

1 **Vertically migrating *Isoxys* and the early Cambrian biological pump**

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10

11 **Abstract**

12 The biological pump is crucial for transporting nutrients fixed by surface-dwelling primary producers  
13 to demersal animal communities. Indeed, the establishment of an efficient biological pump was  
14 likely a key factor enabling the diversification of animals over 500 million years ago during the  
15 Cambrian explosion. The modern biological pump operates through two main vectors: the passive  
16 sinking of aggregates of organic matter, and the active vertical migration of animals. The coevolution  
17 of eukaryotes and sinking aggregates is well understood for the Proterozoic and Cambrian, however  
18 little attention has been paid to the establishment of the vertical migration of animals. Here we  
19 investigate the morphological variation and hydrodynamic performance of the Cambrian  
20 euarthropod *Isoxys*. We combine elliptical Fourier analysis of carapace shape with computational  
21 fluid dynamics simulations to demonstrate that *Isoxys* species likely occupied a variety of niches in  
22 Cambrian oceans, including vertical migrants, providing the first quantitative evidence that some  
23 Cambrian animals were adapted for vertical movement in the water column. Vertical migration was  
24 one of several early Cambrian metazoan innovations that led to the biological pump taking on a  
25 modern-style architecture over 500 million years ago.

26 **Key words:** biological pump; Cambrian; computational fluid dynamics; *Isoxys*; pelagic

27

## 28 Introduction

29 The biological pump is the process by which organic nutrients are transported from shallow ocean to  
30 deep sea [1]. Today this consists of two major vectors: passive sinking of organic matter aggregates  
31 and vertical movement of animals [1] (**Fig 1**). The biological pump, the main driver of the marine  
32 carbon cycle, is responsible for approximately two thirds of the vertical gradient of carbon in the  
33 ocean [2]. While transport of carbon to the deep ocean through vertical mixing of dissolved organic  
34 carbon (DOC) carries significant amounts of carbon to depth, 95% of DOC cannot be used as food by  
35 marine organisms [1,3]. In contrast, aggregates and vertical migrants concentrate organic matter in a  
36 form that can be utilized by demersal animals [1] and are therefore crucial for the establishment and  
37 sustenance of deep-water communities.

38 The biological pump was very different before the Cryogenian Period (>720 Ma).  
39 Cyanobacterial picoplankton (0.2–2.0  $\mu\text{m}$ ) domination led to a stratified and turbid water column  
40 [e.g. 4–6]. Cells were too small to sink in significant numbers, so most nutrients were recycled within  
41 the surface waters, with little export to the deeper ocean [4–6]. The flux of the aggregation vector  
42 (**Fig. 1**) would have increased when primary productivity shifted to be eukaryote-dominated during  
43 the Cryogenian  $\sim$ 650 Ma [7], and further when Cambrian suspension feeding sponges and pelagic  
44 phytoplanktivores applied a size-selective pressure for larger primary producers [8–12]. The  
45 carcasses, sloppy feeding, and faecal pellets of macrozooplankton, including phytoplanktivorous  
46 crown-group crustaceans [e.g. 9], would have further increased the formation and size of  
47 aggregates, and thus the flux along this vector. However, despite the importance of vertical  
48 migration in the transport of nutrients necessary to sustain mesopelagic and deep-water ecosystems  
49 in the modern ocean [e.g. 13], this vector is poorly understood in the Cambrian.

50 Comparisons of the morphology of Cambrian fossil organisms with modern vertically mobile  
51 pelagic animals provides the opportunity to infer whether vertical migration occurred in oceans over  
52 500 million years ago. Qualitative comparisons between the pelagic crustacean *Gnathophausia* and  
53 the nektonic stem group euarthropod [14] *Isoxys* [e.g. 15,16] suggest the latter is a promising  
54 candidate. The 20 *Isoxys* species so far formally described are united by the presence of a bivalved  
55 carapace which bears both anterior and posterior spines [17] (**Fig. 2**), and the genus has a  
56 cosmopolitan distribution [e.g. 18]. The presence of eyes that comprise  $\sim$ 10% of the body length, a  
57 digestive tract with paired serial midgut glands, and a pair of anteriorly positioned raptorial  
58 appendages (**Fig. 2**) support a predatory habit for *Isoxys*, which would have been well-suited for  
59 capturing small soft-bodied invertebrates [19]. *Isoxys* is unusual for Cambrian animals as a pelagic  
60 lifestyle has been proposed [e.g. 18], although some recent workers have suggested a potential

61 hyperbenthic lifestyle (1–10 m above the bottom), with individuals capable of moving small  
62 distances vertically [19,20]. However, while *Isoxys* carapaces appear to show adaptations for  
63 hydrodynamic streamlining, interspecific differences in both carapace asymmetry and spine lengths  
64 (e.g. **Fig. 2**), as well as soft parts, suggest that different species may have occupied distinct niches,  
65 including some much closer to the seafloor [21].

66           Here, we provide the first quantitative assessment of carapace shape variation across the  
67 group and compare *Isoxys* to other Cambrian ‘bivalved’ euarthropods and *Gnathophausia*.  
68 Subsequently, through computational fluid dynamics simulations, we test the importance of the  
69 spines and carapace outline for generating lift and reducing drag, and thus the ability of different  
70 taxa to move vertically through the water column. These analyses support the hypothesis that *Isoxys*  
71 taxa occupied a variety of distinct niches in Cambrian oceans, including vertical migrants.

