

1 *Protomelission* is an early dasyclad alga and not a Cambrian bryozoan

2 Jie Yang^{1*}, Tian Lan^{2*}, Xi-guang Zhang¹✉, Martin R. Smith³✉

3 1. Institute of Palaeontology, Yunnan University, Chenggong, Kunming 650500, China

4 2. College of Resources and Environmental Engineering, Guizhou University, Guiyang 550003, China

5 3. Department of Earth Sciences, Durham University, Durham, DH1 3LE, UK

6 * Joint first authors

7 ✉ Authors for correspondence: XGZ, xgzhang@ynu.edu.cn; MRS,

8 martin.smith@durham.ac.uk

9 **The animal phyla and their concomitant body plans trace their origins to a singular**
10 **burst of evolution in the Cambrian period, over 500 million years ago¹. Phylum**
11 **Bryozoa, the colonial “moss animals”, have been the exception: convincing skeletons of**
12 **this biomineralizing clade have been curiously absent from Cambrian strata, in part**
13 **because potential bryozoan fossils are difficult to distinguish from the modular**
14 **skeletons of other animal and algal groups^{2,3}. At present, the strongest candidate⁴ is the**
15 **phosphatic microfossil *Protomelission*⁵. Here, we describe exceptionally preserved non-**
16 **mineralized anatomy in *Protomelission*-like macrofossils from the Xiaoshiba**
17 **Lagerstätte⁶. Taken alongside the detailed skeletal construction and the potential**
18 **taphonomic origin of ‘zooid apertures’, we consider that *Protomelission* is better**
19 **interpreted as the earliest dasycladalean green alga – emphasizing the ecological role of**
20 **benthic photosynthesizers in early Cambrian communities. Under this interpretation,**
21 ***Protomelission* cannot inform the origins of the bryozoan body plan; despite a growing**
22 **number of promising candidates^{7–9}, there remain no unequivocal bryozoans of**
23 **Cambrian age.**

24 **Introduction**

25 The phyla – the principal subdivisions of the animal kingdom – had diverged by the start of
26 the Cambrian “explosion”¹⁰. The origins of their distinctive body plans can be resolved only
27 by reference to the fossil record, as surviving lineages are characterized by rapid early
28 innovation¹¹ overprinted by half a billion years of subsequent evolution and extinction. The
29 recognition of “extinct phyla” as offshoots of surviving animal lineages^{12,13} has uncovered
30 unexpected connections between disparate animal phyla^{14,15}, and revealed the otherwise
31 unpredictable morphologies that characterize ancestors of modern animal groups^{16,17} –
32 cementing the role of the fossil record as a reliable eyewitness to the origins of animal
33 diversity¹⁸.

34 However, despite the near-simultaneous appearance of other “shelly” bilaterian phyla
35 (Euarthropoda, Mollusca, Brachiopoda and Echinodermata) close to the start of Cambrian
36 Stage 3^{12,19,20}, Phylum Bryozoa lacks convincing representatives in the Cambrian fossil
37 record. Proposed Cambrian bryozoans are putative⁸, disputed^{2,9} or discredited^{7,21}. Presently,
38 the most compelling candidate is the Stage 3– 4 microfossil *Protomelission*^{4,5}, whose
39 reinterpretation as a bryozoan⁴ would complete the Cambrian roster of mineralizing body
40 plans.

41 *Protomelission gatehousei* is a millimetre-scale organism comprising secondarily
42 phosphatized cataphract modules, organized around a central cavity to form a holdfast and an
43 erect axis. The axis is preserved as a bifacial structure, possibly reflecting the early
44 compaction of an originally hollow club-shaped organism. Modules are arranged at 25° to the
45 central cavity, and comprise thin-walled chambers with a small basal aperture on their
46 internal surface; and a large, external distal aperture, whose margin is typically irregular due
47 to breakage, leaving its original shape and size speculative^{4,5} – if indeed an aperture was even
48 present before abrasion. The shape, size, and distal opening of modules have been taken to

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49 indicate a position in the bryozoan total group⁴, but these features are not necessarily
50 sufficient to render this the only, or even the most likely, possible affinity. In fact, the thinness
51 of the walls and the irregular folding of peripheral modules are difficult to square with a
52 bryozoan interpretation^{5,22}.

