The importance of neutral over niche processes in structuring Ediacaran early animal

communities

*Emily G .Mitchell[1], Simon Harris[2], Charlotte G. Kenchington[1,] Philip Vixseboxse[3], Lucy

Roberts[4], Catherine Clark[1], Alexandra Dennis[1], Alexander G. Liu[1], and Philip R. Wilby[2].

[1]Department of Earth Sciences, University of Cambridge, Downing Street, Cambridge CB2

3EQ, UK.

[2]British Geological Survey, Nicker Hill, Keyworth, Nottingham NG12 5GG, UK.

[3]School of Earth Sciences, University of Bristol, Wills Memorial Building, Queens Road,

Bristol, BS8 1RJ, UK.

[4]Department of Zoology, University of Cambridge, Downing Street, Cambridge, CB2 3EJ,

United Kingdom

Correspondence: ek338@cam.ac.uk; 07867783127

Running Title: Neutral ecology of Ediacaran communities

Keywords: Ediacaran, neutral theory, spatial point process analysis, paleoecology, ecology,

paleontology.

Article Type: Letter

Length: 145 words in abstract, 4126 words in main text, 78 references, 3 figures.

Statement of Authorship: EGM conceived the project, designed the research, ran the

analyses and wrote the first draft of the paper. All authors contributed to field data collection.

SH developed the data post-processing protocol. SH and EGM processed field data. All

authors were involved in writing the final manuscript.

Data accessibility statement: Should the manuscript be accepted, the data supporting the

results will be archived in Dryad, Figshare or Hal and the data DOI will be included at the

end of the article.

Abstract

The relative influence of niche versus neutral processes in ecosystem dynamics is a

fundamental question in community ecology, but the extent to which they structured early

animal communities is unknown. The oldest known metazoan-dominated paleocommunities

occur in Ediacaran age (~565 million years old) strata in Newfoundland, Canada and

Charnwood Forest, UK. These comprise large and diverse in-situ populations of sessile

organisms that are amenable to spatial point process analyses, enabling inference of the most

likely underlying niche or neutral processes governing their community structure. We

conducted comprehensive spatial mapping of seven of the largest Ediacaran

paleocommunities using LiDAR, photogrammetry and a laser-line probe. We find neutral

processes to dominate these paleocommunities with limited influence of niche processes.

Our results differ from the niche-dominated dynamics of modern marine ecosystems,

revealing that the dynamics of environmental interactions prompted very different ecosystem

structuring for these early animal communities.

Introduction

Two opposing theories lie at the heart of debate regarding the fundamental mechanisms that govern ecosystem structure and biodiversity: niche and neutral. Niche theory is a central tenet of classical ecological theory, whereby species avoid competitive exclusion by occupying different niches within the ecosystem (1). Smaller niche overlaps result in less competition between taxa, permitting numerous taxa to exist in an area without excluding each other. Species are able to co-exist because they have different requirements. Niche models describe selection-dominated ecosystems, whereby species dynamics operate deterministically as a series of inter-specific interactions, which act as stabilizing mechanisms for the ecosystem (2).

Neutral processes are often referred to as the 'null model' of niche processes: instead of species *differences* enabling co-existence, it is their *similarities* that drive high diversity (3). Within neutral models, species fitness is similar across a community, and so different taxa can co-exist because no single taxon has a significant competitive advantage over any other. Despite this seemingly unrealistic assumption, neutral theories have been able to accurately reproduce certain species-area-distributions (3) and beta diversity patterns (4, 5), sometimes better than niche theories (1).

In recent years, unified or continuous theories have emerged, whereby niche and neutral processes combine to enable species coexistence (2, 6). In these combined models, species can exhibit strong differences and strong stabilizations (niche-type), or similar fitness and weak stabilizations (neutral-type), with the classic niche and neutral models forming the extreme end-members of this continuum model. However, it is often not possible to select

the best-fit niche or neutral model, making it difficult to disentangle the relative influence of

niche- and neutral-type processes within modern, complex ecosystems (6).

In order to investigate how niche and neutral processes contributed to community dynamics

in deep time, we focus on some of the oldest macroscopic metazoan-dominated

paleocommunities that are currently known: those comprising the Avalonian Assemblage of

the Ediacaran macrobiota (7 - 9). The evolution of macroscopic metazoans was coupled with

a transformation in ecosystem dynamics, with paleocommunities evolving from pre-

Ediacaran microbial populations with assumed simple community structure (10), via late

Ediacaran (571–540 Ma) paleocommunities that exhibited both simple and complex

community structures (11), into Cambrian 'modern' metazoan ecosystems with comparable

ecosystem structures to the present day (12). Some of the oldest metazoan-dominated

communities form part of the Avalonian Assemblage of the Ediacaran macrobiota (13), and

are known primarily from Newfoundland, Canada and Leicestershire, UK (Fig. S1).

