

Herpetogaster collinsi from the Cambrian of China elucidates the dispersal and palaeogeographic distribution of early deuterostomes

Xianfeng Yang Corresp., 1, 2, 3, Julien Kimmig Corresp., 4, 5, James D Schiffbauer 6, 7, Shanchi Peng 3

Corresponding Authors: Xianfeng Yang, Julien Kimmig Email address: yangxf@ynu.edu.cn, jkimmig@psu.edu

The Cambrian Radiation represents one of the largest diversification events in Earth history. While the resulting taxonomic diversity is exceptional, relatively few of these novel species can be traced outside the boundaries of a single palaeocontinent. Those species with cosmopolitan distributions were likely active swimmers, presenting opportunity and means to conquer new areas, but this would not have been the case for sessile organisms. Herpetogaster is a genus of sessile, stalked, filter-feeding deuterostomes with two species, H. collinsi and H. haiyanensis, known from the lower to middle Cambrian (Series 2-Miaolingian, Stage 3-Wuliuan) of Laurentia and Gondwana. Here, we expand the distribution of H. collinsi to Gondwana with newly discovered specimens from the Balang Formation of Hunan, China. This discovery raises questions on the origin of the genus and how sessile organisms were able to disperse over such a broad distance in the lower Cambrian.

¹ Yunnan Key Laboratory for Palaeobiology, Yunnan University, Kunming, China

² MEC International Joint Laboratory for Palaeobiology and Palaeoenvironment, Yunnan University, Kunming, China

³ State Key Laboratory of Palaeobiology and Stratigraphy, Chinese Academy of Sciences, Nanjing, China

⁴ Paläontologie und Evolutionsforschung, Abteilung Geowissenschaften, Staatliches Museum für Naturkunde Karlsruhe, Karlsruhe, Germany

The Harold Hamm School of Geology & Geological Engineering, University of North Dakota, Grand Forks, North Dakota, United States

⁶ Department of Geological Sciences, University of Missouri - Columbia, Columbia, Missouri, United States

⁷ X-ray Microanalysis Laboratory, University of Missouri - Columbia, Columbia, Missouri, United States



1 Herpetogaster collinsi from the Cambrian of China elucidates the dispersal and palaeogeographic 2 distribution of early deuterostomes 3 Xianfeng Yang^{1,2,3}, Julien Kimmig^{4,5}, James D. Schiffbauer^{6,7}, Shanchi Peng³ 4 5 6 ¹Yunnan Key Laboratory for Palaeobiology, Yunnan University, North Cuihu Road 2, 650091, 7 Kunming, China ²MEC International Joint Laboratory for Palaeobiology and Palaeoenvironment, Yunnan 8 9 University, 650091, Kunming, China 10 ³State Key Laboratory of Palaeobiology and Stratigraphy, Chinese Academy of Sciences, Nanjing, 11 210008, China 12 ⁴Paläontologie und Evolutionsforschung, Abteilung Geowissenschaften, Staatliches Museum für Naturkunde Karlsruhe, Karlsruhe, 76133, Germany 13 ⁵The Harold Hamm School of Geology & Geological Engineering, University of North Dakota, 14 Grand Forks, ND 58202, USA. 15 ⁶Department of Geological Sciences, University of Missouri, Columbia, MO 65211, USA 16 17 ⁷X-ray Microanalysis Laboratory, University of Missouri, Columbia, MO 65211, USA 18 19 Corresponding Authors: 20 21 Xianfeng Yang 22 Yunnan Key Laboratory for Palaeobiology, Yunnan University, North Cuihu Road 2, 650091, 23 Kunming, China



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24	Email address: yangxf@ynu.edu.cn
25	
26	Julien Kimmig
27	Paläontologie und Evolutionsforschung, Abteilung Geowissenschaften, Staatliches Museum für
28	Naturkunde Karlsruhe, Karlsruhe, 76133, Germany
29	Email address: julien.kimmig@smnk.de
30	
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32	Abstract
UZ	Abstract

The Cambrian Radiation represents one of the largest diversification events in Earth history. While the resulting taxonomic diversity is exceptional, relatively few of these novel species can be traced outside the boundaries of a single palaeocontinent. Those species with cosmopolitan distributions were likely active swimmers, presenting opportunity and means to conquer new areas, but this would not have been the case for sessile organisms. *Herpetogaster* is a genus of sessile, stalked, filter-feeding deuterostomes with two species, *H. collinsi* and *H. haiyanensis*, known from the lower to middle Cambrian (Series 2–Miaolingian, Stage 3–Wuliuan) of Laurentia and Gondwana. Here, we expand the distribution of *H. collinsi* to Gondwana with newly discovered specimens from the Balang Formation of Hunan, China. This discovery raises questions on the origin of the genus and how sessile organisms were able to disperse over such a broad distance in the lower Cambrian.

