1831

### DIAGNOSIS

From Gibson (1981): ‘Heteronemertea with a single pair of horizontal lateral cephalic slits, posteriorly enlarged to form wide bays; ciliated cerebral canals emerge from ventral wall of cephalic bays; proboscis unbranched, containing two (outer circular, inner longitudinal) or three (outer longitudinal, middle circular, inner longitudinal) muscle layers and none, one or two muscle crosses; rhynchocoel circular musculature not interwoven with body wall muscles; dorsal fibre core of cerebral ganglia bifurcated only at rear into upper and lower branches; nervous system with neither neurochords nor neurochord cells; ganglionic cell layer of brain usually not separated from body wall muscles by neurilemma; foregut with or without splanchnic musculature, if present variably composed of circular and/or longitudinal fibres; dermis variable, mostly with distinct connective tissue layer separating glandular zone from body wall muscles; caudal cirrus present; cephalic glands normally well developed, occasionally weakly formed or absent; frontal organ usually present; eyes present or absent; sexes separate.’

This wide diagnosis accommodates species attributed to *Micrura* and the validity of this diagnosis is compromised due to the non-monophyletic nature of the genus. However, the specimens used for the present study agree with this diagnosis, as well as those of Hubrecht (1879) and Bürger (1895), and given the absence of a better system for assigning heteronemerteans to genus, we leave the species in the genus *Micrura*.



**Fig. 2.** *Micrura dellechiajei* (Hubrecht, 1879). (A) photograph of a complete specimen from the Alborán Sea; (B) photograph of a complete specimen from the Menorca channel; (C) photograph of a complete specimen from the Alborán Sea; (D) Bürger's (1895) original drawings of *M. dellechiajei*.

*Micrura dellechiajei* (Hubrecht, 1879)  
(Figures 2–4)

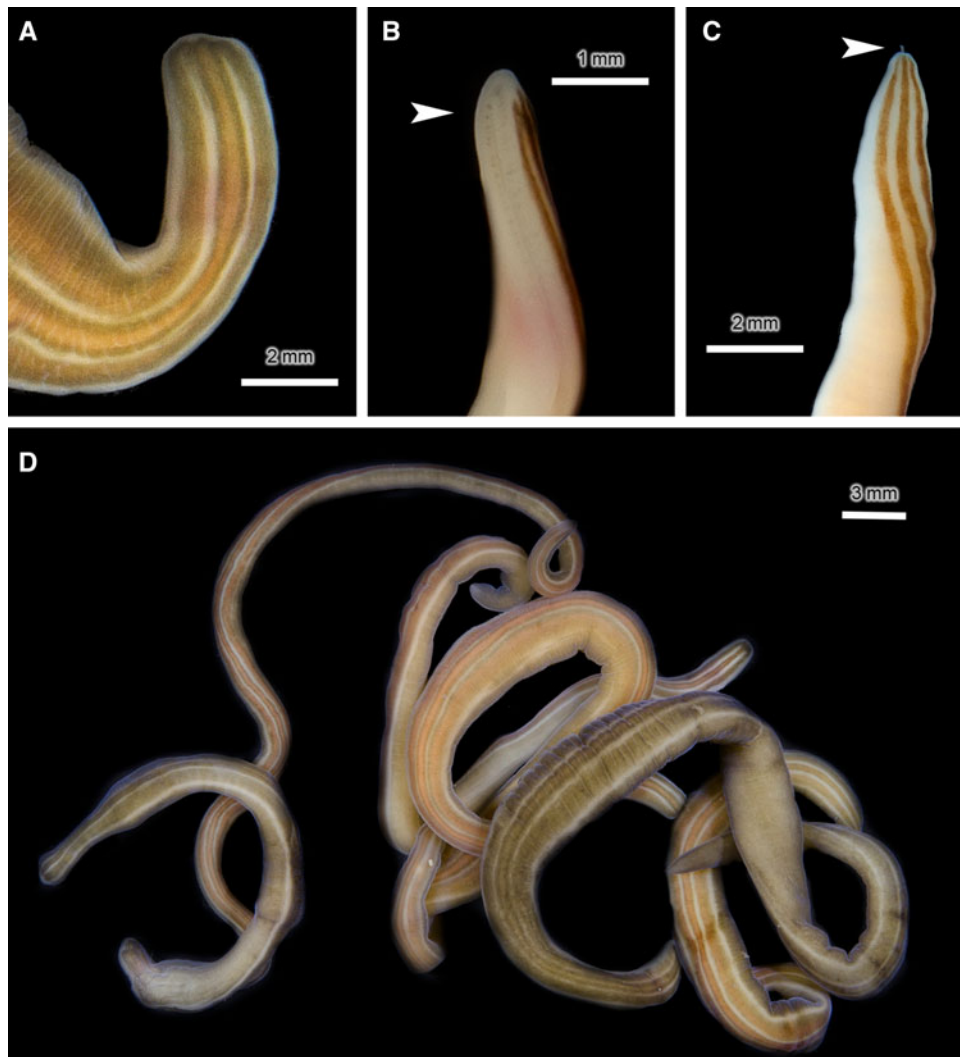
*Cerebratulus bilineatus* (partim.) (Renier, 1804); *Cerebratulus dellechiajei* Hubrecht, 1879; *Meckelia cerebratulus* Diesing, 1850; *Micrura dellachiajei* Bürger, 1895; *Micrura dellechiajei* Bürger, 1892; *Nemertes bilineatus* Örsted (1844); *Nemertes bilineata*; see Gibson (1995).

