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# 1 A novel tool to untangle the ecology and fossil preservation knot

# 2 in exceptionally preserved biotas

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#### 18 Abstract

19 Understanding the functioning of extinct ecosystems is a complicated knot of ecological, 20 evolutionary, and preservational strands that must be untangled. For instance, anatomical and 21 behavioral differences can profoundly alter fossilization pathways. This is particularly true in 22 exceptionally preserved soft-bodied biotas that record the earliest phases of animal evolution 23 during the Cambrian Explosion and the Ordovician Radiation. Herein, a novel method of data 24 partitioning based on probabilistic modelling is developed to examine these processes for the 25 Walcott Quarry, Burgess Shale, Canada (510Ma), and the Fezouata Shale, Morocco (c. 26 475Ma). The modelling shows that the mechanism for soft-tissue preservation in the Walcott 27 Quarry is ecologically selective, favoring the endobenthos. This is not found in the Fezouata 28 Shale. Taken in concert with bioturbation data, a new model of comparative preservation is developed based on sedimentary flow dynamics. This suggests that during the Cambrian 29

- Explosion and Ordovician Radiation the most exceptional fossils sites must still be calibrated
  against each other to understand the unfolding evolutionary events and the ecological
  structuring of ancient animal communities.
- 33
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#### 36 1. INTRODUCTION

37 Early animal communities from the Cambrian and the Ordovician are some of the most 38 complete in the fossil record, with exceptional fossilization of soft tissues giving a 39 comprehensive view of faunal diversity during the Cambrian Explosion and the Ordovician 40 Radiation. The Walcott Quarry (Miaolingian, Wuliuan, Cambrian, Burgess Shale, Canada) 41 bears a diverse assemblage of soft-bodied animals preserved in minute details (Nanglu et al., 42 2020). More than 100 genera have been described from the Walcott Quarry, helping to 43 resolve the enigma of metazoan origins in the Cambrian (Daley et al., 2018; Nanglu et al., 44 2020). The Fezouata Shale offers a more complete view of the transition between the 45 Cambrian Explosion and the Ordovician Radiation than was previously known (Van Roy et 46 al., 2010).

47 The Walcott Quarry and the Fezouata Shale share the same Burgess Shale-type (BST) 48 mode of preservation defined by carbonaceous compressions (Gaines et al., 2008; Saleh et al., 49 2020a). Accessory authigenic mineralization (i.e., pyritization, phosphatization) may occur 50 but this remains secondary to carbonaceous compressions (Gaines et al., 2008; Gaines, 2014; 51 Saleh et al., 2019, 2020b-c). Preservation was aided by rapid sediment influx (Gaines, 2014). 52 Clay minerals may have facilitated the exceptional preservation of soft tissues by slowing 53 down bacterial decay (McMahon et al., 2016; Anderson et al., 2018, 2020; Saleh et al., 2019). 54 The general conditions for BST preservation are broadly shared between the Fezouata Shale 55 and the Walcott Quarry (Saleh et al., 2020a); yet important abiotic and biological differences 56 exist. For instance, the Fezouata Shale is a wave-dominated shallow-marine polar 57 environment (70-100m), whereas the Walcott Quarry is deposited at the bottom of an 58 escarpment in a tropical sea (Martin et al., 2016; Vaucher et al., 2016, 2017; Saleh et al., 59 2021a). While the Fezouata Biota shares some taxa with Cambrian ecosystems (e.g., 60 marrellomorphs, radiodonts, and lobopodians), the general faunal composition of the 61 Ordovician site is markedly different from that of the Walcott Quarry (Van Roy et al., 2015a; 62 Lefebvre et al., 2016, 2019).

Assemblage discrepancies between fossil sites are influenced by many factors: ecology,
evolution, preservation, and even collection bias (Brasier et al., 2011). This is a broad-scale

problem in palaeontology that impacts all time intervals across the whole gamut of 65 66 evolutionary scenarios from the origin of life to the diversification of hominids (Saleh et al., 67 2020b). Using abundance data and alpha diversity metrics is the traditional approach for 68 palaeoecological reconstructions (Caron and Jackson, 2008; Saleh et al., 2018; Whitaker and 69 Kimmig, 2020). However, at many sites, these data can be skewed by collection bias caused 70 by government rules, economic value, and scientific interest (Brayard et al., 2011). Even with 71 rarefaction indices (Caron and Jackson, 2008; Nanglu et al., 2020), it is almost impossible to 72 quantify the effect that these biases have on raw taxon abundance data. When trying to 73 understand fossil preservation, a better approach is one based on occurrences rather than 74 abundances as some taxa have tissues that are more likely to be preserved (e.g., mineralized 75 brachiopods) in the fossil record than other taxa (e.g., soft jellyfish), introducing a 76 preservational bias. Moreover, because most taxa have more than one tissue type, it is best to 77 separate the preservation variable by comparing tissue types between fossil sites. Recent 78 probabilistic modelling of genus distributions and combinations of preserved tissue types has 79 shown considerable advantages over traditional approaches (Saleh et al., 2020b). Probability 80 models showed the preservation of anatomical structures in the Walcott Quarry and the 81 Fezouata Shale are significantly different (Saleh et al., 2020b), and the Fezouata Shale did not 82 preserve entirely cellular organisms (e.g., the body walls of chordates; internal organs such as 83 digestive and nervous tissues) (Saleh et al., 2020b). It is likely the animals were dead and 84 decaying on the seafloor prior to their burial in the Fezouata Shale (Saleh et al., 2021b), and 85 cellular structures in contact with the water column are the fastest to decay (MacGabhann et 86 al., 2020; Saleh et al., 2020b). Herein, this methodology is expanded upon by incorporating 87 data on ecological strategies, to investigate the effect of habitat on fossilization. Consequently, the interplay of taxonomy, preservation, and ecology is analytically 88 89 approached using the Walcott Quarry and the Fezouata Biota as separate, and then 90 comparable case studies. This quantification allows a separation of ecological and 91 preservation effects helping to form a clearer picture of the dawn of animal life in the early 92 Palaeozoic.