Subjects Biodiversity, Developmental Biology, Ecosystem Science, Paleontology, Taxonomy

- 48 Keywords Deuterostomia, Ambulacraria, Herpetogaster, Lifestyle, Dispersal, Cambrian Stage 4,
- 49 Exceptional preservation, Larvae, Palaeoecology, Palaeogeography

Introduction

- During the Cambrian Radiation, hundreds of metazoan species, representing nearly all known
- 54 modern phyla, appear in the fossil record and can be found throughout rich fossil deposits around



the world (Paterson et al., 2016; Fu et al., 2019; Harper et al., 2019; Kimmig et al., 2019; Nanglu, Caron & Gaines, 2020; Yang et al., 2021). While there is an excellent record of soft-bodied fossils from globally distributed Burgess Shale-type Lagerstätten as early as Cambrian Stage 3 (~521–514 Ma, not formally ratified), few species can be found crossing palaeocontinental boundaries (Paterson et al., 2016; Fu et al., 2019; Harper et al., 2019; Kimmig et al., 2019; Nanglu, Caron & Gaines, 2020; Yang et al., 2021). This pattern of species distribution not only begs important questions about modes of reproduction and broader-scale mobility or migration of species in the early Cambrian, but also how the few species with cosmopolitan distributions were able to accomplish this feat. Based on newly discovered fossils presented herein, one species that had successfully trekked and established a foothold on multiple palaeocontinents is *Herpetogaster collinsi* Caron et al. 2010.

Herpetogaster, one of the earliest-known representatives of the invertebrate deuterostomes, is a stalked filter feeder grouped with the informal cambroernid clade. The genus is represented by two species found in the lower to middle Cambrian (Series 2–Miaolingian, Stage 3–Wuliuan) of Gondwana and Laurentia (Caron, Conway Morris & Shu, 2010; Kimmig, Meyer & Lieberman, 2019; Yang et al., 2020; Nanglu et al., 2022; Pari, Briggs & Gaines, 2022). To date, its two species have been observed to be restricted to their respective palaeocontinents: H. haiyanensis to Series 2 deposits in Gondwana and H. collinsi to generally younger Miaolingian deposits in Laurentia (with the exception of the Series 2 Pioche Formation). From all known fossils of the genus, Herpetogaster appears to have been a sessile organism, with a stolon and a basal discoidal holdfast. While the stolon may have been contractible, its likely function was to anchor the organism in place within the sediment. Neither of the two species preserve swimming appendages, and thus the presumed mechanism of expanding its distribution would have been through passive transport



by ocean currents. Recent flume experiments have shown that soft-bodied animals can hypothetically be transported over tens of kilometers by current flow (Bath Enright et al., 2021), but this is an improbable explanation for the occurrence of the same species separated by thousands of kilometers and within deposits of different ages. With specimens of *H. collinsi* reported here from the Balang Formation, Hunan, China, along with possible larval fossils, we propose that these organisms possessed a motile larval stage that provided the means for broad dispersal and migration.

Thus far, we have collected more than 60 new specimens of *H. collinsi* from the lower Cambrian (Series 2, Stage 4) Balang Formation. Here, we provide detail of 19 of the best-preserved new specimens, explore the distribution of the genus in time and space, and postulate mechanisms that might have led to its success.

Materials & Methods

The 19 new specimens of *Herpetogaster collinsi* described here are reposited in the collections of the Yunnan Key Laboratory for Palaeobiology, Yunnan University, Kunming, China (YKLP), with specimen numbers YKLP 14570–14588. Specimens with the prefix ROM are housed in the Royal Ontario Museum, Ontario, Canada, with the prefix KUMIP at the Division of Invertebrate Paleontology, Biodiversity Institute, University of Kansas, and with the prefix YPM at the Yale Peabody Museum of Natural History.

Imaging



The specimens were photographed using a Canon EOS 5D digital SLR camera with a Canon 50 mm macro lens and cross-polarized lighting. Close-ups were captured using a Leica DFC 500 digital camera mounted on a Leica M205-C stereoscope. All specimens were submerged in alcohol to increase contrast. The contrast, colour space, and brightness were adjusted using Adobe Photoshop CC.

Scanning electron microscopy and energy-dispersive X-ray spectroscopy analyses.

The backscattered electron (BSE) imaging and energy-dispersive X-ray spectroscopy (EDS) of uncoated specimens was conducted with a FEI Quanta 650 FEG field emission scanning electron microscope (SEM) at the Yunnan Key Laboratory for Palaeobiology, Institute of Palaeontology, Yunnan University, Kunming, China (YKLP). All imaging analyses were conducted with the following operating conditions: $14 \text{ mm} \pm 1 \text{ mm}$ working distance (minor differences to allow for variation in sample thickness or topography) for basic imaging and EDS, 20 keV beam accelerating voltage, 10 nA beam current, 20 Pa chamber pressure (low vacuum), $50 \text{ }\mu\text{m}$ aperture for imaging, and $40 \text{ }\mu\text{m}$ aperture for EDS analysis.