53 Here we compare phosphatized *Protomelission* microfossils previously described
54 from Wirrealpa (five specimens) and Xiaoyangba (one specimen)^{4,5} with twelve probable
55 *Protomelission* macrofossils from the Xiaoshiba biota (Cambrian Stage 3).

56 **Results**

57 Our new specimens comprise an originally clavate thallus (Fig. 1a, d, e) and a basal holdfast,
58 which is in some cases attached to an animal shell (Fig. 1d, e, h). Compaction of the thallus is
59 reflected by ridges and wrinkling of its surface (Fig. 1a). Its preservation across two separate
60 surfaces within the matrix (Figs 1b, d, 2a, f, g) reveals the internal and external surfaces of
61 the thallus in juxtaposition. One specimen displays budding or branching (Fig. 1b).

62 The thallus comprises quincuncially arranged, slightly rounded modules, each around
63 250 μm long, 200 μm wide (Figs 1–2) and 1 000 μm deep (Fig. 2e). Reflecting their close
64 packing, modules are broadly hexagonal or rhombic in outline, and are slightly elongated
65 parallel to the axis of the thallus. Each module is enclosed by a thin but robust layer
66 associated with elevated concentrations of iron and phosphorous (Fig. 2d; Extended Data Fig.
67 1), which opens internally with a small (5–8 μm) proximal aperture (Figs 1c, 2i) and
68 externally at a large distal aperture whose irregular outline and size reflects partial survival of
69 the original surface layer (Figs 2a–d, 3b, d–e, h; Extended Data Fig. 1a).

70 As with the phosphatized Wirrealpa specimens (Fig. 3c, g, j), the nature of the
71 external aperture varies between modules and between specimens, in a manner consistent
72 with abrasion of a delicate external membrane: in many Xiaoshiba (Fig. 3b, i) and Wirrealpa

73 (Fig. 3c, j) specimens, the majority of the external membrane is absent. Even where the
74 external membrane is at its most complete, the irregular shape, size and position of the sub-
75 elliptical apertures (Fig. 3e, f) do not correspond with the consistent regularity expected of a
76 bryozoan colony, and a taphonomic origin is difficult to dismiss.

77 On the basis of the close correspondence in construction, size, and arrangement of the
78 modules with those of the *Wirrealpa* microfossils (Fig. 3a–c, e–j; Extended Data Table 1), we
79 tentatively assign our material to *Protomelission?* sp. – though we stress that any comparison
80 must account for the complementary preservational pathways of the two deposits. The
81 *Wirrealpa* material was entombed in lime-rich sediment (since removed by acid maceration)
82 and secondarily replaced, soon after death, by phosphate – protecting the material from
83 further compaction^{4,23}. Phosphatized deposits tend to contain early developmental stages or
84 fragments of larger organisms²³, reflecting a taphonomic bias towards small (< 5 mm)
85 specimens. In contrast, Xiaoshiba-style preservation entails rapid burial of relatively
86 complete organisms, followed by substantial post-burial compaction. Fossils are exposed by
87 splitting mudstones along planes of weakness, which typically correspond to internal or
88 external fossil surfaces²⁴.

89 These preservational differences account for certain differences between the material.
90 Firstly, the *Wirrealpa* material presents an external view of (sometimes abraded) specimens,
91 whereas the planes of fracture within Xiaoshiba material variously coincide with internal and
92 external surfaces of modules, or divide modules internally. Secondly, the size difference
93 between the deposits (108–160 × 32–43 mm in Xiaoshiba, 1.8–2.2 × 1.0–1.5 mm in
94 *Wirrealpa*) likely reflects the preservation of larger or more complete specimens in Xiaoshiba
95 – just as palaeoscolecid worms are represented by sub-millimetric fragments of cuticle in
96 *Wirrealpa*⁵ and complete centimetre-scale specimens in Xiaoshiba²⁵. The small size of
97 *Wirrealpa* fragments precludes the recognition of macroscopic features, such as budding.