93

# 94 2. MATERIAL AND METHODS

95 The data matrices for the generic composition and biological tissue occurrences [i.e., 96 biominerals (A) such as brachiopod shells, sclerites (B) such as the headshield of some 97 arthropods, cuticle formed of polysaccharides (C) such as the body walls of priapulids, 98 cellular body walls (D) as in cambroernids, and internal organs (E) such as digestive and 99 nervous systems] in the Walcott Quarry and the Fezouata Shale were taken from a previously 100 published database (Saleh et al., 2020b) based on the collections of the University of Lyon 101 (France), the University of Marrakech (Morocco), and the University of Lausanne 102 (Switzerland), the Royal Ontario Museum (Canada), and the Peabody Museum (USA). Taxa 103 were separated between mode of life categories (i.e., endobenthic, nektobenthic/epibenthic, 104 planktonic/nektonic) based on the literature (Caron and Jackson, 2008; O'Brien and Caron, 105 2015; Nanglu et al., 2020) and our own experience. Raw data is provided in the 106 Supplementary Material 1.

107 The number of times each of the different possible tissue type combinations occurred 108 was identified. Tissue types can occur alone without other tissues as A, B, C, D, or E; and in 109 one of ten possible pairs: AB, AC, AD, AE, BC, BD, BE, CD, CE, and DE. There are also 10 110 different possible three-set intersections ABC, ABD, ABE, ACD, ACE, ADE, BCD, BCE, 111 BDE, and CDE. Finally, there are five different possible associations of four-set intersections 112 ABCD, ABCE, ABDE, ACDE, BCDE, and one five-set combination ABCDE, for a total of 113 31 possible combinations. A simple calculation of the number of taxa belonging to each 114 category introduces a bias in the dataset because each count of a higher-order intersection 115 leads to multiple counts of lower-order intersections. For example, finding an ABC 116 combination causes a count of the AB, AC, and BC categories and consequently, one taxon is 117 counted 4 times. This has to be removed from the data in order to find unique character 118 combinations. This problem can be solved on a five-set Venn diagram and the unique 119 associations can be obtained (Table 1). Automating the process for a five-fold set problem is 120 challenging and software and computational programs are generally limited to solving single

121 four-set problems (Chen and Boutros 2011) or solving the five-set problem indirectly by 122 moving to probability densities of multiset intersections (Wang et al. 2015). It is worth noting 123 here that two of the characters in this analysis are biologically mutually exclusive (C and D; 124 an organism cannot be cuticularized and non-cuticularized) so the problem simplifies 125 somewhat to two intersecting four-fold problems, as CD and all its subsets are empty. 126 Consequently, the five-way intersection ABCDE is impossible. Further subsets of  $C \cap D$  that 127 are also empty are  $A \cap C \cap D$ ,  $B \cap C \cap D$ ,  $C \cap D \cap E$ ,  $A \cap C \cap D \cap E$ ,  $A \cap B \cap C \cap D$ , and  $B \cap C \cap D \cap E$ . 128 Hence, the problem is reduced from 31 to 23 intersections and the highest intersections that 129 can have a value are  $A \cap B \cap C \cap E$  and  $A \cap B \cap D \cap E$  and these can be found directly in the 130 dataset. These values can then be used to calculate the number of taxa that preserve three 131 tissue types because of the identity: 132  $X \cap Y = X \cap Y \cap Z' + X \cap Y \cap Z$ 133 With Z' being everything but Z  $(X \cap Y \cap Z')$  is the association of X and Y with any other tissue

133 With Z' being everything but Z(X||Y||Z') is the association of X and Y with any other tissue 134 type except Z). The equation can be rearranged as:

135  $X \cap Y \cap Z' = X \cap Y - X \cap Y \cap Z$ 

136 Thus, knowing a value for  $A \cap B \cap C \cap E$  and for instance  $A \cap B \cap C$  we can calculate the desired 137 value of  $A \cap B \cap C \cap E'$ . Using the identity:

