

1 **A Cambrian crown annelid reconciles phylogenomics and the fossil record**

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23 **Annelids are among the most disparate animal phyla, encompassing ambush predators,**
24 **suspension feeders and terrestrial earthworms¹. Early annelid evolution remains obscure**
25 **or controversial^{2,3}, partly due to discordance between molecular phylogenies and**
26 **fossils^{2,4}. Cambrian annelid fossils have morphologies indicating epibenthic lifestyles,**
27 **whereas phylogenomics recovers sessile, infaunal and tubicolous taxa as an early**
28 **diverging grade⁵. Magelonidae and Oweniidae (Palaeoannelida¹) are the sister group of**
29 **all other annelids but contrast with Cambrian taxa in both lifestyle and gross**
30 **morphology^{2,6}. We describe a new fossil polychaete, *Dannychaeta tucolus*, from the early**
31 **Cambrian Canglangpu Formation⁷, preserved within delicate, originally organic**
32 **dwelling tubes. The head has a well-defined spade-shaped prostomium with elongate**
33 **ventrolateral palps. The body has a wide, stout thorax and elongate abdomen with**
34 **biramous parapodia with parapodial lamellae. This character combination is shared with**
35 **extant Magelonidae, and phylogenetic analyses recover *Dannychaeta* within**
36 **Palaeoannelida. *Dannychaeta* is the oldest polychaete unambiguously belonging inside**

37 **crown annelids, providing a constraint on the tempo of annelid evolution and revealing**
38 **unrecognised ecological and morphological diversity in ancient annelids.**

39

40 **Systematic Palaeontology**

41 Phylum: Annelida Lamarck, 1809

42 Unranked: Palaeoannelida Weigert & Bleidorn, 2016

43 Family: Magelonidae Cunningham & Ramage, 1888

44 *Dannychaeta tucolus* gen. et sp. nov.

45

46 **Etymology:** *Danny*: referring to Danny Eibye-Jacobsen, for his contributions to our
47 understanding of early annelids; *chaeta* (Latin): bristle; *colus* (Latin): dwelling in; *tubus*
48 (Latin): tube.

49

50 **Holotype:** YKLP 11382 part and counterpart (Fig. 1, Extended Data Fig. 1)

51

52 **Referred material:** YKLP 11383—11402 (Fig. 2-3, Extended Data Figs. 2-6).

53

54 **Locality and Horizon:** Canglangpu Formation, Cambrian Stage 3, Hongjingshao Member
55 (~514 Ma), at southwest of Guanshan reservoir, Chenggong, Kunming, China⁷.

56

57 **Diagnosis for genus:** Elongate, slender polychaetes with organic dwelling tubes. Head with
58 anteriorly tapering spade-shaped prostomium, with paired palps attached ventrolaterally near
59 the mouth. Body heteronomously segmented, with wider thorax containing at least 8
60 chaetigers. Parapodia with lateral lamellae in the posterior part of the abdomen. Abdominal
61 parapodia biramous, unknown in thorax. Capillary chaetae in both rami, occurring in tight
62 parallel bundles.

63

64 **Description:** The holotype (Fig. 1, Extended Data Fig. 1) is incomplete posteriorly
65 (~40 mm long), showing the wider anterior region (thorax; max width 3.9 mm) (Fig. 1a-e) and
66 abdomen (~1.9 mm exclusive of parapodia) (Fig. 1h-i). The number of thoracic segments
67 would consist of ≥ 8 chaetigers, extrapolating from segment spacing (15.5 mm length and 1
68 segment per 1.9 mm). The prostomium is a spade-shaped lobe (Figs 1c-e, 2a-h, Extended Data
69 Figs. 1a-g, 2a-h) and is longer (~4 mm) than wide (~2 mm). The relief of overlapping

70 anatomical features preserved on different planes indicates that the prostomium is dorsal of the
71 palps (Fig. 1d-f, 2b-d). The palps cross over each other in the holotype (Fig. 1d-f), are
72 incompletely preserved anteriorly, but are $\geq 30\%$ the length of the thorax. A specimen in ventral
73 view shows palps inserting ventrolaterally, anterior of a putative burrowing organ (Fig. 2a-d).
74 The prostomium has a faint pair of tapering ridges (Fig. 2d). The gut is preserved as a
75 carbonaceous film (Fig. 3f) that terminates adjacent to palp bases (Fig. 1d-f), indicating palp
76 attachment near the mouth opening.