Geological setting

The specimens described herein were collected from the lower part of the Balang Formation in northwestern Hunan, China (Fig. 1A–D). The Balang Formation is part of a conformable lower Cambrian sequence, positioned between the Niutitang Formation below and the Chinghsutung Formation above (Fig. 1E). The presence of the trilobite *Oryctocarella duyunensis* confirms that the collection interval is positioned within Series 2, Stage 4 of the Cambrian System (Peng et al.,





123	2017; National Commission on Stratigraphy of China, 2018; Zhao et al., 2019; Dai et al., 2021).
124	In the context of other well-known Chinese Cambrian Lagerstätten, the Balang biota is situated in
125	age between the slightly older Chengjiang biota (Series 2, Stage 3) and younger Kaili biota
126	(Miaolingian, Wuliuan). The sampled outcrop, located approximately 32 km south-west of
127	Huayuan town, is composed of finely laminated dark grey calcareous mudstone intervals,
128	interbedded with silty mudstone and silty shale (Fig. 1B).
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130	Results
131	
132	Systematic paleontology
133	
134	Superphylum: Deuterostomia Grobben, 1908
135	Clade: Ambulacraria Metschnikoff 1881
136	Unranked stem-group: Cambroernida Caron, Conway Morris & Shu, 2010
137	
138	Genus Herpetogaster Caron, Conway Morris & Shu, 2010
139	
140	Herpetogaster collinsi Caron, Conway Morris & Shu, 2010
141	
142	Holotype. ROM 58051
143	
144	New Material. YKLP 14570–14588
145	



146	Provenance. Balang Formation, lower Cambrian (Series 2, Stage 4), Oryctocarella duyunensis
147	biozone, Mozi village, Paiwu township, approximately 32 km south-west of Huayuan town, Hunan
148	Province, China. Pioche Formation, Comet Shale Member; lower Cambrian (Series 2, Stage 4),
149	Nephrolenellus multinodus biozone; Ruin Wash, NW 1/4 SW 1/4 sec. 15, R65E T2S, 17 km west of
150	Panaca, Lincoln County, Nevada (see Palmer 1998, and Lieberman, 2003 for greater discussion of
151	the locality). Burgess Shale and Stephen Shale Formations; middle Cambrian (Miaolingian,
152	Wuliuan); Yoho and Kootenay National Parks, British Columbia, Canada.
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154	Diagnosis. Segmented body, coiled dextrally. Short head bearing prominent bilateral anterior
155	dendritic tentacles of sub-equal length and in two-by-two arrangement with pharyngeal structures,
156	possibly lateral pores. Trunk subcylindrical, divided into two subsections, narrowing posteriorly.
157	Ventral and contractile adhesive stolon, sometimes with terminal disc. Digestive tract with anterior
158	mouth, pharynx, voluminous stomach, and narrow intestine with terminal anus. Stomach and
159	intestine of sub-equal lengths, un-looped, with triangular mesenterial insertions (from Caron,
160	Conway Morris & Shu 2010).
161	
162	Description. Most of the specimens are complete and range in length from 8.4 to 53.6 mm and
163	from 2.3 mm to 7.9 mm in width. The body is curved, some specimens show fine segmentation
164	along the trunk (Figs. 2E, and 3C,F), and ends in an anus (Figs. 2D–H and 3C,D,F–H).
165	The head has a maximum width of 7.4 mm and a maximum length of 3.9 mm. The pharynx
166	is visible in 9 specimens. Two tentacles emerge from the corners of the head, they reach up to 14.3
167	mm in length, bifurcate at 0.6 to 5 mm and preserve between 10 and 11 branches.



The trunk ranges from 4.8 to 36.1 mm in length, which equates to about three-quarters of
the total body length of the animal. The other quarter represents the pharynx, 0.7 to 3.6 mm in
length. Some specimens (n = 9; Figs. $\frac{2E,I,J}{a}$ and $3A-C,E,F,I$) preserve a darker internal structure,
which is interpreted as the stomach and digestive tract as it reaches the anus in YXLP 14574 and
YKLP14610 (Fig. 2E,J). The soft-tissue surrounding the stomach and intestine preserves fine
segmentation (Figs. 2E,G,J and 3C,F), separating the trunk into 13 segments.

The stolon extends from the final third of the trunk, around the ninth or tenth segment (Figs. 2E,J and 3C,F), and varies in length (1.5 to 14.3 mm) and width (0.4 to 1.8 mm). This suggests that the stolon may have been contractible. Some specimens preserve a terminal disk at the end of the stolon, interpreted as the holdfast (Figs. 2G,I and 3C,H,I).

The Balang specimens appear to have been gregarious or living in close proximity; up to six specimens have been found on a single slab (Fig. 3C,F).