72

## 73 **Materials and methods**

### 74 Outline analyses

75 Two-dimensional outlines of 20 *Isoxys* species, its sister taxon *Surusicaris elegans* [22], 6 *Tuzoia*  
76 species, and 11 *Gnathophausia* species were constructed in Inkscape from literature sources by SP  
77 (Supplementary Table 1) and imported into R [23] for elliptical Fourier analysis (EFA). DAL  
78 independently constructed 20 *Isoxys* and 1 *Surusicaris* outlines directly from photographs of fossil  
79 specimens to allow assessment of the error introduced in the creation of outlines (Supplementary  
80 Table 2; Supplementary Materials 1). *Tuzoia* was chosen for comparison as it is also common in  
81 Cambrian communities, and has been suggested to be closely related to *Isoxys* based on similarities  
82 in the structure of the eyes and carapace shape [e.g. 24]. *Gnathophausia* was selected because  
83 similarities in the carapace morphology of one species (*G. zoea*) and *Isoxys* have been repeatedly  
84 noted [e.g. 16,19], the carapaces of these animals are similar in size (10–30 mm), and multiple  
85 species of *Gnathophausia* are known to be vertically mobile in the modern ocean, having been  
86 sampled from surface waters and at depths of over 3000 meters [25].

87 Outlines were sampled at the same resolution (64 points provided sufficient detail to  
88 distinguish taxa), centred, scaled by centroid size, and subjected to EFA using the *Momocs* package  
89 [26]. Harmonics describing 99.9% of the variation were retained. EFA results were visualized with a  
90 Principal Components Analysis. A hierarchical clustering analysis (cluster package; [23])  
91 quantitatively grouped similarly shaped carapaces together using all principal components.

### 92 Computational fluid dynamics

93 *Isoxys* species reflecting the variation in carapace shape over both PC1 and PC2 were chosen  
94 for inclusion in computational fluid dynamics simulations, and *Gnathophausia zoea* was analysed for  
95 comparison. We chose to undertake a two-dimensional analysis to conserve computational  
96 resources and minimize errors in the modelled geometries (the exact three-dimensional shape is  
97 unknown for most taxa). This is justified because undeformed *Isoxys* specimens preserved in dorsal  
98 view show a narrow profile [16,21,27,28]. Propulsion during swimming derived from movement of  
99 the ventral appendages, and not from flapping of the bivalve carapace [15, 27]. A lack of adductor  
100 muscles means that *Isoxys* was unable to alter the size of the gape during swimming [15], and the  
101 numerous specimens preserved in ‘butterfly’ orientation are considered exuviae [27]. In addition,  
102 the full variation of the carapaces considered in this study can be visualized in two dimensions. Two-  
103 dimensional analyses are commonly performed on analyses of wing outlines to assess aerodynamic  
104 performance by both biologists and engineers, [e.g. 29,30]. A 2D analysis is suitable for this study as

105 the wake behind the *Isoxys* carapaces is steady at the Reynolds numbers considered, so can be  
106 modelled with two-dimensional simulations [32, 34]. The impact of soft parts such as eyes  
107 protruding from the carapace was analysed for two taxa: *Isoxys acutangulus* and *I. longissimus*.  
108 *Isoxys* was assumed to be negatively buoyant, like *Gnathophausia* [31]. All *Isoxys* species were  
109 assumed to have the same carapace composition and density. The cuticle ornamentation in some  
110 *Isoxys* species is not expected to impact the drag at the low Reynolds numbers considered in this  
111 study, as the roughened surface falls within the slowly moving fluid near the carapace surface [32].

112         Following validation and verification of the model and setup for low Reynolds numbers, and  
113 mesh quality assessments using ANSYS Mesh and ANSYS Fluent (*Ansys® Academic, Release 2020 R2*;  
114 Supplementary Materials 2), outlines of the selected *Isoxys* species and *G. zoea* were exported from  
115 R as .txt files readable by ANSYS DesignModeler (*Ansys® Academic, Release 2020 R2*; Supplementary  
116 Materials 2), and standardized to a dorsal length (chord length) of 25 mm. This allowed size-  
117 independent comparisons of hydrodynamic performance of shapes. While some *Isoxys* taxa (e.g. *I.*  
118 *communis*, *I. longissimus*) can reach up to 50 mm, other adult forms only reach ~20 mm (e.g. *I.*  
119 *glæssneri*, *I. volucris*) [18,27]. A size of 25 mm represents a compromise size for comparison  
120 between these larger and smaller forms, with the influence of larger size able to be assessed by  
121 simulating higher Re (as Re depends on both size and swimming speed).

122         Computational fluid dynamics (CFD) simulations were conducted using the steady-state  
123 laminar solver in ANSYS Fluent (*Ansys® Academic, Release 2020 R2*). The laminar solver performed  
124 best of the three considered during validation and verification (laminar, SST, k-epsilon;  
125 Supplementary Materials 2), as expected for the low Reynolds numbers in this study [32].

126         Coefficients of drag (Cd) and lift (Cl) (Supplementary Materials 2) were calculated under  
127 three flow speeds equating to 0.75, 1.00 and 1.18 body lengths per second (chord Reynolds  
128 numbers, Re, 255, 340, and 400 respectively for saltwater conditions at 0°C; Supplementary  
129 Materials 2). These Re were chosen as swimming speeds of between 75% and 100% of the body  
130 length per second have been observed in adult *Gnathophausia ingens* (carapace mean length ~25  
131 mm; [33]), and as the laminar model was validated against published drag and lift data for NACA  
132 airfoils at exactly Re = 400 [29]. The chord length (25 mm) was taken as the reference area for both  
133 coefficients.