Avalonian soft-bodied organisms were sessile and preserved *in-situ* in deep-water

paleoenvironments dated to ~571–560 Ma (14, 15), beneath volcanogenic/volcaniclastic

event beds (16, 17). As such, bedding-plane surfaces exposed by modern weathering of tuffs

preserve near-complete census paleocommunities (16, 18), although the impact of erosion of

these surfaces needs to be considered (19; cf. 20). Since they were soft bodied, dead

organisms could not accumulate over long time periods, reducing the extent of time-

averaging. Avalonian ecosystems pre-date macro-predation and vertical burrowing, and so

remained in place post-mortem (21–23), with no evidence of locomotion on any of these

bedding-planes. Consequently, the size and position of each specimen can be considered an

accurate record of the organism's life history, including its dispersal and the habitat and

community interactions it was subject to. In common with living communities, it is therefore

possible to use spatial point process analyses (SPPA) to infer the most likely underlying

ecological and biological processes in operation (25).

For sessile organisms, community-scale spatial distributions depend on the interplay of a

limited number of different factors, namely physical environment (which manifests as habitat

associations of a taxon or taxon-pairs; 26), organism dispersal/reproduction (27), competition

for resources (28), facilitation between taxa (29), and differential mortality (30).

To assess the relative influence of niche and neutral processes for sessile communities, niche

processes are identified as intra- or inter-specific habitat associations, and/or intra- and inter-

specific competition and/or facilitation (31). Neutral processes are identified where univariate

distributions exhibit complete spatial randomness (CSR), and by dispersal processes that are

independent of local environment (i.e. habitat heterogeneities; 31-35). Dispersal patterns are

indicated by best-fit Thomas Cluster (TC) or Double Thomas Cluster (DTC) models (31).

CSR indicates neutral processes because there are no biologically or ecologically significant

intrinsic or extrinsic influences on the spatial distribution. TC and DTC aggregations are also

considered neutral since they describe dispersal processes, whereby aggregations arise from

propagules only traveling a limited distance, thus being unable to reach all suitable substrates

regardless of underlying habitat heterogeneities or species requirements (27, 36, 37).

Intra-specific habitat associations are best-modelled by a heterogeneous Poisson model (HP),

or when combined with dispersal limitations, an Inhomogeneous Thomas Cluster model

(ITC; 31, 37). Density-dependent competition, as indicated by size-dependent spatial

segregation (38), indicates a lack of sufficient resources, and is therefore a niche-based

process. The other bivariate or inter-specific interactions between taxa include facilitation,

which is considered niche because the requirement of one taxon relying on another indicates

that the facilitated taxon could not survive independently. Facilitation is best modelled by a

Linked Cluster model (LCM). Habitat associations between taxa are also considered niche

processes because such taxa associations correspond to the underlying habitat variations on

which the species depend, and are best modelled by a shared parents models (SPM) and/or

heterogeneous Poisson models (22). Therefore, for univariate distributions, neutral processes

are indicated by CSR, TC or DTC models, and niche processes by segregation and HP and

ITC models (Fig. S3). For bivariate distributions, neutral processes are indicated by CSR

while niche processes are indicated by segregation, LCM, SPM and/or HP models.

Methods

<u>Data collection and extraction</u>

In this study we assessed the univariate and bivariate spatial distributions of taxa from seven

Avalonian bedding-plane assemblages: the 'D', 'E', and Bristy Cove (X-Ray) surfaces in the

Mistaken Point Ecological Reserve, the St. Shott's surface at Sword Point; the H14 (Johnson)

surface at Little Catalina and Spaniard's Bay all in Newfoundland, Canada; and Bed B in

Charnwood Forest, UK (Fig. S1, Table S1). These spatial analyses require the mapping of

large spatial areas (up to 115m^2), in sufficient resolution to be able to taxonomically identify

the specimens from the resulting digital dataset. The best way to map the surfaces differed

depending on the preservation and dip of the surface. All surfaces were LiDAR scanned

using a Faro Focus 330X to ensure spatial accuracy was maintained over large areas. The

LiDAR scans resulted in a 3D surface mesh of 1 mm resolution. The Spaniard's Bay and

Mistaken Point 'D' and 'E' surfaces were laser scanned using a Faro Scan Arm LLP,

resulting in surface meshes of 0.050 mm resolution. The high-resolution scanning was done

in grids of 1m x 1m. Due to large file sizes, these high-resolution scans could not all be

viewed simultaneously, so control points were marked in both each high-resolution scan, and

in the LiDAR scan, enabling accurate combination of the high-resolution scans with the

LiDAR surface data (done using Geomagic 2015). Taxon identification, position, and fossil

dimensions of disc width, disc length, stem length, stem width, frond length and frond width

were marked up in Inkscape 0.92.3 on a 2D map of the combined dataset as vectors for every

specimen, creating a 2D vector map of the paleocommunity.