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98 Most significantly, the best-preserved of our compression fossils additionally preserve
99 non-skeletal tissue, which is never evident in *Wirrealpa*. Unornamented sub-conical flanges,
100 around 500 μm in length, emerge orthogonally from each module (Figs 1b, f; 2a–e). These
101 structures were originally labile, denoted by depletion in iron and phosphorous (Fig. 2d,
102 Extended Data Fig. 1). They are best displayed at the lateral margins of specimens, where
103 they are preserved parallel to the plane of bedding; because they are oriented perpendicular to
104 the plane of splitting in the medial thallus, their full outline is not visible, though the chemical
105 signature of their bases is evident across the thallus (Extended Data Fig. 1b, c). (Analogous
106 preservation is seen in the centripetal sclerites of certain priapulid worms, which can be
107 difficult to distinguish from somatic tissue except at the edges of fossils – e.g. fig. 8 in
108 ref. 14.)

109 **Discussion**

110 *Protomelission?* flanges lack tentacles or other complex structures, which would be expected
111 if they corresponded to the soft bodies of a colonial metazoan. Regular polygonal modules
112 also characterize the inner surface of certain archaeocyath-like sponges²⁶, but the largely
113 enclosed nature of *Protomelission* modules and the presence of external flanges are difficult
114 to reconcile with an archaeocyath construction. Rather, *Protomelission?* sp. is more
115 convincingly interpreted as the oldest²¹ dasyclad green alga (Extended Data Table 1); a stem-
116 group position is implied by molecular clock estimates (with c. 95% posterior probability) of
117 a post-Cambrian crown-group radiation²⁷.

118 We interpret the phosphatized *Wirrealpa* specimens as representing replacements of
119 skeletal material. Zhang et al.⁴ interpret a wrinkled microstructure as indicating an originally
120 non-mineralized composition, but in view of the absence of other non-mineralized elements
121 in the *Wirrealpa* assemblage⁵, we are reluctant to rule out the possibility that the phosphate

122 replaces original calcium carbonate, with the idiosyncratic microstructure reflecting the
123 deposition of carbonate within the algal cell wall or an encompassing mucilage layer, as in
124 extant dasycladaleans²⁸.

125 Palaeozoic dasycladaleans such as the cyclocrinids preserve tessellating modules
126 corresponding to a robust, often calcified layer that surrounds lateral branches that emerge
127 from a central cavity. Each such module exhibits a small basal aperture opening into the
128 central cavity, and a larger distal opening from which a non-mineralized flange emerges^{3,29},
129 corresponding to the structures seen in our compression fossils (Figs 1–2). As with many
130 fossil dasyclads³⁰, gametophores are not preserved in *Protomelission* – potentially signifying
131 that the fossil material represents vegetative stages, or an endospore taxon.

132 A subset of phosphatic microfossils assigned to Cambroclavida³¹ – a problematic
133 early–mid Cambrian group restricted to open marine environments within the photic zone³² –
134 also comprise articulated arrays of hollow modules with taphonomically abraded distal
135 openings (Fig. 3d, k, l; Extended Data Table 1). In certain taxa, these openings correspond to
136 the base of a compositionally distinct conical flange^{33,34}. The parallel morphology hints that
137 at least some cambroclaves may represent more heavily skeletonized relatives of
138 *Protomelission*-like dasycladaleans, potentially extending the record of these algae to span
139 the stratigraphic gaps between *Protomelission* and other putative³⁵ Cambrian dasyclads^{3,21}.

140 Taken together, this indicates that benthic algae had a greater role in Cambrian
141 communities than previously appreciated. Whereas eukaryotic macroalgae have been a
142 significant component of benthic ecosystems since at least the Ediacaran³⁶, they account for a
143 small proportion of the biomass and diversity in most Burgess Shale-type settings^{37–42}
144 (particularly after excluding ‘algal’ taxa now assigned to cyanobacteria or
145 hemichordates^{43,44}). Whereas photosynthetic microorganisms exploited changing ocean
146 chemistry to secrete carbonates from the base of the Cambrian⁴⁵, *Protomelission* potentially

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147 indicates an onset of mineralization in macroalgae broadly coincident with the proliferation
148 of animal body plans, and the concomitant ecosystem shifts, at the opening of Cambrian
149 Age 3.