Preservation. Balang Formation specimens are usually regarded as carbonaceous compressions preserved within carbonaceous mudstones, similar to those of other Cambrian Burgess Shale-type deposits (e.g., Wen et al. 2019). Two specimens of *Herpetogaster collinsi* were analysed using SEM-EDS (Fig. 4). The first specimen (YKLP 14583) has a distinctive, if not continuous carbon signature (Fig. 4L), whereas the other specimen (YKLP 14573) did not provide a strong carbon signature, but a more continuous film (Fig. 4E). The analyses also show enrichment of iron and phosphate in the host rock, though neither elemental signature appears to be associated with the fossils themselves. This suggests that that diagenesis played a vital role in the preservation of carbon in the Balang Formation, but also suggests that all the *Herpetogaster* specimens underwent the same taphonomic process. The carbon signature in YKLP 14573 correlates spatially with





calcium (Fig. 4C), suggesting that some of the signal is likely derived from the host rock; however, these elemental signatures do not correlate in YKLP 14583. A similar observation to YKLP 14573 has been made with fossils from the Marjum Formation of Utah (Broce & Schiffbauer, 2017; Leibach et al., 2021). These studies inferred that the higher carbon concentration observed locally stemmed from accumulation of carbonaceous material within interstitial spaces between grains of the host rock. The absence of detectable carbonaceous films or local enrichments of carbon in YKLP 14573 might indicate a few possible taphonomic scenarios involving the complete removal of organic carbon. For example, given their infaunal nature, the organisms likely died within the uppermost oxic- to sub-oxic layers of sediment, allowing for efficient decay by oxygenic microbes. Burial occurred rapidly after death, as indicated by their mostly complete preservation and the presence of preserved labile structures (e.g., tentacles, stolon, gut). Organic carbon removal/dispersion could have continued through diagenesis as well.

An interesting feature of the Balang specimens is that several have a relatively uniform preservational coloration (Figs. 2A,C,D and 3G,H) as compared to representatives of the genus in other deposits. This may be a result of additional decomposition of the specimens after burial, leading to decay of the internal organs, and a generally more homogeneous, black-film appearance. Additionally, rather than all dendritic in form, the tentacles are preserved in different morphologies, as opposed to the Burgess Shale and Pioche Formation specimens (Fig. 5). Enhanced decomposition may again be the culprit for this lack of dendritic detail; overlapping tentacles may also reduce the appearance of finer details, though this is not expected to be the case in all the examined Balang specimens.



Remarks. The Balang Formation specimens are assigned to *H. collinsi*. Though there are taphonomic differences as described above, these specimens appear to be nearly morphologically identical to those described from the Burgess Shale (Caron, Conway Morris & Shu 2010) and the Pioche Formation (Kimmig, Meyer & Lieberman 2019), preserving the dextrally coiled body, anterior dendritic tentacles, pharyngeal structures, stolon, terminal disc, and digestive tract. Segmentation is also prominent in some of the Balang specimens (Figs. 2D,E;,G,H,J and 3A,C,E,F), with segmentation lines visible in 9 of 19 closely examined specimens.

These Balang specimens visibly differ from *H. haiyanensis*, which has over one hundred branches per tentacle, a single layered stolon, and preserved inner and outer layers (Yang et al., 2020)—none of which can be explained by taphonomic differentiation though were plausibly beneficial adaptations to the deltaic environment which they occupied (Peng 2009; Saleh et al. 2022).

Discussion

The oldest-known occurrence of the *Herpetogaster* genus is from the lower Cambrian (Series 2, Stage 3) Chengjiang biota of China (Yang et al., 2020, 2021). The Chengjiang specimens, however, belong to *H. haiyanensis*, which were found in the Haiyan Lagerstätte. From the Haiyan locality, a total of eight specimens, including a juvenile specimen, were described (Yang et al., 2020). The subsequent occurrences are all representatives of *H. collinsi*, and include specimens from the Comet Shale, Nevada (Kimmig, Meyer & Lieberman, 2019), a possible specimen from the Parker Quarry, Vermont (Pari, Briggs & Gaines 2022), and now the specimens reported herein from the Balang Formation (Fig. 5). The youngest representatives known to-date are also



specimens of *H. collinsi*, recovered from the Burgess Shale (Caron, Conway Morris & Shu 2010). While this is likely not yet a complete picture of the distribution and diversity of *Herpetogaster* through time and space, it suggests that the genus might have originated in Gondwana sometime around Cambrian Stage 3 and had likely spread globally, at least in the equatorial range (Fig. 1F), by Cambrian Stage 4. The current fossil record, with the oldest representatives from the Chengjiang biota (Yang et al., 2020, 2021), support the origin of the genus in Gondwana; however, the addition of the new specimens from the Balang biota also suggest that *H. collinsi* originated in Gondwana. Li et al. (2023) suggested *Herpetogaster* is the earliest-diverging cambroernid, and phylogenetically positioned at the base of the Ambulacraria, which aligns well with the origin of the genus on the timeline observed.

This discovery of *H. collinsi* in Laurentia and Gondwana around the same time (Cambrian Stage 4) implores consideration of how these sessile organisms, with no known swimming appendages or mechanisms for motility over broad distances, managed to establish themselves on the shelf of two palaeocontinents separated by thousands of kilometers of open ocean (Fig. 1F). There are many challenges that must have been overcome for this migration to have taken place, not least including the distance, but also predation, and changing water temperatures and oxygenation.