DESCRIPTION

**External features.** Specimens 4–10 cm long, 3–4 mm wide when fully extended. Body colour varying from very dark to olive or white (Figure 2). Dorsal surface with longitudinal striped pattern alternating symmetrically in colour nuance, with respect to the dorsal midline. Dark specimens with four white dorsal, longitudinal stripes. Olive and white specimens with three dark longitudinal stripes. Some specimens without striped pattern in posterior. Dorsal midline generally orange with dark brown border. Ventral surface pale in all specimens. Head (Figure 3A) ovate, not distinct from rest of body, except for slight constriction at posterior part of

cephalic slits. Tip of head obtuse. Lateral horizontal slits long, open and shallow, reaching behind cerebral ganglia (Figure 3B). When relaxed, body round in cross section, but anterior end dorsoventrally flattened. Posterior end tapering, bearing minute caudal cirrus, sometimes difficult to observe (Figure 3C). Mouth composed of thin but elongate slit, slightly posterior to end of lateral slits. Black ocelli in rows throughout anterior portion of slits (Figure 3B). Gonads visible through body wall in intestinal region, forming rows on both sides of ventral midline. Specimens contract when exposed to light and when disturbed, coiling in a spiral, releasing mucus and usually fragmenting.

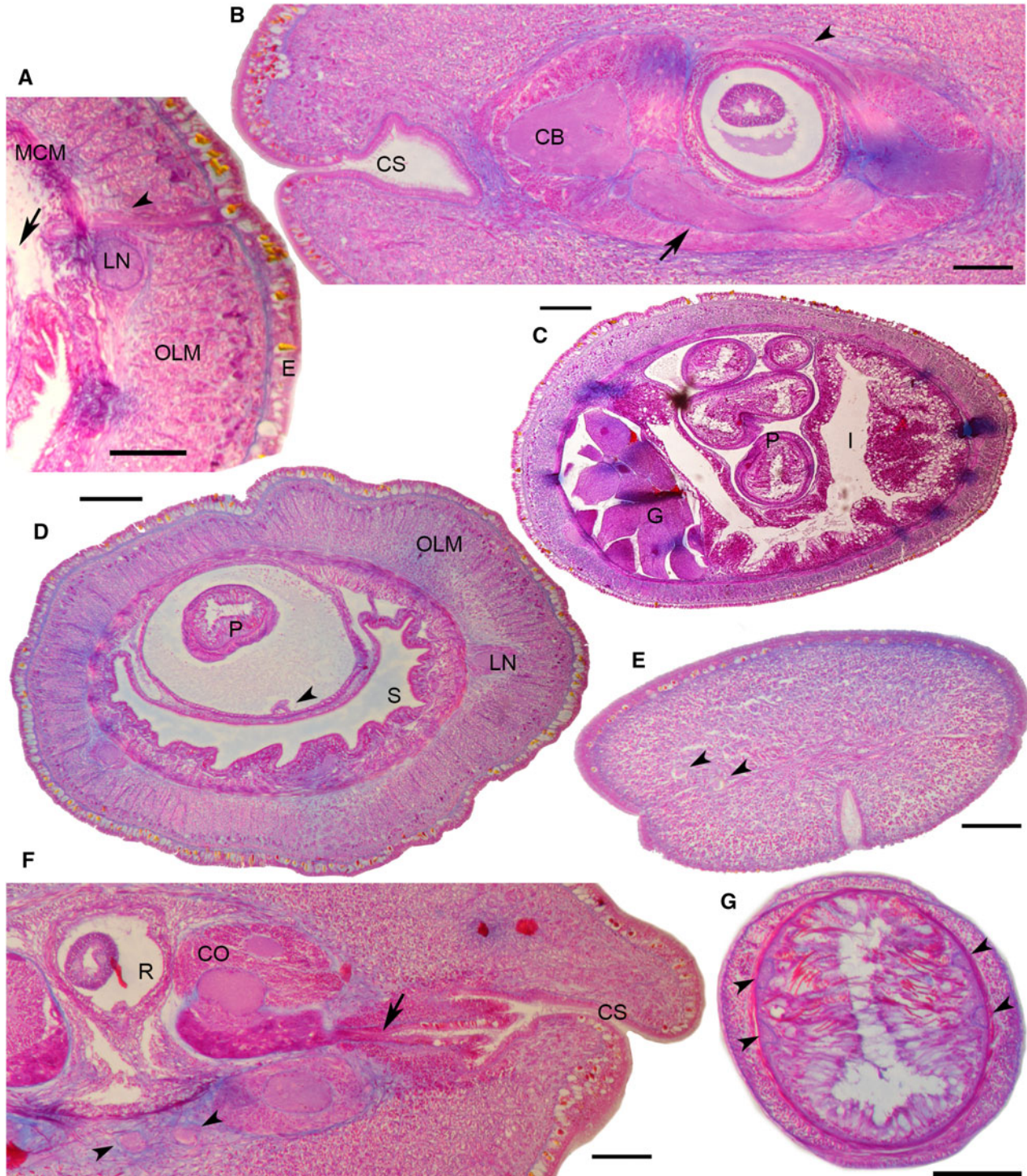
**Body wall, musculature, and parenchyma.** Epidermis ciliated, 35–50  $\mu\text{m}$  thick, dominated by serous goblet cells. Ciliated cells separating goblet cells, predominating in cephalic slits. Mouth with slightly longer epithelial cells. Epidermis resting upon basal lamina. Dermal gland zone as thick (35–50  $\mu\text{m}$ ) as epidermis, placed under the basal lamina. Dermal gland zone consisting of hard-to-detect circular dermal musculature, very thin (15–20  $\mu\text{m}$ , depending on contraction) longitudinal dermal musculature, and with layer of gland