150 Whether or not our compression fossil material ultimately proves synonymous with
151 secondarily phosphatized *P. gatehousei*^{4,5}, it demonstrates that the suite of features used to
152 assign *Protomelission* to Bryozoa do not exclusively characterize that group (Extended Data
153 Table 1): a bryozoan affinity can be considered tentative at best. A growing number of
154 Cambrian fossils, including *Protomelission*, the Harkless bryomorph⁸, *Pywackia*⁹, the
155 overlooked cambroclave *Deltaclavus*³¹, and the disfavoured^{2,21} *Cambroporella* and
156 *Archaeotrypa*⁷, display characteristics that might be reconciled with a bryozoan affinity – but
157 on the basis of presently available material, no taxon can be interpreted with sufficient
158 certainty to document a pre-Ordovician origin of Bryozoa. Taken alongside the early origin of
159 poriferan-like⁴⁶ and cnidarian-like fossils^{47,48} and the continued absence of multiple extant
160 phyla across ever more modes of exceptional fossil preservation^{49–51}, the origin of bilaterian
161 body plans need not necessarily be compressed into a unique period in the heat of the
162 Cambrian ‘explosion’.

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290 **Figure legends**

291 **Figure 1 | *Protomelission*-like dasyclad algae from Xiaoshiba. a–c, YKLP 12446, large**
292 **thallus with bud, showing inner and outer surfaces (at bottom and top of panel, respectively);**
293 **b, enlargement of area B in a, showing budding structure with lateral flanges (white arrows);**

294 **c**, enlargement of area *C* in **a**; inner surface of thallus, showing regular arrangement of
295 individual modules with regularly situated fine holes (arrows); **d**, YKLP 12447, showing
296 clavate form and attachment to brachiopod shell; **e**, YKLP 12448, complete specimen
297 attached to indeterminate shell fragment (orange arrow), showing lateral flanges (white
298 arrows); **f**, YKLP 12449, showing inner surface of thallus, with lateral profile of marginal
299 modules (white arrows), and flanges extending centripetally into matrix (orange arrows); **g**,
300 YKLP 12447; enlargement of area *G* in **d**, showing inner surface of thallus with regularly
301 situated fine holes (arrows); **h**, YKLP 12450, two specimens anchored by holdfasts (arrow) to
302 a brachiopod shell.

303

304 **Figure 2 | Preservation of *Protomelission?* sp. a–e**, YKLP 12451: **a**, entire specimen,
305 showing outer surface (on right), poorly preserved inner surface (on left), and lateral section
306 (box *E*); **b**, enlargement of area *B* in **a**, showing lateral aspect of modules and flanges
307 (arrowed); **c**, fluorescence photograph of area *B* in **a**, showing flanges; **d**, energy dispersive
308 X-ray spectrograph showing iron abundance in area *B* in **a**: abundance elevated in module
309 walls, and depressed relative to the matrix in flanges; **e**, enlargement of area *E* in **a**, showing
310 lateral aspect of modules, and flanges (arrows); **f–g**, YKLP 12452, external surface (majority
311 of **f**) juxtaposed with internal surface (**g**); in which fine holes occur at bases of each module
312 (arrows); **h–i**, YKLP 12453, displaying internal surface of thallus.

313

314 **Figure 3 | Comparison of the Xiaoshiba *Protomelission?* sp. with *Wirrealpa P.***
315 ***gatehousei* and the Cambroclavid *Deltaclavus graneus*. a–b**, Xiaoshiba *Protomelission?*
316 YKLP 12446, 12451, showing cataphract module arrangement; **a**, outer membrane largely
317 intact; **b**, taphonomically expanded apertures perforate the partially preserved membrane
318 (arrows). **c**, *Wirrealpa P. gatehousei* SADME 10470, showing equivalent preservational