134         Solutions were considered converged when residuals were  $<10^{-6}$ . Simulations were run at  
135 numerous angles of attack, to evaluate the hydrodynamic performance of carapaces at multiple  
136 orientations. In tank experiments, *Gnathophausia ingens* has been observed to change angle of  
137 attack to generate more lift or less drag at different swimming speeds [33]. The angle of attack was

138 increased from 0–8° at all Re, until the stall angle could be identified and/or unsteady flow was  
139 observed. If the stall angle was not reached, further experiments were run until the maximum lift  
140 coefficient was obtained. Negative angles of attack were also simulated to assess the negative lift  
141 generated by the different outlines. In all cases, the absolute value of the negative angle of attack  
142 was increased until the drag coefficient was equal to or greater than that of the stall angle. When  
143 unsteady flow was suspected to be the reason that steady state simulations did not converge, the  
144 steadiness of the flow field was determined by carrying out a time-dependent analysis of 100 time  
145 steps, with each time step equal to the flow speed (so, for an inlet velocity of 0.01875 ms<sup>-1</sup>, the time  
146 step = 0.01875 s).

147

## 148 Results

### 149 Outline analysis

150 In the outline analysis, a total of 18 harmonics were retained. Principal coordinates 1 and 2  
151 described 79.3% of the total variation. Carapace asymmetry, narrowness, and spine length increased  
152 as PC1 became more positive, while the length of the anterior spine relative to the posterior spine  
153 corresponded to an increase in PC2 (**Fig. 3**). *Isoxys* occupied the largest area in the morphospace.  
154 Visual overlap of the areas occupied by the genera demonstrated that some *Isoxys* taxa were more  
155 similar in shape to *Gnathophausia* than their Cambrian relatives *Surusicaris* and *Tuzoia*. Confirmation  
156 was provided by the cluster analysis (**Fig. 3**). All six *Tuzoia* species formed a single cluster, with *Isoxys*  
157 taxa spread over the three remaining groups (*Surusicaris* group, *Gnathophausia* groups 1 and 2).  
158 Species clustered with *Surusicaris* have symmetric and deep carapaces and relatively short spines.  
159 Species in the *Gnathophausia* groups displayed narrower outlines whose narrowness, asymmetry  
160 and spine length increased with PC1. The single species of *Isoxys* in *Gnathophausia* group 1, *I.*  
161 *paradoxus*, displayed an anterior spine much more elongate than its posterior one, that contrasted  
162 with the seven species in *Gnathophausia* group 2 with their spines of approximately equal length.

### 163 Computational fluid dynamics

164 Computational fluid dynamics simulations assessed the impact of increasing asymmetry,  
165 spine length, and relative lengths of anterior and posterior spines on hydrodynamic performance.  
166 Inclusion of eyes did not significantly impact the hydrodynamic performance of carapaces  
167 (Supplementary Materials 3). *Isoxys zhurensis*, the most symmetric species chosen for analysis,  
168 created an unsteady wake with Kármán vortex street at the lowest Re considered, and so no drag or  
169 lift coefficients were obtained (Supplementary video). The flow around remaining *Isoxys* outlines  
170 was laminar with a steady wake, and there was no evidence for three dimensionality  
171 (Supplementary Materials 4). Greater asymmetry and narrowness (more positive in PC1, **Fig. 3**)  
172 resulted in lower drag coefficients, as demonstrated by a comparison of the short-spined taxa *I.*  
173 *chilhoweanus*, *I. acutangulus*, and *I. mackenziensis*. The most asymmetric of these forms, *I.*  
174 *mackenziensis*, produced lower drag coefficients than the other two, but ranges of lift coefficients  
175 were similar for all three (**Fig. 4a**). More elongate spines increased the range of lift coefficients  
176 (vertical bars, **Fig. 4a**) and, significantly, negative lift coefficients at negative angles of attack (e.g.,  
177 compare *I. mackenziensis*, *I. communis*, and *I. longissimus*). In *I. paradoxus*, where the anterior spine  
178 is much longer than the posterior one (more positive in PC2, **Fig. 3**), the range of lift coefficients  
179 further increased (**Fig. 4a**). Similar drag coefficients and ranges of lift coefficients were obtained in

180 an analysis of the hydrodynamics of *Gnathophausia zoea*, when either the carapace alone or both  
181 the carapace and abdomen were considered (**Fig. 4b**).



182 **Discussion**

183 Vertical migrations and niche partitioning in Isoxys

184 Functional morphology of *Isoxys* fossil specimens supports an off-bottom (hyperbenthic or pelagic)  
185 life habit for this animal [15,16,18,19,34], based on the eye orientation (forwards, slightly ventral)  
186 and the elongate slender carapace shape of *Isoxys*. Our study combines outline morphometric and  
187 computational fluid dynamics analyses and suggests that *Isoxys* species occupied a variety of niches,  
188 including some as pelagic vertically mobile predators.