**Fig. 3.** *Micrura dellechiajei* (Hubrecht, 1879). (A) macrophotograph of the head of a specimen in dorsal view, showing the characteristic stripe pattern; (B) macrophotograph of the head of a specimen in lateral view showing the cephalic slits and the ocelli (arrowhead); (C) macrophotograph of the posterior end of a specimen in dorsal-lateral view showing the small caudal cirrus (arrowhead); (D) group of specimens for comparison of the differences between the colour nuances of the individuals.

cells embedded on dermal connective tissue stratum. Well-developed (80–150  $\mu\text{m}$ , depending on contraction) body wall with outer longitudinal musculature (OLM) (Figure 4A, D) below dermal zone. Thin peripheral neural

sheath with nerve cords embedded, present under OLM. Well-developed (25  $\mu\text{m}$ ) middle circular musculature (MCM) followed by an inner longitudinal musculature (ILM), 30  $\mu\text{m}$  thick, present under peripheral neural sheath.



**Fig. 4.** *Micrura dellechiajei* (Hubrecht, 1879). (A) detail of a section through the posterior foregut region showing the lateral nerve cord, the lateral blood lacuna (arrow) and one excretory tubule (arrowhead); (B) detail of a section through the cephalic region showing the dorsal cerebral ganglion with the dorsal (arrowhead) and ventral (arrow) commissures; (C) section through the intestinal region showing the deep lateral intestinal diverticula and the gonads; (D) section through the stomach region showing the moderately convoluted wall; note the rhynchocoelic villus (arrowhead); (E) section through the apical region of the body showing the proboscis pore and a pair of eyes (arrowheads); (F) detail of a section through the cephalic region showing the ciliated canal (arrow) that connects the cephalic slit with the cerebral organ; note the buccal nerves (arrowheads); (G) section of the proboscis showing two pairs of proboscis nerves (arrowheads), the muscle crosses, and the differentiated epithelium. Abbreviations: CB, cerebral ganglion; CS, cephalic slit; CO, cerebral organ; E, epidermis; G, gonad (ovary); I, intestine; LN, lateral nerve cord; MCM, middle circular muscle layer; OLM, outer longitudinal muscle layer; P, proboscis; R, rhynchocoel; S, stomach. Scale bars: 100  $\mu\text{m}$ .

Precerebral cephalic musculature present as framework of outer longitudinal muscle fibres mixed with radial and oblique muscle fibres, with interspersed gland cells. Circular muscles around the rhynchodaeal blood lacuna absent. MCM starts at middle of cerebral region, at point of connection between cerebral organ canals and cephalic slits. Dorsoventral muscles absent in the anterior intestinal region.

*Proboscis apparatus.* Proboscis pore ciliated, opening subterminally and ventrally near tip of head (Figure 4E). Proboscis reaches rhynchodaeum as tubular chamber with ciliated epithelium; globet cells absent. Thin basal lamina surrounding pore lumen. Rhynchodaeum musculature consisting of thin dermal circular layer and discrete longitudinal layer. Rhynchocoel extends for most of body length. Circular muscle fibres of rhynchocoel wall very thick throughout cerebral region; these do not intermingle with muscles of body wall. Rhynchocoel CM and body wall MCM in contact dorsally throughout most of length of body, always with fine coat of ILM as separator. Rhynchocoel with circular musculature in mouth region, intertwined with longitudinal musculature of foregut ventrally and throughout part of length of villus. Villus with spongy appearance. Rhynchocoel wall consisting almost exclusively of thin circular muscular coat in midgut region. Proboscis insertion point just anterior to cerebral ganglia, with two proboscis nerves entering from cephalic nerve region. Proboscis significantly longer than body length, unbranched, possessing five distinct layers: thin endothelial lining, inner longitudinal muscle zone, circular muscle layer, neural plexus with two pairs of proboscis nerves discernable (Figure 4G), and outer glandular epithelium. Epithelium of proboscis differentiated into two regions, one anterior and one posterior; one forming a ridge bearing glandular, cup-shaped structures with apical arrays of small rhabditiform granules, and other with wide folds bearing gland cells lacking such granules. Two muscle crosses distinguishable, running from circular muscle layer to endothelium of proboscis, situated perpendicular to opposing pairs of proboscis nerves.