138  $A \cap B \cap C = A \cap B \cap C \cap E + A \cap B \cap C \cap E'$ 

139 Which can be simply rearranged to:

$$A \cap B \cap C \cap E' = A \cap B \cap C - A \cap B \cap C \cap E$$

In this equation, an unknown exclusive three tissue type association  $(A \cap B \cap C \cap E')$  can be calculated from two known quantities which can be recovered directly from the database. It is then trivial to extend this to all the other possible three variable intersections with the corollary that some intersections require the subtraction of both of the four variable intersections if the three-set intersection contains subsets of both of the four variableintersections. For instance:

147

# $A \cap B \cap E \cap C' \cap D' = A \cap B \cap E - (A \cap B \cap C \cap E + A \cap B \cap D \cap E).$

Once the unique associations of three variables are known they can be used, following similar logic, to calculate all the unique two-variable intersections (e.g.,  $A \cap B \cap C' \cap D' \cap E'$ ) and finally, the one variable intersections (e.g.,  $A \cap B' \cap C' \cap D' \cap E'$ ).

A manual approach would be highly prone to error due to the high number of taxa and data entries. Thus, this method has been automated in a single Excel file that contains all of the formulae above such that future analyses need only to construct the data matrix, with data analyses proceeding automatically. This approach is quicker and more reliable, providing numerous opportunities to cross-check the data: the sum of the values of all 23 subsets; equal number of taxa for all calculations; no empty taxa coding; C $\cap$ D intersections remain zero.

157 The data is then analyzed as a discrete probability distribution. The total number of 158 genera having just one character (i.e., A, or B, or C, or D, or E) was plotted against the 159 number of genera that have pairs (e.g., AB), threes (e.g., ABC), or fours (e.g., ABCE) for all 160 modes of life in the Walcott Quarry and the Fezouata Shale (Fig. 1a, b). The probability of 161 occurrences P(A), P(B), P(C), P(D), and P(E) was calculated (Table 1) by taking the set value 162 for "A" and dividing by n to obtain P(A). The average number of tissue types per genus, as 163 derived from the dataset, was then calculated by adding these probabilities for each mode of 164 life in each biota (Table 1).

To examine how exceptional preservation differs between modes of life, further analyses focused on the preservation of soft tissue (E) because all animals have internal tissues and so this category must have been present during the life of the organism (whereas this is not true for all the other categories, for example, it is possible to have no biominerals). The association of soft internal organs (E) with other structures was investigated for all tissuetype categories in the Fezouata shale and the Walcott Quarry. This is achieved by examining 171 conditional probabilities in the subset of A. For example, the probability that a taxon with a172 biomineral also having preserved internal organs/tissues (p(EIA)).

173

$$P(E|A) = (A \cap E + A \cap B \cap E + A \cap C \cap E + A \cap D \cap E + A \cap B \cap D \cap E + A \cap B \cap C \cap E)/A$$

The reverse conditional approach of finding A given that E has occurred p(AlE) was also calculated, following a similar logic (automated in the Supplementary Material 2).

176 In order to investigate the role of mode of life on fossil preservation the data was then 177 partitioned using mode of life categories. This allows a direct link between exceptional 178 preservation of internal tissues (E) to a specific mode of life category (MoL). Therefore 179 P(ElMoL) is the probability of finding soft internal organs (E) given a certain mode of life is 180 required (Table 1). Then, for the Fezouata Shale and the Walcott Quarry separately, the 181 likelihood of producing the distribution of combinations of soft tissues (E) with a specific 182 mode of life assuming that the endobenthos, for example, has the "true" preservation regime 183 was investigated using a binomial distribution  $P(x \ge n)|Bi(n, p)$ , with:

184

$$P(x) = \binom{n}{x} p^{x} q^{n-x} = \frac{n!}{(n-x)! \, x!} p^{x} q^{n-x}$$

185

186 In the Walcott Quarry, p=P(Elendobenthic), q=1-p, n is the number of 187 epibenthic/nektobenthic genera preserving an internal organ (E), and x is the total number of 188 taxa that could potentially preserve E which is, in this case, the total number of 189 epibenthic/nektobenthic genera in the Walcott Quarry (because they all possessed internal 190 organs during life). All calculated probabilities are summed and the probability  $P(x \ge n)|Bi(n, x)| = 0$ 191 p), of producing the value of P(Elnektobenthic) considering that internal organ preservation in 192 the endobenthic community P(Elendobenthic) is the "true" distribution is then obtained, 193 giving the critical interval of the distribution. This was then also performed for 194 nektonic/planktonic taxa, and then repeated entirely for the Fezouata Shale.