For H14, Bristy Cove and St Shott's surfaces, fossil relief was not sufficiently high to permit

accurate capture of all morphological details using the laser-line probe. These surfaces did

have good colour differentiation of the fossils, so a photomap was created by photographing

the specimens along a horizontal and vertical grid, and using Agisoft Photoscan software

v1.3.5 to create a photogrammetric render of the surface. The LiDAR scan was then

imported into Photoscan, and the photographs aligned on the LiDAR scan to ensure large-

scale accuracy. An orthomosaic of the surface was produced within Agisoft PhotoScan, and

the fossils marked up as vectors as above.

Bed B, Charnwood Forest has a dip of 45°, and so is not suitable for *in situ* high-resolution

scanning using our equipment. Instead, we used Reflectance Transformation Images (RTIs)

of casts of this surface (57, 58). Each RTI was marked up as a vector map and imported onto

the LiDAR scan. The LiDAR scan enabled checking of mould deformation, and where

needed was used to retrodeform the vector map.

Upon completion, vector maps were processed using a custom script in Haskell (59), which

output the specimen identification number, taxonomic identification, and specimen

dimensions. This output formed the basic dataset for the spatial analyses.

Taxonomic identification

Specimens were assigned to one of twenty macrofossil taxa/groups, including several 'bin'

groups (60), on the basis of their morphological attributes:

1) *Arborea*, 2) *Aspidella*, 3) *Avalofractus*, 4) *Beothukis*, 5) *Bradgatia*, 6) Brushes, 7) *Charnia*,

8) *Charniodiscus*, 9) "Feather Dusters" which includes *Plumeropriscum* and

Primocandlebrum, 10) *Fractofusus andersoni + F. misrai*, 11) *Hylaecullulus*, 12)

Ivesheadiomorphs, 13) Ostrich Feather, 14) *Pectinifrons*, 15) *Primocandelabrum*, 16)

Thectardis, 17) *Trepassia*, 18) *Vinlandia*, 19) "Holdfast Discs" [all discoidal specimens of

uncertain affinity, with or without associated stems, which lack sufficient detail to identify

the taxon], 20) "Other Species" [rare forms that do not fall into any of the other groups; e.g.,

Hapsidophyllas]. Non-abundant taxa (taken as < 30 specimens) and taphomorphs (organ taxa,

such as *Hiemalora* or the decayed remains of already dead organisms, such as

ivesheadiomorphs) were excluded from analyses, leaving 13 abundant taxa, three of which

(*Charniodiscus*, *Charnia*, *Bradgatia*) occur abundantly on two bedding-planes and one

(*Fractofusus*) on four bedding-planes.

Bias analyses

Differential erosion has the potential to distort spatial analyses (17, 19) so for each surface,

we tested for erosional biases (19) and tectonic deformation, corrected for these factors into

account if they significantly affected specimen density distributions (Fig. S2, Table S2). .

Our data have been tested for the influence of differential erosion using heterogeneous

Poisson models. We modelled possible sources of erosion (cf. 20), fitting at least three

heterogeneous Poisson models to the data, with the models dependent on x (parallel to strike),

y (parallel to dip), and a point chosen on a surface-by-surface basis dependent on the most

likely point of erosion. The St. Shott's, Bed B and H14 surfaces all showed significant fossil

density changes depending on these physical features of the surface, implying that the

observed surface specimen density has been significantly influenced by post-preservational

erosion processes. Such heterogeneous erosional processes were incorporated into

subsequent analyses so that the underlying biological and ecological processes could be

investigated. Tectonically distorted data were retrodeformed by returning elongated holdfast

discs to a circular outline (16, 20).

<u>Spatial Analyses</u>

Initial data exploration, inhomogeneous Poisson modelling and segregation tests were

performed in R (61) using the package spatstat (62-64). Programita was used to find distance

measures and to perform aggregation model fitting (described in detail in references (65-68)).