Even with an expansive fossil record and a wealth of soft-bodied organisms from the global distribution of Konservat-Lagerstätten known in the Cambrian (Muscente et al., 2017), it is rare that the same species is found in both Gondwana and Laurentia. In most cases when a Laurentian species has been proposed to occur in a Gondwanan deposit, or vice versa, it has later been revised and given a new species name, or even a new genus (Yang et al., 2021; Hou et al., 2017). However, most of the genera that are shared between the Gondwanan (e.g., Balang, Chengjiang, Emu Bay,



Guanshan, and Kaili) and the Laurentian (e.g., Burgess Shale, Parker Quarry, Pioche Formation, Rockslide Formation, Sirius Passet, Spence Shale, Wheeler Formation, Marjum Formation, and Weeks Formation) biotas are arthropods. Many of these arthropods were pelagic and motile swimmers, and could have been actively seeking new habitats with new or greater resources (Legg & Vannier, 2013; Kimmig & Pratt, 2015; Robison, Babcock & Gunther, 2015; Foster & Gaines, 2016; Paterson et al., 2016; Hou et al., 2017; Lerosey-Aubril et al., 2018, 2020; Fu et al., 2019; Harper et al., 2019; Kimmig et al., 2019; 2023; Ma et al., 2020; Nanglu, Caron & Gaines, 2020; Yang et al., 2021). However, *Herpetogaster*, as a substrate-anchored deuterostome, requires fundamentally different considerations.

An equally plausible alternative for the palaeogeographic distribution of arthropods considers their larvae; the small size and zooplankton-like habit of the post-embryonic nauplius larval stages of arthropods could have traversed long distances by ocean currents (Müller & Walossek, 1986; Zhang & Pratt, 1993; Waloszek & Dunlop, 2002). Therefore, it follows that we should also consider the life cycle of *Herpetogaster*, or at least what we can infer from modern analogues. Though we have numerous fossils of the genus, much remains unknown about its ontogenesis. We can make some assumptions from modern ambulacrarian species; for example, most reproduce sexually, and many develop through a ciliated, free-swimming and feeding larval stage before settling and attachment. Depending on taxon, this motile larval form goes by numerous names, for example, tornaria for enteropneusts, pluteus for echinoids, auricularia for holothurians, and doliolaria for crinoids. These larvae bear little anatomical resemblance to the adult forms, undergoing considerable morphological change before entering their respective adult life stages (Ettensohn, Wessel & Wray 2004). One could suggest from the close affinity of *Herpetogaster* to modern invertebrate deuterostomes (Caron, Conway Morris & Shu 2010; Nanglu



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et al., 2022) that it might have also had a planktonic larval stage. Some evidence for the presence of planktonic larvae has been found in acorn worms from the Chengjiang biota (Yang et al., 2022), thus making this a tantalizing hypothesis. As *Herpetogaster* has been recovered at the base of the Ambulacrarian tree in recent phylogenies (Li et al., 2023), it suggests that the last common ancestor of the clade might have already had a planktonic larval stage, or alternatively that it developed several times within the Ambulacraria.

In addition to suggestions from phylogenetic affinity, another indication of plausible larval dispersal in *Herpetogaster* may come from their gregarious life habit. As noted, we observed some of the *H. collinsi* specimens from the Balang Formation found together on a single slab (Fig. 3C,F), which lends support to former suggestions of a gregarious lifestyle (Caron, Conway Morris & Shu 2010; Kimmig, Meyer & Lieberman, 2019). A planktonic larval stage is common in many other modern gregarious marine invertebrates (Pechenik, 1999; Toonen & Pawlik, 2001), with dispersal and global connectivity well-modeled from broadcast spawning of corals (Wood et al., 2014), and has been hypothesized for more ancient gregarious taxa as well (Cortijo et al., 2015; Schiffbauer et al., 2016). The gregarious lifestyle of H. collinsi may indicate that it, too, was a broadcast spawner, where fertilization and development of the offspring happened externally, promoting wide dispersal of embryos and larvae and resulting in a broad distribution of offspring. While more evidence is necessary to confirm our interpretation, we observed a single millimetric ovoid structure in close proximity to an adult *Herpetogaster* specimen (Fig. 3A). From hints of a lateral band extending around this ovoid, we offer a tentative suggestion that it may be a possible larva (Fig. 3B), as it is comparable in size and form to those of modern acorn worms or other echinoderms (e.g., Yang et al., 2022). This potential larva is understandably speculative, as it does not preserve enough details to be definitively identified, and other examples within this deposit



have not yet been identified. However, previous authors have argued that larvae (*sensu* Hickmann, 1999; Haug, 2018) were likely the key to the worldwide distribution of deuterostomes from Burgess Shale-type biotas (e.g., Han, Zhang & Liu, 2008; Yang et al., 2022), and larvae are already known to play a vital role in Cambrian arthropod development and dispersal (e.g., Liu et al. 2016; Lerosey-Aubril & Laibl, 2021)—both of which urge the continued search for more possible larval fossils from the Balang.

We suggest that *Herpetogaster* having a planktonic larval stage and reproduction through broadcast spawning would have permitted dispersal of this genus over long distances, enabling them to establish a foothold in both Laurentia and Gondwana and providing a logical explanation for the sum of our observations. The question that remains is whether planktonic larvae were shared across the *Herpetogaster* genus, or if they developed later in *H. collinsi*. Testing this hypothesis, however, will require further discoveries of *H. haiyanensis* outside of the Chengjiang region or additional species elsewhere in the Cambrian. Continued investigation of the Balang biota is necessary to confirm or adapt our provisional inferences based on the presence of the tentative but promising tornaria-like larva.