*Alimentary canal.* Mouth present as ventral slit posterior to cephalic slits, 1 mm long. Interior margin with deep ridges and thick epithelium covered with glandular cells. Foregut wall moderately convoluted, with distinct lining of cilia and microvilli. Subepithelial gland-cell zone ensheathing the stomach at foregut (Figure 4D). Posterior part of foregut marked by change in wall thickness, becoming thinner. Intestinal region with convoluted wall with numerous large spherical glandular cells and bearing deep lateral diverticula for most of length, alternating with gonads in mature specimens (Figure 4C).

*Circulatory system.* Blood lacuna dorsal to the proboscis pore in head region. Rhynchodaeum dividing blood lacuna into two thin-walled cephalic blood vessels. Posterior to proboscis insertion, at level of brain commissures, lateral vessels anastomose to form U-shaped vessel cradling ventral portion of rhynchocoel. In half region of brain, vessel bifurcating, giving rise medially to mid dorsal blood vessel, forming long rhynchocoelic villus, approximately 2–2.5 mm long (Figure 4D). U-shaped vessel widening at level of cerebral organs, progressively becoming bisected by connective tissue. Lateral branches expand forming thin-walled lateral lacunae, bathing posterior portions of cerebral organs, then extending further posteriorly, adjacent to rhynchocoel. Lacunae eventually giving rise to ventrolateral vascular

plexus passing backwards towards margins of buccal cavity and anterior part of foregut, extending to excretory tubules. At end of foregut, upper branches of vascular plexus joining dorsal lacunae to form pair of main dorsolateral blood vessels. In intestinal region, blood supply consisting of paired ventrolateral (running alongside intestine) and dorsolateral (close to rhynchocoel) vessels, and single small mid-dorsal vessel.

*Nervous system.* Brain well developed, with paired dorsal cerebral ganglia larger than ventral ones. Cerebral ganglia pink, visible through dorsal and ventral body wall. Dorsal cerebral commissure shorter and narrower than ventral (25 and 35  $\mu\text{m}$  in diameter, respectively) (Figure 4B). Both dorsal and ventral ganglia enclosed by connective tissue. Outer and inner neurilemma absent around ganglia and lateral nerve cords. Dorsal ganglia bifurcated posteriorly, forming upper and lower neuropil. Ventral ganglia more elongate, extending slightly past dorsal lobes. Ventral ganglia confluent with lateral nerve cords (Figure 4A, D). Neurochords and neurochord cells not distinguished in nervous system. Numerous dorsal cephalic nerves leading forward from anterior borders of brain lobes, innervating cephalic gland region. Two buccal nerves arising from near rear of ventral ganglia (from inner lateral margin), and extending into foregut wall (Figure 4F). Dorsal nerve arising from dorsal commissure, extending posteriorly adjacent to body wall MCM. Peripheral neural sheath encircles the body wall MCM.

*Sensory organs.* Eyes 25–30  $\mu\text{m}$  in diameter, consisting of rudimentary reddish pigment-cup ocelli (Figure 4E). Ocelli ( $N = 20–25$ ) irregularly but longitudinally distributed along lateral cephalic margins, almost in rows, below and above cephalic slits. Two apical sense organs observed on anterior tip of head at level of proboscis pore, appearing as small ciliated pits opening independently, lacking epidermal glandular cells. Pits surrounded by glandular cells interspersed between cephalic musculature fibres in anterior part of head. Lateral horizontal cephalic slits beginning at tip of head and reaching posterior part of cerebral organs, at posterior brain region. Neuroglandular cerebral organs ovoid (maximum 230–250  $\mu\text{m}$  in diameter), with typical lineid structure, attaching to rear of dorsal cerebral ganglia. Cerebral organ canal opening into small chamber of cephalic slit anterior to end (Figure 4F). Epithelium of cerebral organ canal surrounded by nerve cells, bearing cilia, lacking glandular cells. Glandular cells surrounding blind terminus of canal.

*Excretory system.* Well developed, confined to foregut region of body, consisting of several thick-walled longitudinal collecting tubes (20–30  $\mu\text{m}$  in diameter) running close to blood lacunae, extending anteriorly and posteriorly and opening as two lateral nephridiopores near posterior part of foregut region (Figure 4A).

*Reproductive system.* All specimens mature. Only females sectioned, sexes are assumed separate. Gonads distributed in lateral rows on each side of intestine (Figure 4C), distributed between diverticula from dorsal to ventrolateral margins. All ovaries bearing up to 10 or more oocytes (170–230  $\mu\text{m}$  diameter), with nucleus (30–40  $\mu\text{m}$  diameter). No oviducts observed.

#### ECOLOGY

Specimens were found on maërl beds, constituted mainly by *Lithophyllum racemus* (Lamarck) Foslie, 1901,

*Lithothamnion philippii* Foslie, 1897, and *Phymatolithon calcareum* (Pallas) Adey & McKibbin, 1970, and in organically enriched sediments consisting mainly of molluscan shells, with sessile animal communities including species of Porifera (*Spongia* Linnaeus, 1759, *Faciespongia* Burton, 1934 and *Axinella* Schmidt, 1862) and Cnidaria (*Alcyonium* Pallas, 1766, *Epizoanthus* Gray, 1867, *Eunicella* Verrill, 1869, *Paramuricea* Koelliker, 1865 and *Leptogorgia* Milne-Edwards, 1857). Among the more conspicuous and numerous species associated with the samples containing *M. dellechiajei* were the echinoderms *Parastichopus regalis* (Cuvier, 1817) and *Hacelia attenuata* Gray, 1840, as well as the polychaete *Hyalinoecia tubicola* (O.F. Müller, 1776). Specimens were sexually mature between June and September.