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319 pattern. **d**, *Deltaclavus* IAGS-BC-88-30181, showing cataphract module arrangement;
320 taphonomically expanded apertures perforate the partially preserved membrane. **e–f**
321 Xiaoshiba *Protomelission?* YKLP 12454: **e**, specimen with most complete preservation of
322 external membrane; **f**, enlargement of area *F* in **e** shows irregularly located apertures (white
323 arrows) in thin outer membrane of modules, with inconsistent size and shape, and similar
324 damage lying across the boundaries of modules (orange arrows), presumably of taphonomic
325 origin. **g**, *Wirrealpa P. gatehousei* SADME 10470-3; and **h**, Xiaoshiba *Protomelission?* YKLP
326 12452: both showing flat inner surface of modules with small basal aperture (arrows). In **g**,
327 modules of the facing surface (white arrows) do not leave an impression on the inner surface
328 of opposing modules, consistent with their separation, as in **h** (white arrows), by a central
329 cavity in life, prior to compaction. **i**, Xiaoshiba *Protomelission?* YKLP 12451; enlargement of
330 box 3I in Fig. 2a; and **j**, *Wirrealpa P. gatehousei* SADME 10470-2, both showing three-
331 dimensional relief of outer surface, with apertures (asterisks) enclosed by partially preserved
332 membrane (arrows). **k**, *Deltaclavus* IAGS-BC-88-30181, lateral view showing lateral profile
333 of modules. **l**, *Deltaclavus* IAGS-BC-88-30178, showing three-dimensional relief of outer
334 surface, with apertures (asterisks) enclosed by bounding ridge (arrows). Images of
335 *Deltaclavus* courtesy S. Conway Morris; images of *Wirrealpa Protomelission* reproduced
336 from ref. 4 under license (<https://creativecommons.org/licenses/by/4.0/>).

337 **Methods**

338 Twelve specimens from the Xiaoshiba biota were collected from outcrop of the Hongjingshao
339 Formation, near Kunming, dated to the *Yunnanocephalus–Chengjiangaspis–Hongshiyanaspis*
340 biozone, lower Canglangpuan Stage, Cambrian Series 2, Stage 3)^{6,52}. Specimens were imaged
341 under visible light under a LEICA M205-C stereomicroscope; and with fluorescence
342 photography using a LEICA DFC 7000T monochrome digital camera attached to a LEICA

343 M205 FA fluorescence stereomicroscope under visible light. Backscatter electron microscopy
344 and energy-dispersive X-ray spectroscopy were conducted using a FEI Quanta 650 scanning
345 electron microscope under low vacuum and a 30 kV accelerating voltage.

346 **Statistics and Reproducibility.** Photographs and micrographs have been selected
347 after examination of all available material in order to best represent the features under
348 description.

349 **Data availability.** Specimens are accessioned at the Institute of Palaeontology,
350 Yunnan University (YKLP12436–42).

351 **Code availability.** No custom software or code was used in this study.

352 **Methods references**

353 52. Hou, J.-B., Yang, J., Zhang, X.-G., Hughes, N. C. & Lan, T. [Trilobite-based](#)
354 [biostratigraphy of the Xiaoshiba Lagerstätte](#). *Fossils and Strata* **64**, 173–191 (2019).

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360 **Author contributions.** X.G.Z., M.R.S. and T.L. designed the project. X.G.Z. M.R.S.
361 and T.L. carried out the morphological and anatomical analyses, and contributed to the final
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363 approved the final manuscript. M.R.S. and X.G.Z. wrote the manuscript with input from other
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366 **Competing interests.** The authors declare no competing interests.

367 **Additional information.** Correspondence and requests for materials should be
368 addressed to XGZ or MRS. Reprints and permissions information is available at
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370

371 **Extended Data Figure 1 | Elemental abundances in *Protomelission?* sp.** Reflected light,
372 backscatter electron, and energy-dispersive X-ray spectroscopy images of **a**, central thallus of
373 YKLP 12451, corresponding to region 3I in Fig. 2a; **b–c**, flanges at thallus margin in **b**,
374 YKLP 12451, corresponding to region B of Fig. 2a; **c**, YKLP 12446, corresponding to region
375 of Fig. 1b.

376

377 **Extended Data Table 1 | Anatomical comparison between *Protomelission* and possible**
378 **relatives.** Comparison of pertinent morphological features in *Protomelission gatehousei* from
379 Wirrealpa and Xiaoyangba^{4,5}, *Protomelission?* sp. from Xiaoshiba (this study), the
380 cambroclave *Deltaclavus*³¹, and representatives of the dasycladalean algae and Bryozoa. Bold
381 type emphasizes differences from the bryozoan body plan.