Conclusions

The discovery of *H. collinsi* from the Balang Formation of China represents the first report of this species from Gondwana. Most specimens are complete and the detailed preservation, which required exceptional depositional circumstances, makes an assignment to the species unquestionable. However, the presence of the same sessile species in Laurentia and Gondwana poses the question as to how these organisms managed to colonize two distant parts





328	of the world. We suggest that the most likely scenario sees <i>Herpetogaster</i> with a planktonic
329	larval stage and reproduction through broadcast spawning, which would have permitted dispersal
330	over long distances and enabled <i>H. collinsi</i> to establish a foothold in both Laurentia and
331	Gondwana. As <i>Herpetogaster</i> is already known from Cambrian Stage 3 in Gondwana, it would
332	suggest a possible origination of the genus there, but the limited record of the genus at that time
333	does not yet allow for a definite conclusion.
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335	Competing Interest
336	The authors declare no competing interest.
337	
338	Author Contributions
339	Xianfeng Yang conceived and designed the study, collected specimens, performed the data
340	analyses, analyzed the data, prepared figures and/or tables, authored or reviewed drafts of the
341	article, and approved the final draft.
342	
343	Julien Kimmig and James D. Schiffbauer conceived and designed the study, analyzed the data,
344	prepared figures and/or tables, authored or reviewed drafts of the article, and approved the final
345	draft.
346	
347	Shanchi Peng collected specimens, analyzed the data, authored or reviewed drafts of the article,
348	and approved the final draft.
349	
350	Data availability





351	All data used in the analyses are as part of the text or electronic supplementary material.
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569 **Figures** 570 Figure 1: Distribution of *Herpetogaster*. 571 (A) Location of the studied section of the Balang Formation located approximately 32 km south-572 west of Huayuan town, Hunan Province, South China. (B) Generalized stratigraphy of the Balang 573 574 Formation at this location. (C) Palaeogeographical distribution of *Herpetogaster* during Cambrian Stages 3-4 and during the Wuliuan (modified from Streng and Geyer 2019). 1. H. hanyanensis 575 (Chengjiang biota, China, Stage 3), 2. H. collinsi (Balang biota, China, Stage 4), 3. H. collinsi 576 577 (Burgess Shale, Canada, Wuliuan), 4. H. collinsi (Pioche Formation, Nevada, USA, Stage 4), 5. H. sp. (Parker Formation, Vermont, USA, Stage 4). Abbreviations: Ni., Niutitang Formation; Chi, 578 579 Chinghsutung Formation. 580 Figure 2: Complete specimens of Herpetogaster collinsi Caron et al. 2010 from the Balang 581 Formation of China. 582 (A) YKLP 14570 a small, likely juvenile specimen, preserving showing slim paired tentacles and 583 the extended stolon. (B-C) YKLP 14571 and YKLP 14572 part and counterpart of a likely sub-584 585 adult specimen, the trunk is almost completely preserved as a black carbonaceous film, with the detail of dendritic symmetrical tentacle and a stolon. (D) YKLP 14573, a large adult specimen, 586 with the showing the branching tentacles. Co-occurring with a relative individual. (E) YKLP 587 588 14574, adult specimen with prominent segments and segment boundaries (indicated by arrows). (F) YKLP 14575, adult specimen coiled with stolon. (G) YKLP 14576, adult specimen with well-589 590 preserved anus and terminal disc. (H) YKLP 14577, adult specimen with both tentacles and well-591 preserved anus. (I) YKLP 14578, sub-adult specimen with well-preserved branching tentacle. (J)



592 YKLP 14610, adult specimen with showing segments and segment boundaries. Scale bars: (A) 2mm; (B-J) 5mm. Abbreviations: an, anus; in, intestine; p?, putative pharyngeal pores; ph, 593 pharynx; seg, segment boundary?; st, stolon; stom, stomach; td, terminal disc; te, tentacle. 594 595 596 Figure 3: Gregarious specimens of Herpetogaster collinsi from the Balang Formation of 597 China. (A) YKLP 14580, large adult specimen is found co-occurring with associated plankton, possibly 598 larvae. (B) YKLP 14581, close-up of the associated tonaria-like structure, which shows similarities 599 600 to extant early stage ambrulacarian larvae. (C) YKLP 14582, at least six specimens of H. collinsi on a single slab. (D) YKLP 14584, two adult specimens with almost complete paired tentacles. (E) 601 602 YKLP 14585, two adult specimens. (F) YKLP 14583, counterpart of (C). (G) YKLP 14586, two 603 complete adult specimens. (H) YKLP 14587, three specimens with almost complete paired tentacles, one preserving the stolon and terminal disc. (I) YKLP 14588, at least two adult 604 specimens. Scale bars: (B) 1mm; (A, C–I) 5mm. Abbreviations: an, anusph, pharynx; seg, segment 605 606 boundary?; st, stolon; td, terminal disc; te, tentacle. 607 608 Figure 4: SEM micrograph and SEM-EDS elemental maps of Herpetogaster collinsi from the **Balang Formation of China.** 609 (A–G) YKLP 14573. (A) Picture of the specimen indicating the analyzed area (dashed rectangle). 610 611 (B) Detailed view of the analyzed area. (C-G) SEM-EDS elemental maps of Ca, Fe, C, P, S, respectively. (H-N) YKLP 14583. (H) Picture of the specimen indicating the analyzed area 612 (dashed rectangle). (I) Detailed view of the analyzed area. (J–N) SEM-EDS elemental maps of Ca, 613 614 Fe, C, P, S, respectively. Scale bars: (A, H) 5mm; (C-G, J-N) 1mm.