77 Abdominal parapodia are distinct lobes, projecting $\sim 300 \mu\text{m}$ from the body (Fig. 3c-e).
78 Anterior abdominal chaetae in the holotype are $\sim 500 \mu\text{m}$ long (Fig. 1h). In narrower midbody
79 fragments, chaetae are $\sim 800 \mu\text{m}$ long (e.g. Fig. 3d), suggesting variable chaetation along the
80 body. Abdominal chaetae are in tight fascicles (Figs 1j, 3c-e, Extended Data Fig. 3k) and are
81 most clear using fluorescence microscopy (Fig. 3d). The chaetae are directed slightly anteriorly
82 (Fig. 1i) which is used to orient fragmentary specimens. In two specimens, lateral lamellae
83 occur adjacent to chaetal bundles (Fig. 3g). Lamellae are crescent-shaped and approximately
84 half the body width in length, with a dorsolateral (Extended Data Fig. 1k) to dorsal (Extended
85 Data Fig. 5f-h) placement. As the rami are often parallel, biramous morphology is revealed by
86 subparallel chaetal bundles (Fig. 3d) or rare oblique views (Fig. 3h, Extended Data Fig. 5).
87 Fine details of chaetal morphology are obscure but are consistent with capillaries in the
88 abdomen (e.g. Fig. 3d).

89 The pygidium is never well-preserved, but one specimen has putative pygidial cirri
90 (Extended Data Fig. 5j, k). An ovoid structure between chaetiger one and three is of uncertain
91 identity but resembles a blood lacuna (Fig. 1d-f, Extended Data Figs. 1h-j, 4g).

92 Eight specimens are preserved within a structure that extends beyond the body margin
93 and chaetae (Fig. 3a-f, i-k, Extended Data Figs. 2-4). This is parallel to the body axis,
94 approximately four times the body width (excluding parapodia) (Fig. 3a) with a sharp boundary
95 with the matrix, which is visible using light (Fig. 2f) and fluorescence (Fig. 3h) microscopy
96 and elemental maps for iron (Extended data Fig. 3f). This structure is consistent with a dwelling
97 tube, or tube-lined burrow. The tube has a slightly darker appearance relative to the matrix and
98 lacks identifiable agglutinated bioclasts or grains. Tubes have slight relief (Extended Data Fig.
99 3g) and sometimes have thick walls at their margins (Extended Data Fig. 5c-e), due to
100 compaction. Like the body, the tube contains iron (Fig. 3f) localised to small grains in the
101 matrix that appear bright in SEM backscatter images (Extended Data Fig. 3f). This is consistent
102 with an organic composition, with organic material acting as a substrate for pyrite formation⁸.
103 Specimens preserved in tubes vary from well-preserved with delineated parapodia/chaetae

104 (Fig. 3c-e, Extended Data Fig. 3) to highly effaced, indicating in situ decay⁹ (Extended Data
105 Fig. 4,6). Preservation quality varies along individual specimens (e.g. Extended Data Figs. 3a-
106 d).

107

108 **Discussion**

109

110 Due to low preservation potential⁹, annelid body fossils are rare and distributed discontinuously
111 and unevenly through geological time¹⁰. Diverse fossil polychaetes are known from the early
112 and middle Cambrian deposits (e.g. Sirius Passet^{3,11} and Burgess Shale¹²), but rare from
113 China^{13,14}. Cambrian annelids typically are generalised polychaetes in morphology, with well-
114 developed biramous parapodia (suggesting motility¹⁵), elongate chaetae and a pair of
115 palps^{2,3,6,11}. They cannot be assigned to extant annelid subclades^{6,12} and lack proposed annelid
116 synapomorphies (e.g. the prostomium-peristomium head structure³ and typically pygidial
117 cirri¹²) and are interpreted as stem-group annelids⁶. The lack of Cambrian crown annelids has
118 prompted hypotheses of relatively late crown group diversification, perhaps late
119 Cambrian/Ordovician⁴, when jawed polychaetes become diverse and abundant¹⁶.