189 Lift and drag coefficients of *Isoxys* carapaces indicate variation in the depth range and  
190 swimming speeds of these species. *Isoxys* taxa clustering with *Surusicaris* (**Fig. 3**) generate positive  
191 lift, but do not generate negative lift (**Fig. 4**). This supports suggestions of previous workers that  
192 these *Isoxys* species may have occupied a hyperbenthic (1–10 m above the seafloor) and/or  
193 nektobenthic [21] niche, perhaps moving vertically short distances in the water column [15,19,35].  
194 Vertical movement would be achieved by altering the angle of attack to produce lift force greater  
195 than (ascent), equal to (horizontal swimming), or less than (descent) the impact of their negative  
196 buoyancy. Drag reduction associated with streamlining would have allowed some taxa (e.g., *I.*  
197 *mackenziensis*) to capture faster-moving prey animals, as faster swimming speeds could have been  
198 maintained over longer distances for the same metabolic cost.

199 *Isoxys* species clustering with *Gnathophausia* show convergent adaptations to moving  
200 vertically in the water column, such as asymmetric carapaces with elongate anterior and posterior  
201 spines (**Figs. 3, 4**). This does not preclude elongate spines from also acting as anti-predatory  
202 deterrents, as suggested by [15]. These adaptations provided hydrodynamic benefits that would  
203 have allowed the *Isoxys* species to operate over a wider bathymetric range. A streamlined carapace  
204 facilitates not only faster movement, but also more efficient swimming, beneficial for migrations  
205 over a long distance. The carapace shapes of *I. longissimus* and *I. paradoxus* generate lift coefficient  
206 ranges and minimum drag coefficients comparable to the modern crustacean *Gnathophausia zoea*  
207 (**Fig. 4**), which has been recovered at depths ranging from surface waters down to ~3000 m in the  
208 modern ocean [25]. These results also suggest that an elongate abdomen in *G. zoea* reduces the  
209 drag experienced by the animal slightly but does not greatly impact on the range of lift coefficients  
210 (**Fig. 4b**), though the abdomen may also play a physical role. Animals that move vertically in the  
211 water column do not have to cover the entire distance from the surface ocean to demersal  
212 communities, and instead sometimes migrate across only a shorter vertical distance. Thus, *Isoxys*  
213 taxa with the broadest ranges of lift coefficients (*I. longissimus* and *I. paradoxus*) likely covered a  
214 wider depth range than those with smaller ranges of lift coefficients (e.g., *I. communis*).

215 Corroborating evidence for variation in bathymetric range for different *Isoxys* species comes  
216 from the fossil record itself. Members of different groups as resolved in the cluster analysis (convex  
217 hulls, **Fig. 3**) co-occur with different relative abundances in Cambrian deposits preserving soft-  
218 bodied fossils. In general, species with inferred vertically migrating lifestyles are much rarer than  
219 those that lived close to the seafloor. In the Chengjiang Biota, *Isoxys auritus* (*Surusicaris* group)  
220 greatly outnumbers both *I. paradoxus* and *I. curvirostratus* (*Gnathophausia* groups 1 and 2,  
221 respectively) (**Fig. 3**; [15,21,36–38]). A similar pattern of relative abundances can be observed in the  
222 two Burgess Shale taxa (**Fig. 2**): in the Walcott Quarry, *I. acutangulus* (*Surusicaris* group) comprises  
223 nearly 0.5% of the total community, vastly outnumbering the extremely rare inferred vertical  
224 migrant *I. longissimus* (*Gnathophausia* group 2; [16,39]). The relative abundances of these *Isoxys*  
225 species can be partly explained by the differences in lifestyle predicted by the carapace outline and  
226 soft anatomy. The less hydrodynamic taxa (those with higher drag coefficients and narrower ranges  
227 of lift coefficients) likely lived near the seafloor, with the more streamlined species living in the  
228 water column and occupying a broader bathymetric range. This broader bathymetric range would  
229 have included more open water settings, beyond the maximum depth of the shelf where Cambrian  
230 deposits preserving soft-bodied fossils occur – *Gnathophausia zoea* for example has been found at  
231 depths of up to 3000 metres [25]. As modern euarthropod carapaces disarticulate quickly after  
232 death [e.g. 40], the preservation potentials for pelagic euarthropods living high in the water column  
233 are lower than for those living closer to the seafloor. In addition, animals which occupy an ecological  
234 niche in the open water are less likely to find themselves over shelf environments such as those  
235 which preserve soft-bodied fossils or be trapped and transported by an obrution event responsible  
236 for the preservation of soft bodied communities in these settings. The small numbers of vertically  
237 mobile *Isoxys* individuals observed may have been at the bottom of their vertical migrations and/or  
238 been transported horizontally by currents. *Isoxys* species are not globally distributed [41]. Many  
239 species (for example those clustering with *Surusicaris*) appear suited to hyperbenthic habits, and so  
240 would be expected to have provincial distributions. The limited geographic distribution of *I.*  
241 *longissimus* and *I. paradoxus* is most likely due to a combination of factors. Firstly, deposits where  
242 *Isoxys* is expected to be preserved are not evenly distributed in time and space – Stage 3 deposits  
243 are mostly in South China, while Wuliuan and younger are mostly in Laurentia [42], though the  
244 absence of the Chengjiang species *I. paradoxus* from Sirius Passet is notable. Secondly, the lower  
245 preservation potential of pelagic (compared to hyperbenthic) species means that they are rare even  
246 in Tier 1 Burgess Shale-type Lagerstätten (*sensu* [42]). However, despite its rarity, the Burgess Shale  
247 species *I. longissimus* has a wider known geographic range than the co-occurring *I. acutangulus*. The  
248 former has also been reported from the Wheeler Formation, House Range, Utah, USA [43]. The

249 situation appears more complex in the Emu Bay Shale, where the more hydrodynamic species *I.*  
250 *communis* greatly outnumbers the less streamlined *I. glaessneri* [27]. However, the Emu Bay Shale is  
251 not a traditional Burgess Shale-type deposit, instead it represents a localized deep water micro basin  
252 on the inner shelf [44]. Here fluctuating oxygen levels may have periodically deoxygenated the water  
253 column, possibly killing pelagic taxa like *I. communis* in great numbers and creating a taphonomic  
254 bias that preferentially preserves pelagic taxa.