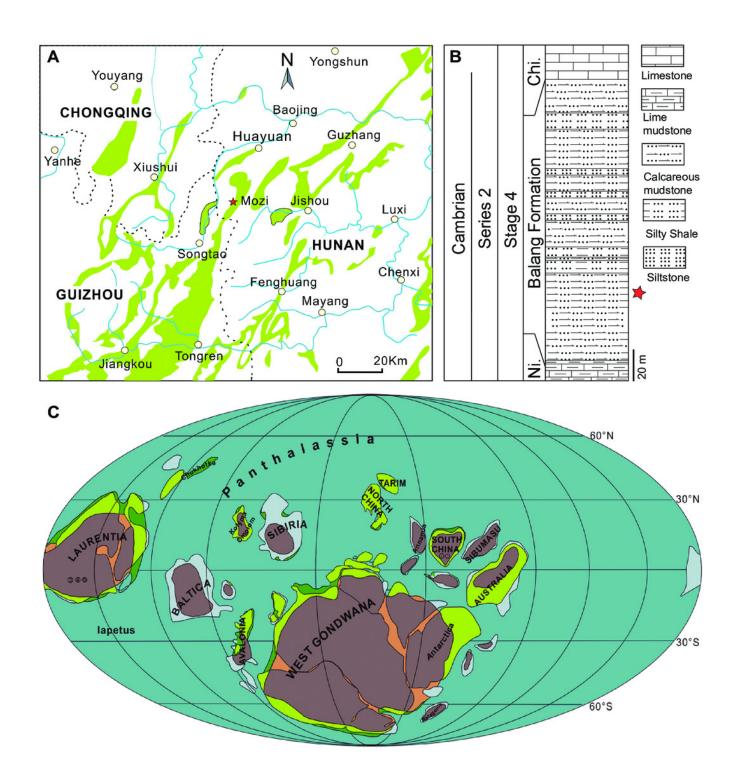
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616	Figure 5: All known Herpetogaster species from Gondwana and Laurentia.
617	(A) YKLP 14404, holotype of <i>Herpetogaster haiyanensis</i> from the Chengjiang biota (Cambrian;
618	Series 2; Stage 3) of Yunnan, China; (B) YPM IP 239054, ?Herpetogaster sp. from the Parker
619	Formation (Cambrian; Series 2; Stage 4) of Vermont, USA; (C) YKLP 14576, Herpetogaster
620	collinsi from the Balang Formation (Cambrian; Series 2; Stage 4) of China; (D) KUMIP 482878,
621	Herpetogaster collinsi from the Pioche Formation (Cambrian; Series 2; Stage 4) of Nevada, USA;
622	(E) ROM 58051, holotype of Herpetogaster collinsi from the Burgess Shale (Cambrian;
623	Miaolingian; Wuliuan) of British Columbia, Canada. Scale bars: 5mm.





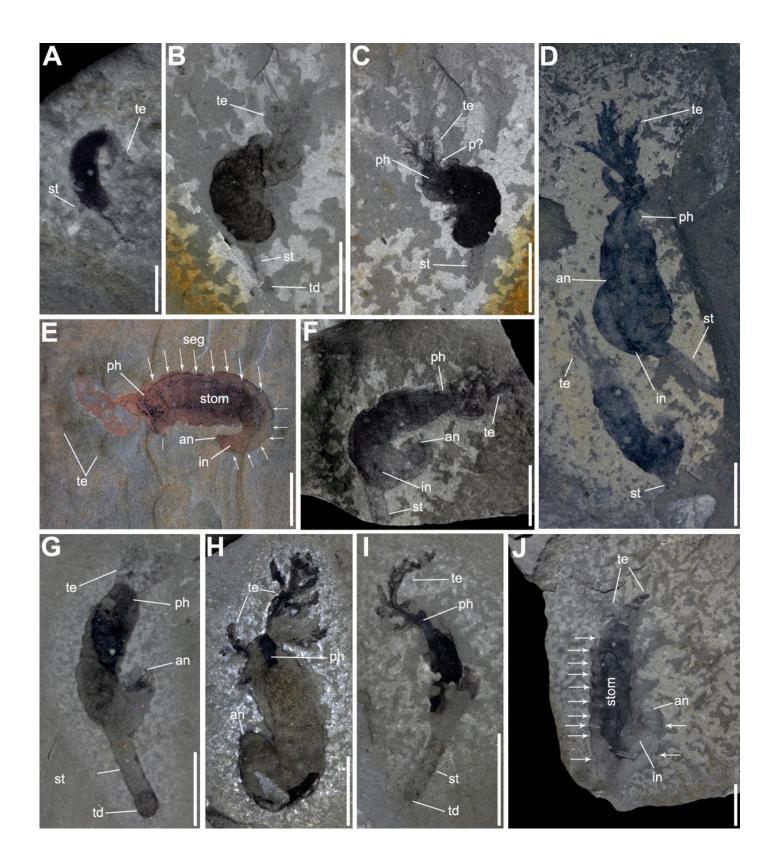
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626	Supplementary Table 1. Measurements of the Balang Formation specimens
627	

Distribution of Herpetogaster.