120 Both molecular^{1,5,17} and morphological¹⁸ phylogenies have converged on scenarios
121 where annelids evolved from polychaete-like ancestors. However, reconciling fossil and
122 phylogenomic evidence has been challenging^{2,6,12,15}. Molecular phylogenies recover a grade of
123 infaunal, sessile and tube-dwelling taxa as deep branches^{1,5}, including Magelonidae,
124 Oweniidae, Chaetopteriformia and Sipuncula. These groups differ from Cambrian polychaetes
125 in terms of gross morphology and inferred mode of life⁶. A tube-dwelling annelid ancestor has
126 been proposed¹⁹ (although see²⁰), which is contradicted by interpretations of the fossil record¹⁵,
127 and morphological hypotheses regarding the origin of annelid body plan features, e.g.
128 segmentation and parapodia^{18,20}. Fossil specimens interpreted as sipunculans (which
129 phylogenomic studies recover within Annelida^{1,5}) are known from the early Cambrian
130 Chengjiang Lagerstätte²¹ but are rare and poorly known. If correctly interpreted, the oldest
131 fossil crown annelids therefore belong to taxa that have lost most annelid synapomorphies⁶,
132 including segmentation²². Magelonidae and Oweniidae (Palaeoannelida¹) are recovered as the
133 sister group of all other annelids and so have featured prominently in recent discussions of the
134 annelid ancestor^{15,17,19}. These families are unusual among polychaetes, as they lack nuchal
135 organs, possess monociliated epidermal cells and simple nervous systems^{1,17}.

136 *Dannychaeta* is dissimilar in gross morphology to that of previously known Cambrian
137 polychaetes, but shares derived characters with extant Magelonidae. A spade-shaped

138 prostomium with ventrolateral palps (Fig. 2) is characteristic for Magelonidae, which also have
139 a differentiated thorax composed of eight or nine chaetigers^{23,24}. Other well-known Cambrian
140 annelids lack a clearly demarcated head and prostomial lobe³ (but have lateral palps¹⁵),
141 suggesting the typical annelid head structure evolved after the origin of a segmented body,
142 parapodia and palps³. The presence of a differentiated head in *Dannychaeta* is unique among
143 Cambrian annelid fossils, indicating in itself a phylogenetic position proximal to or within the
144 annelid crown group³. The significance of the putative blood lacuna is uncertain, but small ring
145 shaped vessels occur in *Magelona*²³ and a larger lacuna occurs in *Poecilochaetus*²⁵. However,
146 both are placed more anteriorly, nearer the prostomium.

147 Although not widespread (~7% of species²⁴), several magelonid species live in tube-
148 lined burrows²⁶ with an organic/parchment-like composition, some adhered sediment
149 grains/bioclots. These tubes are similar in inferred construction materials and dimensions
150 relative to the body as in *Dannychaeta*. The presence of both obliquely and parallelly oriented
151 specimens (e.g. Extended Data Figs. 2, 4) suggests at least some transport prior to burial.

152 We reconstruct *Dannychaeta* as a sessile, infaunal polychaete that fed in the water
153 column using elongate palps (as in extant Magelonidae^{24,26}; Fig. 4a, b). Our phylogenetic
154 analyses recover *Dannychaeta* in the magelonid stem group (Fig. 4c, Extended Data Figs. 6,
155 7). Parapodial and chaetal morphology differ in some details in *Dannychaeta* and Magelonidae.
156 In extant *Magelona*, the chaetae of the abdomen are hooded hooks, which are generally shorter
157 than the thoracic capillary chaetae and occur in rows²⁴, whereas abdominal chaetae in
158 *Dannychaeta* resemble capillaries held in bundles (Fig. 3c-e). Hooded hooks in certain families
159 share details of ultrastructure and formation²⁷ resulting in a proposed close relationship
160 between Capitellidae, Spionidae and Magelonidae²⁷, which is not supported by
161 phylogenomics¹. The absence of hooded hooks in *Dannychaeta* may therefore provide
162 consistent evidence of convergent chaetal evolution in these families. Parapodial lamellae also
163 differ in some details in *Dannychaeta*, as they are dorsolaterally placed, but occur partially in
164 inter-ramal space in some species in *Magelona*²⁴. *Dannychaeta* is larger than extant magelonids
165 that are typically less than 1 mm wide²⁴, although tubicolous species (e.g. *M. alleni*) achieve
166 the largest widths and share a more robust anterior region (thorax) with *Dannychaeta*²⁴.