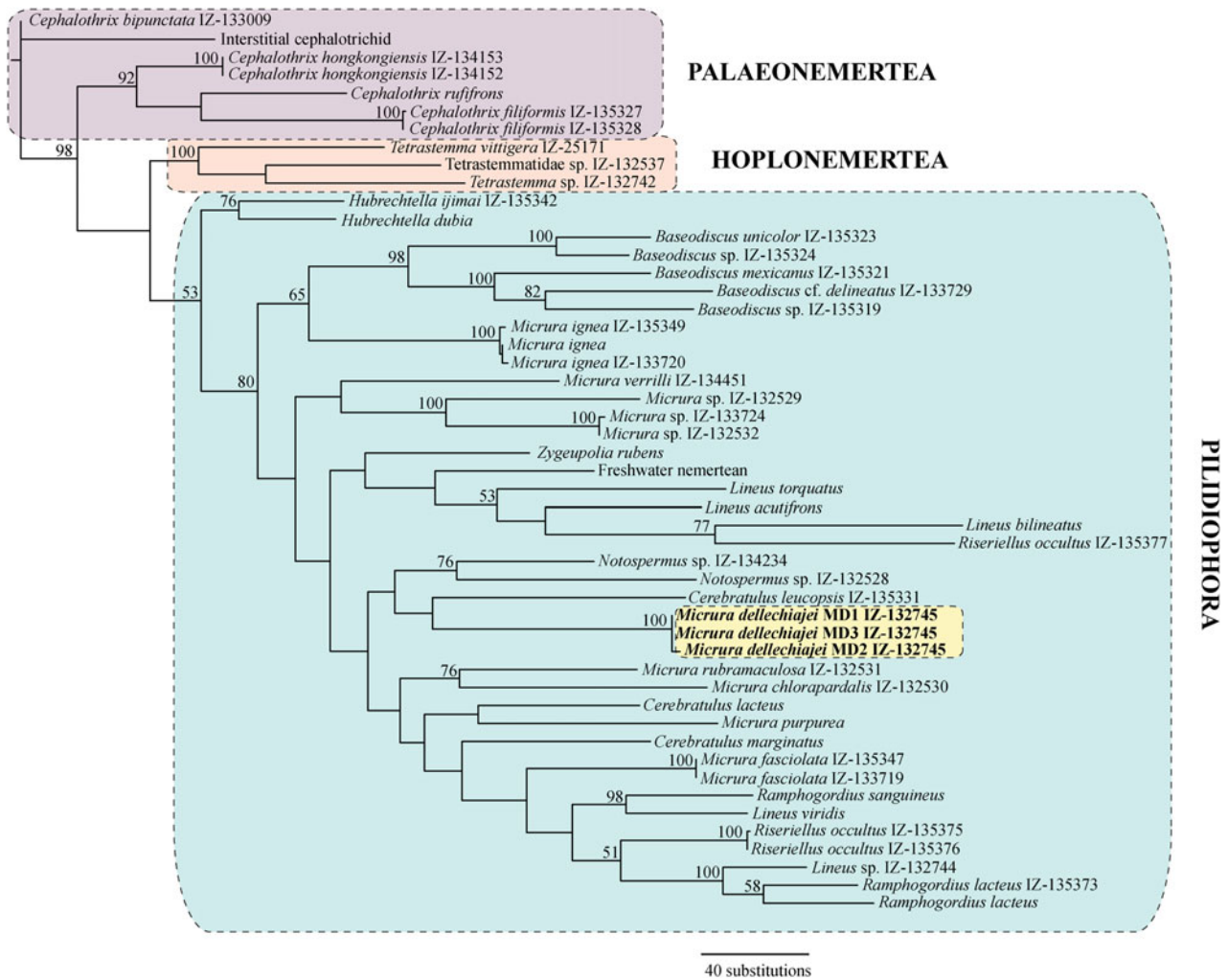
#### PHYLOGENY

The maximum likelihood (ML) analysis resulted in a tree with a final GAMMA-based lnL score of  $-21869.649765$  (Figure 5) and the parsimony analysis (MP) resulted in two equally parsimonious trees at 5392 steps and a compound retention index

and consistency index of 0.508 and 0.255, respectively (the strict consensus tree is presented in Figure 6). Except for some minor rearrangements, mainly within Palaeonemerteae, the ML and MP trees are fully congruent but the ML tree receives slightly higher overall bootstrap support. Both ML and MP trees based on the sequences included in this dataset have previously been discussed at length (Kvist *et al.*, 2014) and, therefore, we here focus mainly on the phylogenetic placement of the included specimens of *Micrura dellechiajei*. In both analyses, the included specimens form a monophyletic group with full support. However, in the ML tree, this clade forms the sister group of two specimens of *Notospermus* Huschke, 1829 (Figure 5), whereas the clade is the sister group of *Cerebratulus leucopsis* (Coe, 1901) in the MP tree. Neither of these positions receive resampling support in the analyses, but the *M. dellechiajei* + *C. leucopsis* clade recovered by the MP analysis is the sister group of the same specimens of *Notospermus*, suggesting that there is some affiliation between these species and *Micrura dellechiajei*. As is typical for a Sanger-based approach (Andrade *et al.*, 2012; Kvist *et al.*,