To provide environmental context, the bioturbation index (BI) was measured (Taylor and Goldring, 1993) for one interval from each site that yielded diverse exceptionally preserved fossils – the phyllopod bed from the Walcott Quarry, and Z-F4m from the Fezouata

198 Shale. BI is equal to zero if bioturbation is absent. BI=1 if the percentage of bioturbation is 199 between 1 and 4% with distinct bedding, few discrete traces and/or escape structures (Taylor 200 and Goldring, 1993). BI=2 if bioturbation percentage is between 5 and 30% with low trace 201 density and common escape structures (Taylor and Goldring, 1993). BI=3 if between 31 and 202 60% of the sediment is bioturbated with a rare overlap of traces (Taylor and Goldring, 1993). 203 BI=4 if bioturbation is high (61-90%) with a common overlap of traces and primary 204 sedimentary structures are mostly erased (Taylor and Goldring, 1993). BI=5 is characterized 205 by intense bioturbation (91-99%) (Taylor and Goldring, 1993) and sediment with almost 206 completely disturbed bedding. BI=6 when the sediment is fully bioturbated (Taylor and 207 Goldring, 1993).

208 Table 1. Summary statistics for the Walcott Quarry and the Fezouata Shale. A = biominerals, B = sclerites, C = 209 cuticle, D = cellular body walls, and E = internal organs. Numbers of genera belonging to each category (e.g., A, 210 AB, ABE) for each mode of life in all exceptionally preserved biotas are summarized. The proportion of each 211 type of tissue in all categories (e.g., P(A), P(C)) are calculated. The sum of these probabilities shows the number 212 of tissues per taxon. Probabilities of finding internal tissue in association with other structures (e.g., P(AIE); 213 P(EIB)) are calculated as well. For each site and for a certain probability (e.g., P(EIB), water query), comparable numbers 214 between modes of life are highlighted in red. Note that for the Walcott Quarry, the obtained numbers for the 215 epibenthic/nektobenthic and the nektonic/planktonic communities are more similar to each other than to the 216 endobenthos. There is no clear pattern between modes of life in the Fezouata Shale. The probability of finding

	Walcott Quarry			Fezouata Shale			
	Endobenthic	Epibenthic Nektobenthic	Nektonic Planktonic	Endobenthic	Epibenthic Nektobenthic	Nektonic Planktonic	
	Tissue-type occurrences						
N	12	77	12	6	131	24	
А		14		4	82	5	
В		5	2		18	15	
С				1	2		
D			1				
E							
AB		6			3		
AC		2					
AD		1			1		
AE		1			8	1	
BC		7	1		7		
BD		1					
BE		2			2	1	
CE	3	1					
DE	1	9	2				
ABC		2					
ABE					5		
ACE	6	2		1			
BCE	2	20	6		4	2	
BDE		2					
ABCE		1			3		
ABDE		1					
	Proportion of each tissue-type						
P(A)	0,5	0,38	0	0,83	0,77	0,25	
P(B)	0,17	0,61	0,75	0	0,3	0,75	
P (C)	0,92	0,45	0,58	0,33	0,12	0,08	
P(D)	0,08	0,18	0,25	0	0	0	
P (E)	1	0,5	0,67	0,17	0,17	0,17	
N(tissue/taxon)	2,66	2,14	2,25	1,33	1,36	1,25	
<b>D</b> ((1))		Proba	abilities of tis	sue co-occurren	ices	0.44	
P(AIE)	1	0,16		0,2	0,15	0,16	
P(BIE)	1	0,55	0,66	0	0,35	0,116	
P(CIE)	1	0,71	0,85	0,5	0,43	1	
P(DIE)	1	0,85	0,66	0	0	0	
P(EIA)	0,5	0,12	0	1	0,72	0,25	
P(EIB)	0,16	0,74	0,75	0	0,63	0,75	
P(E C)	0,9	0,64	0,75	1	0,31	0,5	
P(EID)	0,08	0,3	0,25	0	0	0	
	Probabilities of internal organ association with a certain mode of life						
P(EIMoL)	1 1	0.5	0.67	0.17	0.17	0.17	

217 internal organs considering that a certain mode of life has been found in also investigated (e.g., P(ElMoL)).