167 Other non-pleistoannelid polychaetes also share some characters with *Dannychaeta*.
168 *Spiochaetopterus* has elongate palps, a differentiated anterior region and organic tubes²⁸.
169 However, chaetopterids have ridge-like parapodia (tori) with short, hooked chaetae (uncini²),
170 which are distinct from laterally projecting, lobate parapodia in *Dannychaeta*. The head of
171 *Spiochaetopterus* is neither a distinct anterior lobe nor spade shaped. Nevertheless,

172 chaetopterids branch proximally to Palaeoannelida⁵. A close relationship between
173 *Dannychaeta* and chaetopterids would also suggest the presence of tubicolous, early branching
174 crown group annelids in the early Cambrian.

175 Annelid tube fossils are well-documented¹⁰. Some late Ediacaran and early Cambrian
176 fossil tubes are tentatively assigned to annelids or described as of ‘annelid grade’³⁰, but lack
177 diagnostic features that would allow confident phylogenetic placement. Tube dwelling has
178 evolved several times among extant annelids, and ‘annelid mimicking’ taxa with lophophorate
179 affinities are well known from the early Palaeozoic²⁹, indicating caution should be exercised
180 when assigning fossil tubes. The controversial late Ediacaran tubular fossil *Cloudina* has been
181 tentatively reconstructed as an annelid based on the presence of a tubular gut³⁰.

182 Regardless of the phylogenetic position and fossil record of sipunculans^{4,21,22},
183 *Dannychaeta* confirms crown annelids as minimally early Cambrian in age, revealing early
184 exploration of sessile ecological niches, over a hundred million years prior to other
185 unambiguous examples of tubicolous annelids^{10,29}. *Dannychaeta* reveals that stem-group
186 annelids coexisted with members of the crown group in the early Cambrian and exhibited a
187 diversity of life modes, including epibenthic¹⁵ and sessile forms.

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278

279 **Figure 1 | *Dannychaeta tucolus*, holotype specimen YKLP 11382. a, b**, part and counterpart,
280 direct light. **c**, interpretative drawing of counterpart, colour scheme detailed in **f**. **d**, anterior
281 region of part. **e**, anterior region of counterpart under fluorescent light. Specimen mirrored for
282 easier comparison with part. **f**, interpretative drawing of the anterior, based on both part and
283 counterpart. **g**, Fe map from SEM-EDX analysis. **h, i**, abdominal region, counterpart, direct
284 light (**h**), fluorescent light (**i**). **j**, close-up of parapodium and chaetae. Abbreviations: ch –
285 chaetae, gu – gut, mo – mouth, pl – palp, pa1 – parapodium of segment one, pa8/9 –
286 parapodium of segment eight or nine, pa – parapodium.

287

288 **Figure 2 | *Dannychaeta tucolus*, anterior region. a**, YKLP 11390b, anterior fragment with
289 head and palps. **b**, anterior region of YKLP 11390b, low angle illumination from the northwest.
290 **c**, region as in **b**, fluorescent light. **d**, interpretative drawing outlining anatomical features
291 (colour scheme as in Fig. 1f). **e**, YKLP 11393b specimen preserving anterior region, dwelling
292 tubes and a partial abdomen. **f**, YKLP 11393a, part. **g**, YKLP 11393b, details of anterior region,
293 white arrowheads indicate tube margins. **h**, interpretative drawing of **g**. Abbreviations: pl –
294 palps, pa – parapodium, pro- prostomium, bo – burrowing organ, rd – prostomial ridge.