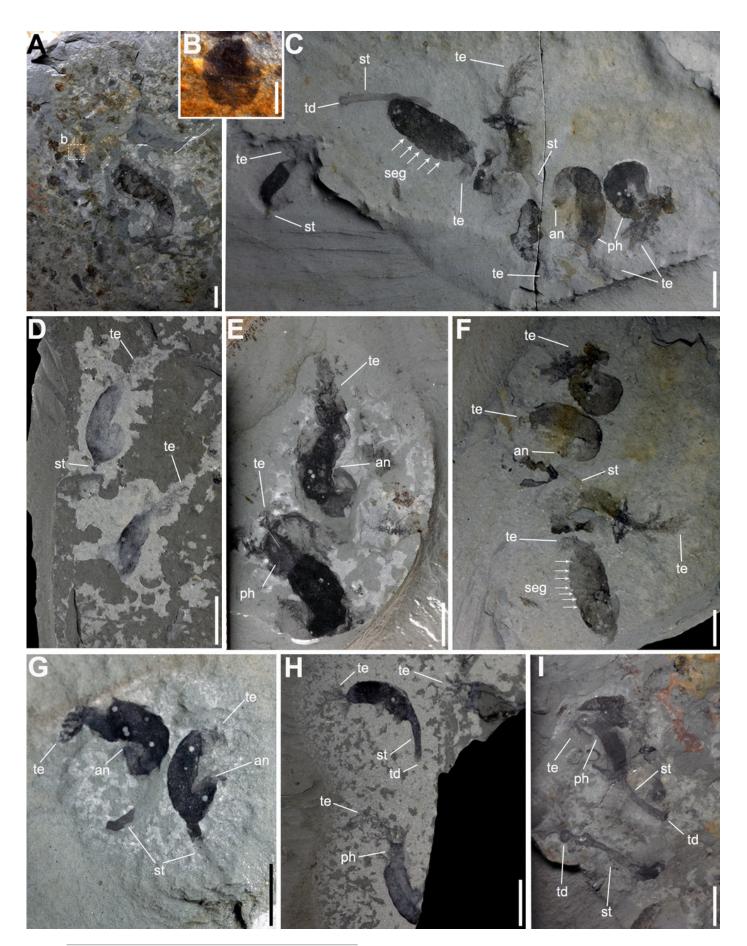


Complete specimens of *Herpetogaster collinsi* Caron et al. 2010 from the Balang Formation of China.



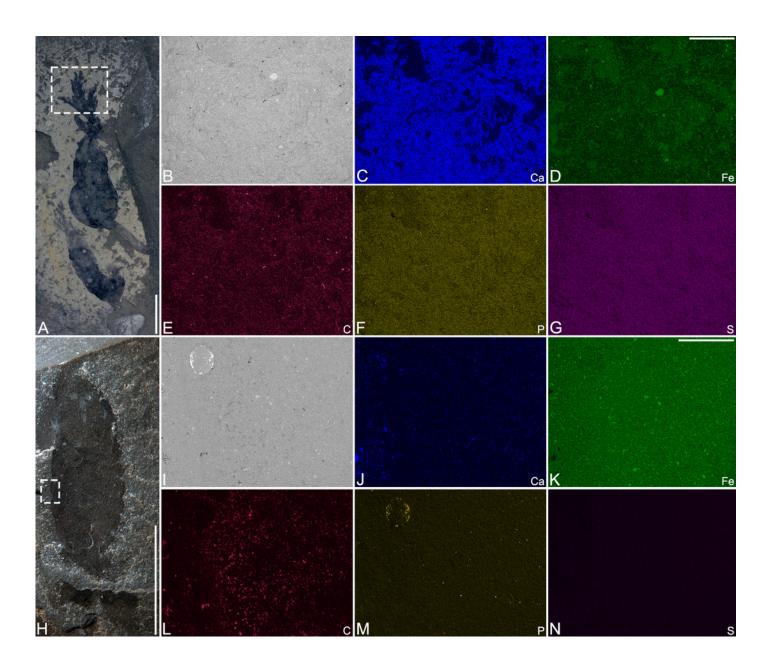


Gregarious specimens of Herpetogaster collinsi from the Balang Formation of China.



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SEM micrograph and SEM-EDS elemental maps of *Herpetogaster collinsi* from the Balang Formation of China.



All known Herpetogaster species from Gondwana and Laurentia.