**Fig. 5.** Best scoring tree from the maximum likelihood analysis of COI + 16S rRNA dataset (lnL =  $-21869.649765$ ). Likelihood bootstrap values  $>50\%$  are shown to the left and above each node. Specimens sequenced for the present study are denoted in bold font and are encircled by a yellow background, and IZ numbers refer to the morphological voucher ID deposited in the Department of Invertebrate Zoology collection of the MCZ.



**Fig. 6.** Strict consensus of two equally parsimonious trees (length: 5392 steps; consistency index: 0.255; retention index: 0.508) produced by TNT for the COI+16S rRNA dataset. Likelihood bootstrap values > 50% are shown to the left and above each node. Specimens sequenced for the present study are denoted in bold font and are encircled by a yellow background, and IZ numbers refer to the morphological voucher ID deposited in the Department of Invertebrate Zoology collection of the MCZ.

2014), most of the genera within Heteronemertea, including *Micrura*, *Cerebratulus* Renier, 1804 and *Lineus* Sowerby, 1806, are found to be paraphyletic. It should be noted that *Lineus bilineatus* (discussed below) is not affiliated closely with *M. dellechiajei*, as it places in a remote region of the tree – these species were probably confused by Delle Chiaje (1829).

## DISCUSSION

The taxonomic history of *Micrura dellechiajei* (Hubrecht, 1879) is complex, much like other nemerteans described in the 19th century. Delle Chiaje (1829, 1841) recorded two species with longitudinal stripe patterns: *Cerebratulus bilineatus* (now *Lineus bilineatus*, Renier, 1804) and *Polia bilineata*, Delle Chiaje, 1841. The latter species was referred to as *Nemertes bilineata* by Örsted (1844), *Meckelia cerebratulus* by Diesing (1850), and *Cerebratulus dellechiajei* by Hubrecht (1879). However, Hubrecht (1879: 214) also mentioned in his description of *C. dellechiajei* that ‘A small caudal appendage similar to that of other species which have formerly been united in the genus *Micrura* is present in most of the

specimens’, which ultimately lead Bürger (1895) to provide an extensive description and finally cementing the placement of the species within the genus *Micrura*, under the currently valid (see Norenburg & Gibson, 2013) specific epithet *Micrura dellechiajei*.

Some of the taxonomic confusion surrounding this species may stem from the stark variation in colour portrayed by its specimens. As evident in Bürger’s (1895) drawings (see Figure 2D of the present paper), as well as in the photographs of the specimens collected in Spanish waters (Figure 2A–C), the colouration ranges from very dark (Menorca Channel), to olive or white (Alborán Sea). In addition, Hubrecht (1879) mentioned that the eyes are disposed in four series on the lateral margins of the head, and recorded a maximum of 120 eyes. Our newly studied Spanish specimens also show this disposition, but the number of eyes is much lower ( $N = 30$ ), which could potentially serve as a further point of taxonomic confusion between *M. dellechiajei* and other heteronemertean taxa. Importantly, nephridia were not observed by Bürger (1895), whereas we here show that a well-developed excretory system is present for this species (Figure 4A). A



caveat of our study may be the fact that neither type material nor material from the type locality were available for study (histological or molecular) and, therefore, there is a risk that Bürger (1895) referred to a different species. However, no specimens of this rare animal have been reported from or near Capri in 120 years, and thus our study can serve to fix a name to well-studied specimens, although we take no nomenclatorial action.