255 Further support for the Chengjiang taxon *Isoxys auritus* occupying a niche closer to the  
256 seafloor than *I. curvirostratus* comes from a comparison of the soft anatomy (soft parts are unknown  
257 for *I. paradoxus*) [21]. The stout endopods of *I. auritus* appear well-suited for interacting with the  
258 substrate, while exopods with broad fringing lamellae and a sophisticated vascular system in the  
259 more streamlined *I. curvirostratus* suggest it was a more powerful swimmer, providing additional  
260 support for a pelagic habit [21].

261 Lastly, a compendium of fossil, geochemical, and phylogenetic data show that vertically  
262 mobile *Isoxys* species would have had access to a variety of pelagic prey items and an oxygenated  
263 water column to travel in. Cambrian oceans were not stratified, instead displaying wedge-shaped  
264 oxygen minimum zones broadly comparable to modern oceans [45]. *Isoxys* prey size range (~5–20  
265 mm; [19]) includes Cambrian phytoplanktivores, such as crustaceans with setae and filter plates  
266 from Sirius Passet and Mount Cap (15–50 mm) [4,8,9,12], as well as crown group branchiopods and  
267 copepods and total group ostracods [46], as well as bradoriids, some of which were also likely  
268 pelagic [47,48]. The presence of planktonic larvae, another possible prey item, can be inferred from  
269 tip-dated phylogenetic analyses that support the evolution of metamorphosis in euarthropods by  
270 the Cambrian [49]. Thus, a range of different data sources suggest multiple *Isoxys* taxa were  
271 vertically mobile, and that the Cambrian ocean could support such an ecology.

272

### 273 Metazoans and the Cambrian biological pump

274 The presence of the likely vertically mobile *Isoxys paradoxus* in the Chengjiang Biota makes it the  
275 oldest confidently identified euarthropod vertical migrant, and likely among the first animals to  
276 employ this life habit. For this vector to be significant by the Cambrian Stage 3, a large biomass of  
277 *Isoxys* would need to move vertically. While pelagic animals have a lower preservation potential  
278 than benthic ones (for example very few fossil copepods are known [50]), *Isoxys* species with  
279 inferred (hyper)benthic habits are extremely abundant in both the Chengjiang and Burgess Shale  
280 [15,16,21,36–38]), and their pelagic counterparts *Isoxys longissimus* and *I. paradoxus* may have been  
281 similarly numerous. The Chengjiang and coeval Qingjiang biotas also preserve the earliest evidence

282 for gelatinous zooplankton, which move vertically small distances in the modern ocean [51,52].  
283 However, an active swimming *Isoxys* would have covered greater distances more rapidly.  
284 Furthermore, the presence of a through gut would have increased processing time for food, vital for  
285 transporting nutrients consumed in surface waters to the deep ocean as faecal pellets. Vertical  
286 migration was one of many important eukaryotic and metazoan innovations key to establishing a  
287 modern-style biological pump. A series of metazoan innovations which appear in the fossil record in  
288 quick succession during the early Cambrian gave the biological pump a modern-looking structure  
289 (**Fig. 5**), which was strengthened during the Phanerozoic.

290 The shift to primary production dominated by eukaryotes, and the innovation of active  
291 suspension feeding, would have ventilated the oceans, cleared organic matter in the water column,  
292 and increased transfer of oxygen and plankton of increasing sizes from the surface to the sediment-  
293 water interface [e.g. 6]. The first of these events occurred during the Cryogenian [7], but while  
294 benthic passive suspension feeders are known from the Ediacaran [e.g. 53], early Cambrian stem and  
295 crown-group sponges represent the oldest benthic active suspension feeders [10,11]. As active  
296 suspension feeders, sponges were able to transport large volumes of water between benthic and  
297 pelagic realms [e.g. 54], and the most abundant sponges from the Cambrian, the reef building  
298 archaeocyaths, display pore size differentiation within reef systems [55], presumed to be evidence  
299 of prey size-selectivity. This illustrates that predator–prey feedbacks were present in the Cambrian  
300 Stage 2 (**Fig. 2**), presenting a potential driver towards a larger size of plankton, increasing the sinking  
301 speed and efficiency of this part of the biological pump, and ventilating the water column.

302 The invasion of the pelagic realm by eumetazoan zooplankton provides the next step  
303 towards a modern-style biological pump. These zooplankton would have further cleared surface  
304 waters and contributed faecal pellets to organic aggregates sinking to the deep ocean, and also  
305 increased oxygen levels at depth [e.g. 6,54]. The small shelly fossil (SSF) record provides a source of  
306 evidence for the invasion of the pelagic realm by eumetazoans. Terreneuvian SSFs include the  
307 possible chaetognath *Protoherzina* and the molluscs *Watsonella*, *Aldanella*, and *Oelandiella*, whose  
308 widespread distributions are suggestive of a pelagic lifestyle, or at least a planktonic larval stage (**Fig.**  
309 **5**) [56]. Euarthropods, likely early occupants of the plankton [35], are represented in the SSF record  
310 from the Cambrian Stage 3 by millimetre scale bradoriids among others (**Fig. 5**) [e.g. 47,57,58]. The  
311 first macroscopic nektonic suspension feeders, such as the radiodont *Tamisiocaris*, also appear at  
312 this time [59], while the first centimetre-scale phytoplanktivores are identified close to the Stage 3–4  
313 boundary (**Fig. 5**) [12]. These data suggest that there was an increase in the diversity of millimetre-  
314 scale zooplankton at or close to the base of Stage 3, very close in time to the appearance of the first  
315 vertical migrants. Most bradoriids are considered benthic, however *Anabarochilina* increased its