# 218 **3. RESULTS**

219 In general, most tissue types exist in all mode of life categories (Table 1). However, a few 220 exceptions exist. In the Fezouata Shale, cellular structures in direct contact with seawater (D) 221 are not found in any of the animal groups (Table 1), and sclerotized structures (B) are absent 222 from the endobenthic community (Table 1). Biominerals (A) are not present in the nektonic 223 community of the Walcott Quarry (Table 1). In the Fezouata Shale all mode of life categories 224 show comparable patterns of preservation (Fig. 1a) with a similar number of tissues per taxon 225 (Table 1). The endobenthic, epibenthic/nektobenthic, and nektonic/planktonic communities 226 preserved 1.33, 1.36, and 1.25 tissue per taxon respectively (Table 1). In the Walcott Quarry, 227 there are generally more tissues preserved per taxon than in the Fezouata Shale, and these 228 values show more variation between the three modes of life (Fig. 1b). Endobenthic taxa in the 229 Walcott quarry preserve the highest number of tissues per taxon (i.e., 2.66 tissues per taxon, 230 with the epibenthic/nektobenthic and the nektonic/planktonic communities 2.14 and 2.25 231 tissues per taxon respectively) (Table 1). In the Walcott Quarry, internal tissue associations 232 (E) with other biological structures (A, B, C, and D) are more similar between 233 nektonic/planktonic and epibenthic/nektobenthic communities than to endobenthic taxa (red 234 highlight in table 1). The entire endobenthic community preserved internal organs while it 235 was only a half for epibenthic/nektobenthic, and a third for nektonic/planktonic taxa (Table 236 1). The endobenthic community is significantly different from both the 237 epibenthic/nektobenthic (i.e.,  $P = 9.19 \times 10^{-5}$ ) and the nektonic/planktonic (i.e.,  $P = 4.56 \times 10^{-5}$ ) 238 communities. The latter two habitats are not statistically significantly different from each 239 other (i.e., P= 0.19). In the Fezouata Shale, less than one-fifth of all animal communities 240 preserved internal organs (Table 1, P(ElMoL)). There is no clear pattern of similarity between 241 the three mode of life communities in terms of internal tissue associations with other 242 biological structures (Table 1). Moreover, there are no significant differences in tissue 243 associations between the three mode of life categories in the Fezouata Shale (i.e., 244 endobenthic-epibenthic/nektobenthic (P= 0.56), endobenthic-nektonic/planktonic (P=0.61), 245 and epibenthic/nektobenthic-nektonic/planktonic (P=0.61)).

The two units also markedly differ in terms of bioturbation, with the Walcott Quarry being essentially unburrowed (i.e., BI= 0) (Fig. 1c), whereas the Fezouata Shale is extensively bioturbated (i.e., BI= 4-5) (Fig. 1d) containing *Taenidium* (Ta), *Rhizocorallium* (Rh), *Phycosiphon* (Ph), *Nereites* (Ne), and *Arenicolites? (Ar?)* (Fig. 1d). In the Fezouata Shale discrete trace fossils are overprinted to an undifferentiated burrow mottling, indicative of a tiering structure and a well-developed mixed layer (Fig. 1d).

252



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Figure 1. Differences in proportions of genera (Y-axis) between single (e.g., A, B, C), paired (e.g., AB, BC,
CE), triple (e.g., ABC, ACE) and quadruple (e.g., ABCE) character categories (marked as 1s, 2s, 3s, and 4s on
the X-axis) and modes of life in the Fezouata Shale (a), and the Walcott Quarry (b). Bioturbation index (BI) for
the undisturbed Walcott Quarry sediments (c) and the heavily bioturbated Fezouata Shale (d) which containing
Taenidium (Ta), Rhizocorallium (Rh), Phycosiphon (Ph), Nereites (Ne), and Arenicolites? (Ar?).

259

# 260 4. DISCUSSION

#### 261 4.1. Linking ecology to preservation

The absence of biominerals (A) from the nekton/plankton of the Walcott Quarry (Table 1) reflects the original absence from this site, as there is no evidence of nektonic/planktonic taxa with mineralized bodies during the Middle Cambrian (Klug et al., 2010). Nektonic/planktonic animals at this time consisted mainly of non-mineralized taxa, such as ctenophores and arthropods (Hou et al., 2004; Vinther et al., 2014; Lerosey-Aubril and Pates, 2018; Whalen and Briggs, 2018; Daley, 2019; Zhao et al., 2019). After accounting for this evolutionary
signal, the taphonomic signal still exists for internal tissue and the three mode of life
categories (Table 1).

270 In the Walcott Quarry, different mode of life categories preserved contrasted 271 proportions of internal organs (Table 1), even though these organs are equally present in all 272 animals. These difference in preservation between habitats can be explained by the 273 depositional regime. Classical interpretations suggest organisms were transported from their 274 habitat to the environment where they were preserved (Conway Morris, 1986; Gaines, 2014). 275 In this scenario, nektonic, planktonic, epibenthic, and nektobenthic taxa were not affected by 276 obrution events in the same way as the endobenthos. Many endobenthic taxa construct 277 permanent domiciles and are able to vertically adjust their burrows accordingly for both 278 continuous (equilibrium traces) or episodic (escape traces) sedimentation. However, the 279 pristine preservation of the primary fabric in Walcott Quarry shows the thickness of the 280 emplaced sediment was sufficient to prevent the transported endobenthos from escaping (Fig. 281 2a). Furthermore, even if endobenthic animals were dead and decaying prior to their transport, 282 their decay would have been less than in other animal communities because the body of 283 endobenthic taxa is naturally at least partially buried and has less contact with seawater. 284 Therefore, the preserved endobenthic community in the Walcott Quarry is dominantly formed 285 of organisms that were entrained alive and/or fresh carcasses explaining the soft tissue 286 preservation (Fig. 2a). In contrast, nektonic taxa that live high in the water column were 287 above the incoming sediment flow (Fig. 2b), and might have been able to escape. This 288 hypothesis can explain the rare occurrence of complete organisms belonging to taxa such as 289 radiodonts in the Walcott Quarry (Daley et al., 2009, 2013). Though radiodonts are diverse in 290 the Burgess Shale, complete carcasses such as the nektonic predator, Anomalocaris, are 291 extremely rare, and fossils mainly represent molting products or decomposed carcasses 292 (Daley et al., 2009, 2013, 2014). Hurdia is thought to swim close to the seafloor and is ten 293 times more abundant at this locality than Anomalocaris (Daley et al., 2009, 2010, 2013, 294 2014). Molting products and carcasses fall to the seafloor and are passively transported by 295 event deposits to the preservation site (Fig. 2c), and taxa living low in the water column have