Out of the nearly 500 species of Pilidiophora (Gibson, 1995; Kajihara *et al.*, 2008), four genera contain an overwhelming majority of the diversity, namely *Cerebratulus*, *Lineus*, *Micrura* and *Baseodiscus* Diesing, 1850. Cladistic analyses based only on morphological characters have been ineffective in resolving the phylogenetic relationships among the genera of Pilidiophora, due to an extensive amount of homoplasy (Schwartz & Norenburg, 2001; Schwartz, 2009; Strand *et al.*, 2014). Inadequate descriptions of new species currently presents one of the main challenges to nemertean taxonomy and systematics. Molecular studies on the phylogenetic relationships of nemerteans (Sundberg & Saur, 1998; Sundberg *et al.*, 2001; Tholleson & Norenburg, 2003; Strand *et al.*, 2005; Andrade *et al.*, 2012, 2014; Kvist *et al.*, 2014) have tried to overcome this issue, but have instead found that a staggering amount of genera are non-monophyletic – however, most of these studies agree that nemertean orders (Palaeonemertea, Heteronemertea, Hoplonemertea) and sub-orders (Monostilifera, Polystilifera) are natural (i.e. monophyletic) groups. The trees shown for the present study corroborate this finding (see Figures 5 & 6), even when based solely on mitochondrial genes. The choice of genes was guided by the fact that these loci often show higher rates of nucleotide evolution, as compared with the commonly used 18S rRNA and 28S rRNA, such that they hold the potential to more readily tease out potential cryptic differences between the sequenced specimens (e.g. Chen *et al.*, 2010). In addition, whereas we here use phylogenetic inference to, principally, test the monophyletic status and amount of nucleotide changes between our specimens, *M. dellechiajei* has previously been robustly placed in a phylogenetic context (see Kvist *et al.*, 2014) using six molecular loci – both nuclear and mitochondrial. The single specimen used in that study (MCZ IZ-132745) is also represented herein. Seeing as the three specimens included here show zero nucleotide variation in the relatively fast-evolving COI and 16S rRNA, this warrants the assumption that all three specimens used herein would form a clade, as the sister group to specific members of the genus *Notospermus* (see Figure 3 in Kvist *et al.*, 2014). Importantly, the use of only COI and 16S rRNA by the present study also recovers the species as sister group to a clade of *Notospermus*. Despite the above, it remains that poorly defined genera, such as *Lineus*, *Micrura* and *Cerebratulus* are in dire need of a taxonomic revamp and restructured classification. Currently, the best approach to delimiting and diagnosing nemertean taxa seems to be a combination of morphological characters (including traditional histology) and modern molecular techniques (e.g. Schwartz, 2009; Junoy *et al.*, 2010; Puerta *et al.*, 2010; Kajihara *et al.*, 2011). It is now crucial that future studies use rigorous taxonomic protocols and focus their efforts on resolving the paraphyletic nature of these genera, under the umbrella of molecular phylogenetics and next-generation sequencing, if we are to fully understand the natural history of these enigmatic and charismatic groups.

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## REFERENCES

- Andrade S.C., Montenegro H., Strand M., Schwartz M., Kajihara H., Norenburg J.L., Turbeville J.M., Sundberg P. and Giribet G. (2014) A transcriptomic approach to ribbon worm systematics (Nemertea): resolving the Pilidiophora problem. *Molecular Biology and Evolution* 31, 3206–3215.
- Andrade S.C.S., Strand M., Schwartz M., Chen H-X., Kajihara H., Von Döhren J., Sun S-C., Junoy J., Thiel M., Norenburg J.L., Turbeville J.M., Giribet G. and Sundberg P. (2012) Disentangling ribbon worm relationships: multi-locus analysis supports traditional classification of the phylum Nemertea. *Cladistics* 28, 141–159.
- Bürger O. (1892) Zur Systematik der Nemertinenfauna des Golfs von Neapel. Vorläufige Mittheilung. *Nachrichten von der Königlichen Gesellschaft der Wissenschaften und der Georg-Augusts-Universität zu Göttingen* 5, 137–178.
- Bürger O. (1895) Die Nemertinen des Golfes von Neapel und der angrenzenden Meeres-Abschnitte. *Fauna und Flora des Golfes von Neapel* 22, 1–743.
- Chen H., Strand M., Norenburg J.L., Sun S., Kajihara H., Chernyshev A.V., Maslakova S.A. and Sundberg P. (2010) Statistical parsimony networks and species assemblages in cephalotrichid nemerteans (Nemertea). *PLoS ONE* 5, e12885.
- Delle Chiaje S. (1829) *Memorie sulla storia e notomia degli animali senza vertebre del regno di Napoli*, Vol. 4. Napoli: Società Tipografica.
- Delle Chiaje S. (1841) *Descrizione e notomia degli animali invertebrati della Sicilia citeriore osservati vivi negli anni 1822–1830*, Vol. 3. *Molluschi, Acefali, Bracciopedi, Cirropedi, Crostacei, Anellosoi*. Napoli: C. Batelli e Comp.
- Diesing C.M. (1850) *Systema Helminthum*, Vol. 1. Vienna: W. Braumüller.
- Ehrenberg C.G. (1828–1831) Phytozoa turbellaria Africana et Asiatica in Phytozoorum Tabula IV et V delineata. In Hemprich F.G. and Ehrenberg C.G. (eds) *Symbolae physicae, seu icones et descriptiones corporum naturalium novorum aut minus cognitorum quae ex itineribus per Libyam, Aegyptium, Nubiam, Dongalam, Syriam, Arabim et Habessiniam, pars zoologica II, animalia evertibrata exclusis insectis*. Berlin: Officina Academica, pp. 53–67, pls IV–V (plate published in 1828, text in 1831).