316 distribution in three phases, providing complementary evidence for a steady strengthening of the  
317 pump during the Cambrian. In Epoch 2 *Anabarochilina* was coupled with benthic assemblages, by  
318 the Wuliuan it spread to a wider spectrum of lithofacies, and by the Guzhangian two species became  
319 widely distributed [48].

### 320 Significance of vertical migration for the early Cambrian radiation of animals

321 The metazoan innovation of vertical migration would have impacted both demersal and  
322 pelagic communities. The strength of the impact depends on the amount of biomass undertaking  
323 vertical migration. Models based on Cambrian environmental parameters predict that vertical  
324 migration would have increased the efficiency of the carbon pump by around 7% [60], however,  
325 more significantly, vertical migrants transport organic nutrients to the deep sea more quickly than  
326 aggregates, with a different nutritional balance, and repackage decaying sinking organic matter (**Fig.**  
327 **1**) [1,13,61–64]. In addition, vertical migrants are major contributors to ocean mixing and  
328 ventilation, spreading oxygen and nutrients throughout the water column [e.g. 54,65]. These effects  
329 likely played a role in contributing to the rapid rate of diversification during the Cambrian explosion,  
330 interwoven with numerous evolutionary and ecological feedbacks. For example, the higher  
331 metabolic needs and nutrient requirements of large biomineralizing animals and motile predators  
332 [66] may have been facilitated by increased quality and quantity of nutrient transport in the  
333 biological pump, and resulted in increased oxygenation of bottom waters. In turn, the increasing size  
334 and motility of predators would have provided a further ecological pressure for animals to ‘escape’  
335 into the pelagic realm.

336 The establishment of a biological pump with a modern-style architecture by the Cambrian  
337 Stage 3 does not mean that the fluxes along the Aggregation and Vertical Migration vectors (**Fig. 1**)  
338 remained constant to the modern day. Indeed it likely strengthened through the Palaeozoic with an  
339 increase in biomass (from an increased number of taxa, individuals, and size of individuals). Fossil  
340 evidence points to additional metazoan innovations during the Palaeozoic that would have affected  
341 the fluxes along these vectors and strengthened the pump. For example, the Aggregation vector  
342 would have been strengthened following the evolution of centimetre-scale and decimetre-scale  
343 phytoplanktivores later in the Cambrian [12,67], and the major radiation of plankton and the  
344 evolution of metre-scale nektonic suspension feeders during the Great Ordovician Biodiversification  
345 Event [68–70]. The flux of nutrients along the Vertical Migration vector would have increased as  
346 pelagic and vertically migrating animals diversified and increased in size – an innovation that would  
347 also have increased the mixing of waters and ocean ventilation. For example, the evolution of large,

348 fast moving fish as part of the Devonian nekton revolution is expected to have been especially  
349 significant [54,71].

350 In summary, the innovation of vertical migration in some *Isoxys* species was one of several  
351 interwoven and coevolutionary feedbacks during the early Cambrian that increased the efficiency  
352 and altered the architecture of the biological pump, likely contributing to the rapid expansion in  
353 metazoan diversity at this time.

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364 **Supplementary materials**

365 Are available through the Open Science Framework: [doi.org/10.17605/OSF.IO/2JDRS](https://doi.org/10.17605/OSF.IO/2JDRS)

366 **Author contributions**

367 SP conceived the study. SP and DAL collected data on *Isoxys* carapaces. SP conducted the outline  
368 analyses. SP conducted the CFD analyses with input from IAR. SP drafted the article and figures. All  
369 authors interpreted the data, critically revised the article, and approved the final version.