296 less chance of escaping the incoming sediment gravity flow. The rarer preservation of internal 297 organs for the nektonic community in comparison to the endobenthos cannot be solely linked 298 to taxonomic discrepancies between these categories even if the nektonic community had 299 originally more molting animals than the endobenthic one. In our database, the best-preserved 300 fossil for each genus was inventoried (i.e., for a specific genus, even if more than 100 molts 301 have been found, as soon as one specimen shows internal organs, this genus is inventoried as 302 preserving internal organs, such as the case for Anomalocaris). Thus, another aspect of the 303 data is that for 33% of nektonic genera (P(ElMoL)= 0.67; Table 1), no single animal was 304 captured alive or at the same stage of decay as endobenthic taxa in the Walcott Quarry. The 305 absence of complete carcasses of some nektonic taxa and the abundance of their molting 306 products could be a body size effect with a natural weight limit on entrainment in the gravity 307 flows. Further investigation of flow dynamics may help resolve these issues.

308 The most intriguing part of our data is that epibenthic animals in the Walcott Quarry 309 show a similar pattern of preservation to nektonic ones (red highlight Table 1). The fact that 310 epibenthic taxa show a lower degree of preservation in comparison to the endobenthos (Table 311 1) suggests that some epibenthic taxa were able to either escape (1) burial sediments during 312 transport or (2) before being entombed. The Walcott Quarry deposits were initially interpreted 313 as the products of dilute turbidity currents (Piper, 1972). However, recent studies have shown 314 deposition from pulsating, dense, mud-rich slurry flows, transitional between turbidity and 315 debris flows (Gabbott et al., 2008). These currents are extremely fast and involved 316 progressive aggradation rather than en-masse freezing. Mud-rich slurry flows may have been 317 triggered by physical instabilities, such as delta front collapse, storms, or earthquakes (Allison 318 and Brett, 1995; Gabbott et al., 2008). It is most likely that epibenthic taxa were able to sense 319 these instabilities and escape from the environment (Fig. 2d) rather than escaping from fast-320 moving flows once they are captured. The escape of animals prior to instabilities is well 321 documented (Grant and Halliday, 2010; Grant et al., 2011; Freund and Stolc, 2013) and this 322 behavior may have first evolved during the Cambrian (Kirschvink, 2000; Langenhof and 323 Komdeur, 2018). Escaping the environment prior to the arrival of event deposits has the same 324 implications on preservation as the avoidance of turbulences by nektonic taxa (both lead to

325 the non-preservation of living animals; Fig. 2b, d). In both cases, only decaying carcasses and 326 molting products are preserved (Fig. 2e) explaining the resemblance between epibenthic and 327 nektonic communities and their lower exceptional preservation potential when compared to 328 the endobenthos (Table 1). However, this explanation should not be generalized because if all 329 taxa from the epibenthic community were able to avoid transported sediments, the potential of 330 exceptional preservation would have been much lower in the Walcott Quarry. Currently, at 331 least 50% of living epibenthic/nektobenthic taxa were not able to avoid burial as they show a 332 similar pattern of preservation as the endobenthos (P(E|MoL) = 0.5). Therefore, the 333 mechanism for soft-tissue preservation in the Walcott Quarry is ecologically selective and 334 preserved animal communities in this site exhibit different stages of decay (Fig. 2f). The 335 endobenthic community is the least decayed (Fig. 2f) while the nektonic, nektobenthic, and 336 epibenthic preservation contain a mix of decaying carcasses and living organisms caught in 337 the flow. Animals that were entrained into the flow alive died either during transport or at the 338 time of deposition because no evidence of escape traces are present in the Walcott Quarry 339 (Fig. 1c) (Allison and Brett, 1995; Mángano et al., 2019).