- Gibson R.** (1981) Nemerteans of the Great Barrier Reef 3. Anopla Heteronemertea (Lineidae). *Zoological Journal of the Linnean Society* 71, 171–235.
- Gibson R.** (1995) Nemertean genera and species of the world: an annotated checklist of original names and description citations, synonyms, current taxonomic status, habitats and recorded zoogeographic distribution. *Journal of Natural History* 29, 271–561.
- Goloboff P.A., Farris J.S. and Nixon K.C.** (2008) TNT, a free program for phylogenetic analysis. *Cladistics* 24, 774–786.
- Hubrecht A.A.W.** (1879) The genera of European nemerteans critically revised, with description of several new species. *Notes from the Leyden Museum* 1, 193–232.
- Junoy J., Andrade S.C.S. and Giribet G.** (2010) Phylogenetic placement of a new hoplonemertean species commensal on ascidians. *Invertebrate Systematics* 24, 616–629.
- Kajihara H., Chernyshev A.V., Sun S.-C., Sundberg P. and Crandall F.B.** (2008) Checklist of nemertean genera and species published between 1995 and 2007. *Species Diversity* 13, 245–274.
- Kajihara H., Olympia M., Kobayashi N., Katoh T., Chen H.-X., Strand M. and Sundberg P.** (2011) Systematics and phylogeny of the hoplonemertean genus *Diplomma* (Nemertea) based on molecular and morphological evidence. *Zoological Journal of the Linnean Society* 161, 695–722.
- Katoh K. and Standley D.M.** (2013) MAFFT multiple sequence alignment software version 7: improvements in performance and usability. *Molecular Biology and Evolution* 30, 772–780.
- Kvist S., Laumer C., Junoy J. and Giribet G.** (2014). New insights into the phylogeny, systematics and DNA barcoding of Nemertea. *Invertebrate Zoology* 28, 287–308.
- Lanfear R., Calcott B., Ho S.Y.W. and Guindon S.** (2012) PartitionFinder: combined selection of partitioning schemes and substitution models for phylogenetic analyses. *Molecular Biology and Evolution* 29, 1695–1701.
- Leasi F. and Norenburg J.L.** (2014) The necessity of DNA taxonomy to reveal cryptic diversity and spatial distribution of meiofauna, with a focus on Nemertea. *PLoS ONE* 9, e104385.
- Miller M.A., Pfeiffer W. and Schwartz T.** (2010) Creating the CIPRES Science Gateway for inference of large phylogenetic trees. In Anonymous (ed.) *Proceedings of the Gateway Computing Environments Workshop (GCE)*. New Orleans, LA: Institute of Electrical and Electronics Engineers, pp. 1–8.
- Norenburg J. and Gibson R.** (2013) World Nemertea database. Accessed through: World Register of Marine Species at <http://www.marinespecies.org/aphia.php?p=taxdetails&id=122553>.
- Örsted A.S.** (1844). *Entwurf einer systematischen Eintheilung und speziellen Beschreibung der Plattwürmer auf mikroskopische Untersuchungen gegründet*. Copenhagen: C.A. Reitzel.
- Puerta P., Andrade S.C.S. and Junoy J.** (2010) Redescription of *Lineus acutifrons* (Nemertea: Pilidiophora) and comments on its phylogenetic position. *Journal of Natural History* 44, 2363–2378.
- Renier S.A.** (1804) *Prospetto della classe dei vermi*. An uncompleted work issued in 1804 (see Meneghini's introduction to Renier, 1847), pp. XV–XXVII.
- Schultze M.S.** (1851) *Beiträge zur Naturgeschichte der Turbellarien*. Greifswald: C.A. Koch.
- Schwartz M.L.** (2009) *Untying a Gordian knot of worms: systematics and taxonomy of the Pilidiophora (phylum Nemertea) from multiple data sets*. PhD thesis. George Washington University, Washington, DC, USA.
- Schwartz M.L. and Norenburg J.L.** (2001) Can we infer heteronemertean phylogeny from available morphological data? *Hydrobiologia* 456, 165–174.
- Stamatakis A.** (2006) RAxML-VI-HPC: maximum likelihood-based phylogenetic analyses with thousands of taxa and mixed models. *Bioinformatics* 22, 2688–2690.
- Strand M., Herrera-Bachiller A., Nygren A. and Kínneby T.** (2014) A new nemertean species: what are useful characters for ribbon worm descriptions? *Journal of the Marine Biological Association of the United Kingdom* 94, 317–330.
- Strand M., Hjelmgren A. and Sundberg P.** (2005) Genus *Baseodiscus* (Nemertea: Heteronemertea): molecular identification of a new species in a phylogenetic context. *Journal of Natural History* 39, 3785–3793.
- Strand M. and Sundberg P.** (2011) A DNA-based description of a new nemertean (phylum Nemertea) species. *Marine Biology Research* 7, 63–70.
- Sundberg P. and Saur M.** (1998) Molecular phylogeny of some European heteronemertean (Nemertea) species and the monophyletic status of *Riseriellus*, *Lineus*, and *Micrura*. *Molecular Phylogenetics and Evolution* 10, 271–280.
- Sundberg P. and Strand M.** (2010) Nemertean taxonomy – time to change lane? *Journal of Zoological Systematics and Evolutionary Research* 48, 283–284.
- Sundberg P., Thuroczy Vodoti E. and Strand M.** (2010) DNA barcoding should accompany taxonomy – the case of *Cerebratulus* spp. (Nemertea). *Molecular Ecology Resources* 10, 274–281.
- Sundberg P., Turbeville J.M. and Lindh S.** (2001) Phylogenetic relationships among higher nemertean (Nemertea) taxa inferred from 18S rDNA sequences. *Molecular Phylogenetics and Evolution* 20, 327–334.
- Thollesson M. and Norenburg J.L.** (2003) Ribbon worm relationships: a phylogeny of the phylum Nemertea. *Proceedings of the Royal Society of London B: Biological Sciences* 270, 407–415.

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