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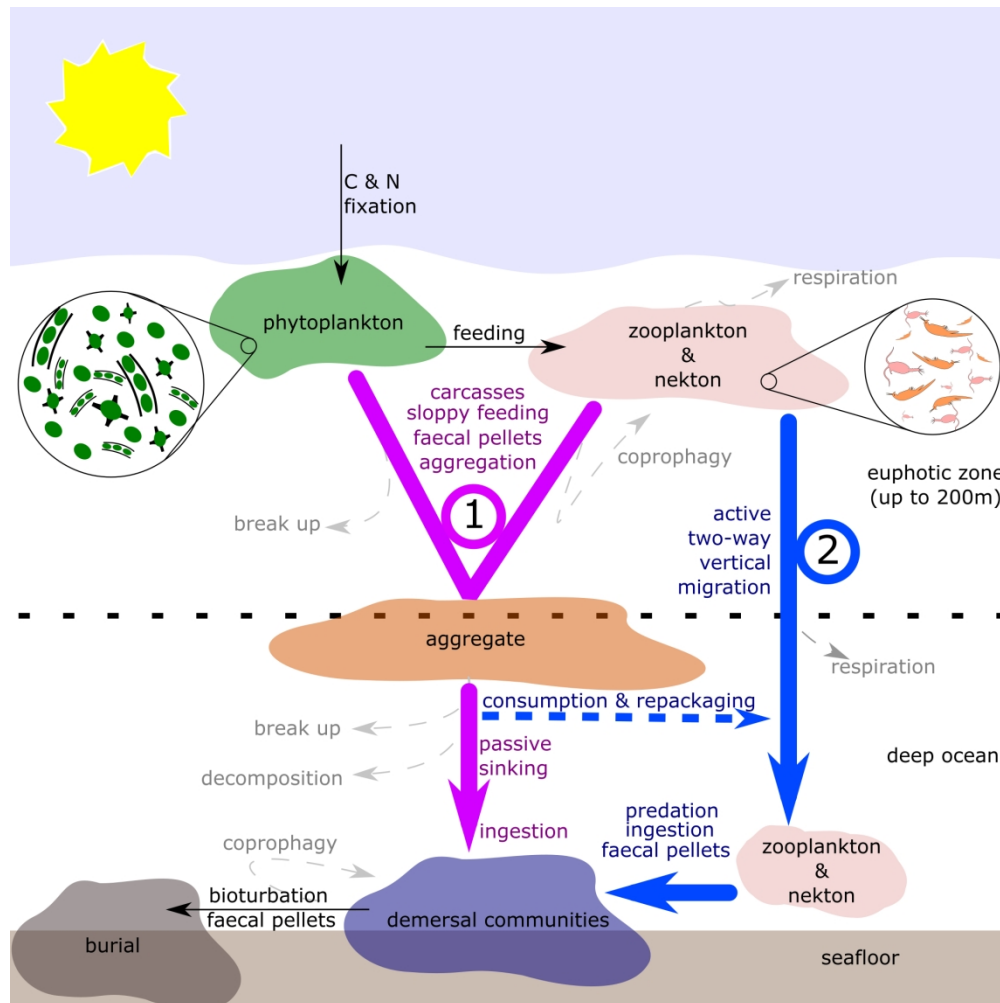


Figure 1. Simplified architecture of the modern biological pump. Arrows trace pathway of carbon (C) and nitrogen (N) and other nutrients. Numbers indicate two main biologically mediated vectors that transport nutrients fixed by phytoplankton from the surface ocean to demersal communities. 1) Aggregation vector of phytoplankton, faecal pellets, and other organic matter which sinks passively through the water column. 2) Vertical Migration vector driven by active two-way migration by metazoans.

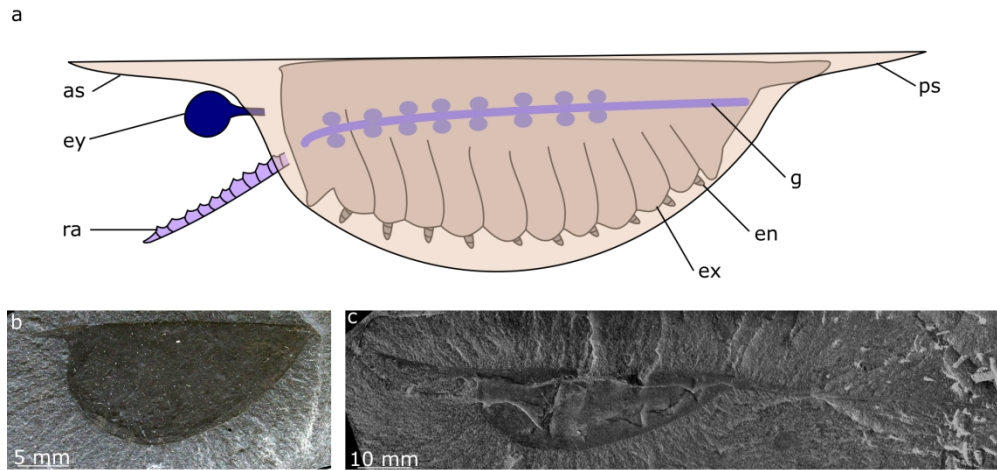


Figure 2. Morphology of *Isoxys*. (a) line drawing of idealised *Isoxys* illustrating known soft parts. (b) YPM IP 005804, *Isoxys acutangulus* from the Burgess Shale, British Columbia, Canada (Cambrian: Wuliuan) (credit: W. K. Sacco). (c) USNM PAL 189170, *Isoxys longissimus* from the Burgess Shale, British Columbia, Canada (Cambrian: Wuliuan). Image courtesy of the Smithsonian Institution (EZID: <http://n2t.net/ark:/65665/m372f2a644-97c3-441c-87e2-24b1dcc2e8c>, credit: Xingliang Zhang). Abbreviations: as, anterior spine; en, endopod; ex, flap like exopod; ey, eye; g, gut with paired diverticulae; ps, posterior spine; ra, raptorial appendage.