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298 **Figure 3 | *Dannychaeta tucolus*, morphological details. a-f**, YKLP 11383. **a**, abdominal
299 region within dwelling tube **b**, interpretative drawing of **a** (colours as in Fig 1). Anteriorly,
300 black filled arrowheads indicate the body margin and white filled arrowheads indicate the tube
301 edge. **c**, close up showing the segmented body and parapodia. **d**, close up of chaetal fascicle.
302 **e**, interpretative drawing outlining chaetal fascicles. **f** close up and SEM-EDX maps. **g-h**,
303 YKLP 11389. **g**, close up showing parapodia and lateral lamellae. **h**, close up of YKLP 11384a
304 showing biramous parapodia, chaetae (white arrowhead indicates tube edge). Abbreviations:
305 ch – chaetae, gu – gut, pa – parapodium, plm – parapodial lamella.

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308 **Figure 4 | Reconstruction of *Dannychaeta*. a**, technical drawing showing proportions, body
309 regions and gross anatomy, colour scheme is as in Fig 1f, dorsal view (left) and right lateral
310 view (right). Parapodia in the thorax not observed but inferred shown in grey. **b**, life
311 reconstruction showing *Dannychaeta tucolus* living in buried tubes, artwork by Bob Nicholls.
312 **c**, Bayesian phylogenetic analysis (365 characters, 143 taxa, mki + gamma model) incorporating
313 *Dannychaeta tucolus*. Numbers at nodes are posterior probabilities with scalebar in number of
314 substitutions per site (see Extended Data figure 7 + 8 for full results and additional
315 information).

316

317 **Methods**

318 The specimens were studied, photographed at the Yunnan Key Laboratory for Palaeobiology,
319 Yunnan University (YKLP) and at the University of Exeter, and are deposited in YKLP.
320 Photographs were taken using a Canon EOS 5DSR coupled with a MACRO 100 mm lens, and
321 a Leica DFC7000T linked to a Leica M205 FA fluorescent microscope. The excitation
322 wavelength of GFPL is 480 nm, and the excitation wavelength of RFPL is 546 nm. Images
323 used a gain value of 3.3, saturation value of 52.00, gamma value of 0.92. The external light
324 source of the fluorescence microscope was a LEICA KL 300 LED, used for taking white light
325 images. Scanning electron microscope images were collected using a FEI Quanta 650 FEG
326 using an accelerating voltage of 25kv and a working distance of 12.4 mm. Energy dispersive
327 X-ray (EDX) analyses used an EDAX Pegasus using accelerating voltages of 25-30kv with a
328 working distance of 12.4-13 mm.

329 Phylogenetic analyses were based on a previously published character matrix for
330 annelids and their close relatives⁶ which has been updated successively with the addition of
331 new taxa and fossil data^{14-16,31}. We performed Bayesian analyses using MrBayes 3.2.6 using
332 the mki model with the Lewis correction for the scoring only of informative characters³², with
333 default priors for all parameters (ie.g.e. all trees given equal prior probability). Bayesian
334 analyses with and without topological constraints based on phylogenomic trees were used to
335 investigate if the conflicting topologies recovered from morphological and molecular data
336 impacted the phylogenetic position of *Dannychaeta tucolus*. These constraints were
337 constructed by incorporating results from the most recent¹⁷ and taxon rich^{33,34} phylogenomic
338 analyses of annelids and are outlined in detail in the supporting information. We followed ref¹⁵
339 by excluding *Arkonips* and *Guanshanchaeta* from the character matrix as they contain
340 redundant character scores. For both analyses 100 million generations were requested, with
341 the analysis stopping once the average deviation of split frequencies dropped below 0.01.
342 Convergence was then assessed using ESS (>200) and PSRF (~1.0) values for all model
343 parameters. Parsimony analyses without topological constraints were conducted in TNT 1.5³⁵
344 (courtesy of the Willi Hennig Society), using both equal weights and implied weights with
345 $k=10$. Bremer support and jackknife and bootstrap frequencies from 1000 replications were
346 inferred for equal weights trees and frequency differences were inferred from 1000 replicates
347 of symmetric resampling for the implied weights trees.

348

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363 **Author contributions**

364 LAP, HC, JV and XYM designed the study and interpreted the fossil specimens and their
365 anatomy. HC, DYZ and XGH collected the specimens. HC prepared and photographed all

366 specimens and performed the EDAX elemental analysis. LAP made the figures, performed the
367 phylogenetic analyses and composed the first draft of the manuscript with substantial input
368 from all co-authors.