340 The probability models confirm the results of previous sedimentary studies suggesting 341 that the mechanism of soft part preservation in the Fezouata Shale are not identical to those in 342 the Walcott Quarry (Martin et al., 2016; Vaucher et al., 2016, 2017; Saleh et al., 2020a, 343 2021a). All mode of life categories in the Fezouata Shale preserved a lower proportion of 344 internal organs than the Walcott Quarry (Table 1). This was previously attributed to the 345 relatively long exposure of carcasses to pre-burial decay in the Fezouata Shale (Saleh et al., 346 2020a, b) (Fig. 2g). This pre-burial decay explains as well the absence of cellular structures in 347 direct contact with seawater (D) from this site (Table 1). However, the major difference 348 between the Walcott Quarry and the Fezouata Shale is that all modes of life in the Ordovician 349 site share a comparable number of structures per taxon and the same proportion of internal 350 tissues (Table 1). This means that burial in the Fezouata Shale is not ecologically selective. In 351 this site, animals were buried in-situ during storms and were not transported (Fig. 2g) (Saleh 352 et al., 2021a, b). Furthermore, escaping the effect of this *in-situ* burial in the Fezouata Shale is 353 not limited to the nekton as there are abundant escape trace fossils cross-cutting event deposits (Saleh et al. 2020a). Even infaunal taxa, such as *Wosekella*, were not affected by the distal storms, as they adjusted their position in the sediments and continued to grow after the thin storm deposits (Saleh et al., 2018) (Fig. 2g). Furthermore, because the mechanism for soft-tissue preservation in the Fezouata Shale does not rely on transport, the size of an animal is not significant in defining what can be preserved (Fig. 2k). Large (articulated) carcasses of taxa such as the radiodont *Aegirocassis* can be found in the Fezouata Shale (Van Roy and Briggs, 2011; Gaines et al., 2012; Van Roy et al., 2015b) (Fig. 2h).

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363 Figure 2. Taphonomic scenarios in the Walcott Quarry and the Fezouata Shale. Walcott Quarry: endobenthic 364 taxa are entrained alive and cannot escape the produced mud-rich deposit and are freshly preserved (a). 365 Nektonic/planktonic taxa living high in the water column can avoid transported sediments (b). Molting products 366 fall passively to the seafloor, might get transported (c). Some epibenthic/nektobenthic animals might have been 367 able to sense physical disturbances and escape the environment prior to the arrival of burial material (d). 368 Carcasses of these taxa are preserved (e). The preservation of the endobenthos is higher than other animal 369 communities (f). Fezouata Shale: carcasses decay on the seafloor (g). When storms cause sediment influx, most 370 living animals escape resulting in the preservation of non-fresh carcasses for all mode of life categories (h).

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# 372 4.2. Implications in understanding Cambrian ecosystems

The ecologically-selective mechanism of preservation present in the Walcott Quarry is absent in the Fezouata Shale which might have wider evolutionary implications, particularly when comparing Cambrian BST deposits. For instance, epibenthic/nektobenthic and 376 nektonic/planktonic taxa, other than arthropods, such as ctenophores (e.g., Batofasciculus, Galeactena, Gemmactena, Maotianoascus, Sinoascus, Thaumactena) (Fig. 3a, b) and 377 378 Cathaymyrus, Haikouichthys, Myllokunmingia, Zhongjianichthys, chordates (e.g., 379 Zhongxiniscus) (Fig. 3c, d) belonging to the D category (i.e., cellular body walls), are 380 regularly reported from the Chengjiang Biota (Cambrian Stage 3, China) (Shu et al., 1996, 381 1999; Hou et al., 2004; Zhao et al., 2019; Saleh et al., 2020b), while they are much rarer in the 382 Walcott Quarry (i.e., 4 genera only) (Nanglu et al., 2020; Saleh et al., 2020b). Furthermore, 383 the Cambrian Stage 3 Qingjiang Biota (China) is apparently dominated by ctenophores and 384 medusoid cnidarians (Fig. 3e, f) (Fu et al., 2019). In the Walcott Quarry, cnidarians remain 385 scarce and are mainly represented by polypoid forms (e.g., Mackenzia) (Conway Morris, 386 1993) (Fig. 3g). These discrepancies probably result from the ecologically-selective 387 preservation bias for the Walcott Quarry. Preservation potential for ctenophores, chordates, 388 and medusoids is low in the Walcott Quarry as: (1) Organisms may avoid transported 389 sediments (Fig. 2b, d); (2) The preserved nektonic/planktonic community consists partially of 390 carcasses and molting products (Fig. 2c, e); (3) Cellular structures in direct contact with 391 seawater are among the fastest to decay (MacGabhann et al., 2019; Saleh et al., 2020b); and 392 (4) Some of these organisms have a long lifespan (Petralia et al., 2017) and do not molt in a 393 similar way to ecdysozoans. The preservation of a diverse ctenophore and chordate 394 assemblage in the Chengjiang Biota and their absence from the Walcott Quarry remains 395 poorly understood. A mechanism is required that would allow the death and fall of 396 nektonic/planktonic taxa from the water column to the sea floor increasing the chance for a 397 fresh carcass to be captured by the flow (e.g., oxygen fluctuations in the water column; 398 Hammarlund et al., 2017; Qi et al., 2018) (Fig. 3h). Moreover, anoxia may increase the rate of 399 decay for some cnidarian tissues (Hancy and Antcliffe, 2020). In this sense, preserved 400 cnidarians in the Qingjiang Biota might be correlated with more oxygen availability than in 401 the Walcott Quarry (Hancy and Antcliffe, 2020) (Fig. 3i). Geochemical and sedimentological 402 data is not yet available to evaluate these hypotheses for the Qingjiang Biota. Yet the 403 Chengjiang and Qingjiang Biotas are particularly important in completing our understanding 404 of Palaeozoic ecosystems owing to their preservation of both hard-to-preserve tissues (i.e.,