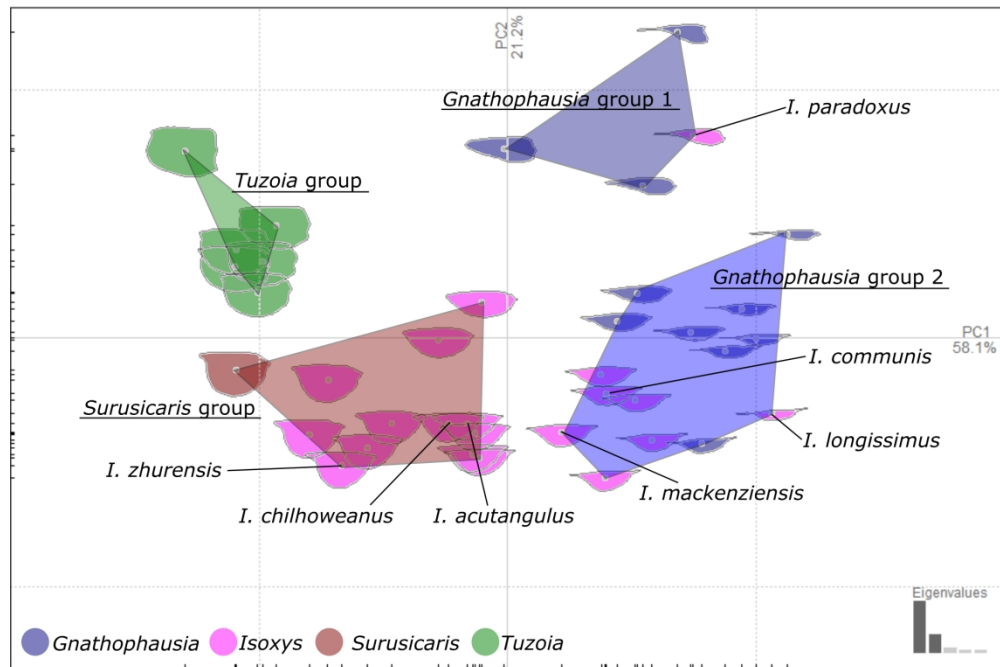


Figure 3. Principal Component Analysis of results of elliptical Fourier analysis conducted on the outlines of 11 species of Gnathophausia, 20 Isoxys, 1 Surusicaris, and 6 Tuzoia. Convex hulls indicate optimum four groupings as recovered by clustering analysis. Labelled Isoxys species chosen for subsequent hydrodynamic analysis.

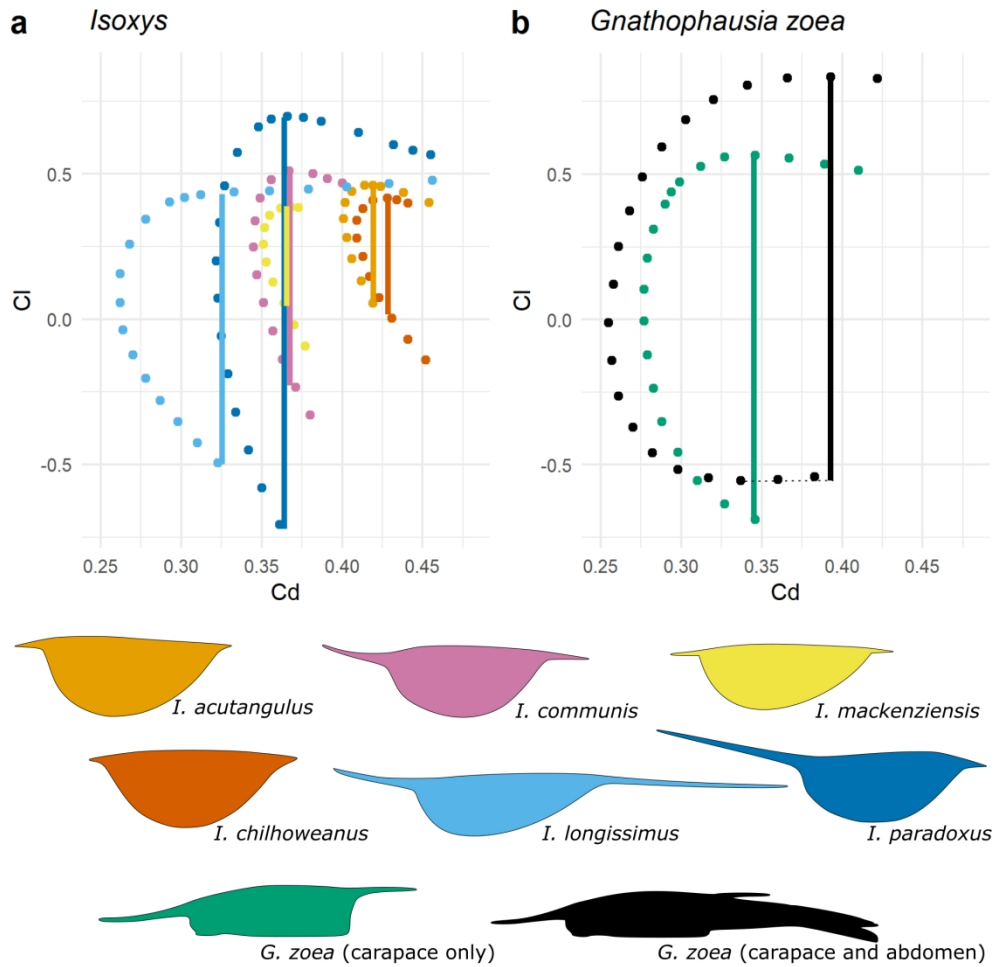


Figure 4. Drag polars (plot of  $C_d$  against  $C_l$ ) of taxa analysed at  $Re=255$  (0.75 body lengths per second for an animal 25 mm long). Each point corresponds to a single simulation at a different angle of attack. Vertical bars show range of lift coefficients. Note that flow was unsteady for *Isoxys zhurensis* at  $Re=255$ , and so no quantitative lift or drag coefficients were recorded. Drag polars at faster flow speeds and raw data presented in Supplemental Materials 3.