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375 **ETHICS DECLARATIONS**

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377 **Competing interests.**

378 The authors declare no competing interests.

379

380 **DATA AND CODE AVAILABILITY STATEMENT**

381

382 All data analysed in this paper are available as part of the published paper, the extended data
383 or supplementary information. The phylogenetic dataset and commands and topological
384 constraints necessary to run the MrBayes analyses are included as NEXUS formatted files. The
385 nomenclatural acts in this publication have been registered at ZooBank (LSID:
386 urn:lsid:zoobank.org:pub:5BC89E47-2955-4539-94FD-D400E8C947FB).

387

388 **Extended Data Figure Captions**

389

390 **Extended data Figure 1 | Additional details of holotype specimen. a**, YKLP 11382a entire
391 specimen, including possible posterior region. **b**, possible posterior region from region shown
392 in **a**. **c**, YKLP 11382a, box indicates region used for fluorescence microscopy. **d**, interpretative
393 drawing of specimen shown in **c**. **e**, YKLP 11382b showing regions used in SEM-EDX
394 mapping and fluorescence microscopy. **f**, close up of anterior region showing prostomial lobe
395 and palp attachment. **g**, interpretative drawing of region shown in **f** showing major anatomical
396 features, colour scheme as in Fig 1, and palp attachment inferred also with additional
397 information from the counterpart. **h**, SEM-EDX elemental maps, element names for each map
398 are shown in the bottom right corner. **i**, fluorescence image showing the anterior region of the
399 body in YKLP 11382b including the gut, palps and putative blood lacuna. **j**, fluorescence image

400 of putative blood lacuna in YKLP 11382a. **k**, close up of posterior fragment associated with
401 the holotype with parapodial lamellae. Abbreviations: bl – blood lacuna, ch – chaetae, gu –
402 gut, mo – mouth, pl – palps, plm – parapodial lamella, pro – prostomium.

403

404 **Extended data Figure 2 | Specimen of *Dannychaeta tucolus* YKLP 11393 preserved in**
405 **dwelling tube. a**, YKLP 11393b specimen preserving anterior region, dwelling tube and a
406 partial abdomen. **b**, YKLP 11393a specimen preserving anterior region, dwelling tubes and a
407 partial abdomen. **c**, YKLP 11393b, close up of anterior region (boxed area shown in **a**), white
408 arrowheads indicate the tube margins. **d**, interpretative drawing of region shown in **c**. **e** close
409 up of prostomium and palps of **c**. **f**, YKLP 11393a, close up of anterior region (boxed area
410 shown in **b**), white arrowheads indicate the tube margins. **g**, close up of same region as in **f**,
411 using fluorescence microscope. **h**, interpretative drawing of region shown in **f** and **g**. **i**, poorly
412 preserved abdominal region, from region shown in **a**, direct light. **j**, same region as in **i**, using
413 fluorescence microscope. **k**, poorly preserved abdominal region, from region shown in **b**, direct
414 light. **l**, same region as in **k**, using fluorescence microscope. **m**, thoracic chaetiger showing
415 parapodia and chaetae from region shown in **b**. **n**, thoracic chaetiger showing parapodia and
416 chaetae from region shown in **a**. Abbreviations: ch – chaetae, gu – gut, pa – parapodium, pro –
417 prostomium, pl – palp.

418

419 **Extended data Figure 3 | Additional details of specimen YKLP 11383, preserved inside**
420 **dwelling tube parallel to bedding. a**, YKLP 11383a midbody fragment preserved inside
421 dwelling tube, part. **b**, interpretative drawing of specimen shown in **a**, regions demarcated by
422 black and blue brackets represent decayed and well preserved regions of the body fossil,
423 respectively. **c**, YKLP 11383b, midbody fragment preserved inside dwelling tube, counterpart.
424 **d**, interpretative drawing of specimen shown in **c**. **e**, close up of well preserved region of YKLP
425 11383a in **a**, showing 11 chaetigers preserved inside the dwelling tube. **f**, SEM backscatter
426 image of similar region to EDX maps shown in Figure 3f, showing bright grains associated
427 with the tube and body fossil, arrowheads indicate the left tube margin and pyritised tube wall.
428 **g**, section of fossil shown in **a** photographed under low angle light to indicate the relief of the
429 dwelling tube. **h**, section of fossil shown in **c** photographed under low angle light to indicate
430 the relief of the dwelling tube. **i**, close up of three chaetiger region of region shown in **e**. **j**, same
431 region as in **i**, photographed using fluorescence microscope. **k**, close up of individual

432 parapodium shown in **j**, photographed using fluorescence microscope. Abbreviations: ch –
433 chaetae, gu – gut, pa – parapodium.