405 cellular) and ecologies (e.g., nektobenthic, nektonic). Moreover, even if the Walcott Quarry 406 preserves a lower generic richness of chordates than in the Chengjiang Biota (Fig. 3), one 407 should keep in mind that 57 specimens of *Pikaia* were discovered in the Walcott Quarry 408 (Nanglu et al., 2020), while less than 10 specimens are known for each chordate genus in the 409 Chengjiang Biota. Such numbers do not question the robustness of the probability model 410 presented in this study. On the contrary, the probability model gives a new and 411 complementary perspective. The abundance of certain chordate genera in the Walcott Quarry 412 can be explained by a specific ecology of these taxa (e.g., swimming closer to the seafloor), 413 that might have been different for other chordates (i.e., swimming higher in the water 414 column). Furthermore, Chordates from the Walcott Quarry are limited to only a specific level, 415 within the 55 cm fossiliferous interval. This local abundance of chordates could be explained 416 as well by a timely change in sedimentary flows within the Walcott Quarry.



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418 Figure 3. Ecological and preservational discrepancies between Cambrian Burgess Shale-type deposits. 419 Ctenophores from the Chengjiang Biota: (a) Galeactena hemispherica (YKLP13810, 5mm scale-bar) (Ou et al., 420 2015), and (b) Maotianoascus octonarius (RCCBYU10217; 1.5mm scale-bar) (Ou et al., 2015). Chordates from 421 the Chengjiang Biota: (c) Myllokunmingia fengjiaoa (RCCBYU-10200a; 1mm scale-bar) (Caron et al., 2010), 422 and (d) Zhongxiniscus intermedius (YIGSHef64682; 1mm scale-bar) (Huilin et al., 2010). (e) Medusoid 423 cnidarian (3mm scale-bar) and (f) ctenophore (4mm scale-bar) from the Qingjiang Biota (Fu et al., 2019). (g) 424 Polypoid cnidarian from Walcott Quarry (Mackenzia costalis; ROM61115; 12mm scale-bar) (Conway Morris, 425 1993). The preservation of a diverse ctenophore, chordate, and medusoid assemblage might have been facilitated 426 by regular oxygen fluctuations (Hammarlund et al., 2017; Qi et al., 2018) augmenting the chances of capturing a 427 fresh carcass in the Chengjiang Biota (h) and by oxygen availability reducing the decay rate of some cnidarians 428 in the Qingjiang Biota (i) (Hancy and Antcliffe, 2020).

429

#### 430 5. CONCLUSIONS

431 A new method using data partitioning in a probability model was developed to decode the 432 ecology of ancient ecosystems. By quantifying the role of mode of life in the preservation 433 processes it is possible to reconstruct ecological selectivity in the fossil record. The focus on 434 how sedimentary flows interact with different modes of life categories (biostratinomy) 435 complements previous studies focusing on early diagenesis (decay and mineralization). The 436 Walcott Quarry best preserved the endobenthos. Endobenthic animals in the Walcott Quarry 437 were captured alive and were not able to escape the flow resulting in their preservation in 438 minute details. Some active organisms living in the water column were able to avoid 439 transported sediments, and decaying carcasses and molting products of these taxa are 440 generally preserved. When compared to the Fezouata Shale, the Walcott Quarry preserved a 441 higher proportion of soft tissues and internal organs. However, in the Fezouata Shale, the 442 mechanism for preservation is not ecologically selective as endobenthic, 443 nektobenthic/epibenthic, nektonic/planktonic taxa are equally preserved. Only carcasses were 444 preserved in the Fezouata Shale, and many animals including endobenthic taxa were able to 445 escape if buried alive. When compared to other Cambrian deposits with abundant soft-tissue 446 preservation, such as the Chengjiang and the Qingjiang biotas, the Walcott Quarry is 447 underrepresenting some taxonomic groups such as ctenophores, chordates, and cnidarians. 448 This appears to be a result of preservation rather than evolutionary processes. Consequently,

449 the mechanism and conditions for soft-tissue preservation in the Cambrian are not universal 450 across all BST sites. Subtleties of each unique local setting have a significant effect on the 451 composition of the preserved fossil biota. Finally, it is crucial to emphasize that BST deposits 452 offer a unique opportunity to understand evolutionary events during the Palaeozoic. However, 453 there is no single fossil site that reflects an entirely accurate picture of past ecosystems. Fossil 454 data from numerous localities should be taken in concert and comparatively analyzed, to 455 untangle the preservation, ecology, and evolution knot when studying the origin of complex 456 animal communities.

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