434

435 **Extended data Figure 4 | Specimens YKLP 11384a, YKLP 11387 and YKLP 11401**
436 **showing effaced specimens preserved in dwelling tubes. a**, YKLP 11384a, anterior fragment
437 preserving thorax and abdomen preserved inside dwelling tube. **b**, YKLP 11387a, anterior
438 fragment preserving thorax and abdomen. **c**, YKLP 11387b, anterior fragment preserving
439 thorax and abdomen. **d**, close up of abdominal chaetigers in YKLP 11387a, from region shown
440 in **b**, white filled arrows indicate the tube margins. **e**, same region as in **d** photographed using
441 fluorescence microscope. **f**, YKLP 11401, effaced specimen preserved in dwelling tube
442 preserving the putative blood lacuna. **g**, close up of region shown in **f**, showing gut and possible
443 blood lacuna. Brackets on **a**, **b**, **c** and **f** indicate the position of the thoracic region.
444 Abbreviations: bl – blood lacuna, ch – chaetae, gu – gut.

445

446 **Extended data Figure 5 | Additional details of specimen YKLP 11389 showing details of**
447 **parapodia, parapodial lamellae and the posterior region. a**, YKLP 11389b, counterpart,
448 posterior fragment preserving parapodia and chaetae. **b**, YKLP 11389a, part showing
449 preservation of lateral parapodial lamellae. **c**, close up of five chaetigers from region shown in
450 **a**. **d**, close up of chaetiger from region shown in **c**. **e**, close up of chaetiger in **d**. **f**, close up of
451 chaetiger preserving parapodial lamellae from region shown in **b**. **g**, chaetigers preserving
452 parapodial lamellae from region shown in **b**. **h**, same region as in **g** showing parapodial
453 lamellae, photographed using fluorescence microscope. **i**, posterior region as shown in **a**, with
454 putative pygidial cirri. **j**, close up of putative pygidial cirri in **i**. **k**, same region as in **j**,
455 photographed with fluorescence microscope. Abbreviations: ch – chaetae, gu – gut, plm –
456 parapodial lamella, pyc – pygidial cirri.

457

458 **Extended data Figure 6 | Specimen YKLP 11384, a decayed specimen preserved in dwelling**
459 **tube. a**, YKLP 11384a, whole specimen in dwelling tube, white arrowheads indicate the tube
460 margin. **b**, YKLP 11384b, whole specimen in dwelling tube, white arrowheads indicate the
461 tube margin. **c**, details of YKLP 11384b, showing putative blood lacuna and gut, white and
462 black arrowheads indicate the tube and body margins respectively. **d**, same view as in **c**, but
463 imaged using fluorescence microscopy, note the thick appearance of the tube margin. **e**, detail
464 of YKLP 11384a, showing preservation of the tube wall, indicated by white arrowheads.

465

466

467 **Extended data Figure 7 | Full results of Bayesian phylogenetic analyses.** **a**, full results of
468 unconstrained analysis under the mki + gamma model, topology is a majority rule consensus
469 tree and the scale bar is in units of expected number of substitutions per site, numbers at nodes
470 are posterior probabilities. **b**, full results of analysis with constraints from phylogenomics under
471 the mki + gamma model, topology is a majority rule consensus tree and the scale bar is in units
472 of expected number of substitutions per site, numbers at nodes are posterior probabilities.

473

474 **Extended data Figure 8 | Results of parsimony analyses under equal and implied**
475 **weighting.** **a**, Strict consensus tree (length 1071) of parsimony analysis without topological
476 constraints, numbers at nodes are support from bootstrapping, jackknifing and Bremer decay. **b**,
477 strict consensus of trees inferred using implied weighting at k=10 (tree score 46.14190),
478 numbers at nodes are relative frequencies from symmetric resampling.

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