The univariate spatial distribution of each taxon on each bedding plane was described using

pair correlation functions (PCFs). A PCF = 1 indicates a distribution that was completely

spatially random (CSR); PCF > 1 indicates aggregation; and PCF < 1 indicates segregation

(25, 32, 39). Univariate and Bivariate pair correlation functions (PCFs) were calculated from

the population density using a grid of 10cm x 10cm cells on all surfaces except Bristy Cove,

where a 1cm x 1cm cell size was used. To minimise noise, a smoothing was applied to the

PCF dependent on specimen abundance: This smoothing was over three cells with all

surfaces except Bristy Cove which had a 5 cell smoothing. To test whether the PCF

exhibited complete spatial randomness (CSR), 999 simulations were run for each univariate

and bivariate distribution, with the 49th highest and lowest values removed (69). CSR was

modelled by a Poisson model on a homogeneous background where the PCF = 1 and the fit

of the fossil data to CSR was assessed using Diggle's goodness-of-fit test (32, 39). Note that

due to non-independence of spatial data, Monte-Carlo generated simulation envelopes cannot

be interpreted as confidence intervals. If the observed data fell below the Monte-Carlo

simulations, the bivariate distribution was interpreted to be segregated; above the Monte-

Carlo simulations, the bivariate distribution was found to be aggregated.

If a taxon was not randomly distributed on a homogeneous background, and was aggregated

(Fig. S3, Table S3), the random model on a heterogeneous background was tested by creating

a heterogeneous background created from the density map of the taxon under consideration,

being defined by a circle of radius R over which the density is averaged throughout the

sample area. Density maps were formed using estimators within the range of $0.1m < R < 1m$,

with R corresponding to the best-fit model used. If excursions outside the simulation

envelopes for both homogeneous and heterogeneous Poisson models remained, then Thomas

cluster models were fitted to the data as follows:

1. The PCF and L function (70) of the observed data were found. Both measures were

calculated to ensure that the best-fit model is not optimized towards only one distance

measure, and thus encapsulates all spatial characteristics.

2. Best-fit Thomas cluster processes (71) were fitted to the two functions where PCF>1. The

best-fit lines were not fitted to fluctuations around the random line of PCF=1 in order to aid

good fit about the actual aggregations, and to limit fitting of the model about random

fluctuations. Programita used the minimal contrast method (32, 39) to find the best-fit model.

3. If the model did not describe the observed data well, the lines were re-fitted using just the

PCF. If that fit was also poor, then only the L-function was used.

4. 99 simulations of this model were generated to create simulation envelopes, and the fit

checked using the O-ring statistic. (64)

5. p_d was calculated over the model range. Very small-scale segregations (under 2 cm) were

not included in the model fitting, since they likely represent the finite size of the specimens,

and a lack of specimen overlap.

6. If there were no excursions outside the simulation envelope and the p_d-value was high,

then a univariate homogeneous Thomas cluster model was interpreted as the best model.

For each bivariate distribution displaying segregation, the size-classes of each taxon were

calculated, the bivariate PCFs of the smallest size-classes and largest size-classes were

plotted, with 999 Monte Carlo simulations of a complete spatially random distribution and

segregation tests performed. The most objective way to resolve the number and range of size

classes in a population is by fitting height-frequency distribution data to various models,

followed by comparison of (logarithmically scaled) Bayesian information criterion (BIC)

values (72), which we performed in R using the package MCLUST (73). The number of

populations thus identified was then used to define the most appropriate size classes.   A BIC

value difference of >10 corresponds to a "decisive" rejection of the hypothesis that two

models are the same, whereas values <6 indicate only weakly rejected similarity of the

models (72-77). Once defined, the PCFs for each size class were calculated. Although it was

necessary to set firm boundaries for each size class, the populations are normally distributed

and therefore overlap. As a result, the largest individuals of the small population are grouped

within the middle size class, while some of the smallest of the medium population are

included within the small size class. As such, the medium population was excluded from

analyses.

Results

Across the seven surfaces and the 19 taxon univariate distributions examined, eight taxon

distributions were best modelled by CSR (Figs. 1 and 2, Table S3). Of the non-CSR taxon

distributions, 10 were best modelled by TC (or DTC). Only *Trepassia* on Spaniard's Bay

was best modelled by an ITC model (Fig. 2, Table S3). The only taxa which had univariate

spatial distribution with a HP best-fit model was *Beothukis* on the 'E' and Spaniard's Bay

surfaces (Figs. 1 and 2).

In order to gain some indication of whether taxa behave differently over large-spatial scales

we compared the univariate spatial distributions of individual taxa at different sites to

represent paleocommunities separated by large spatial and temporal scales. Four taxa are

abundant on multiple bedding-planes (*Bradgatia*, *Charnia*, *Charniodiscus* and *Fractofusus*),

and they all exhibit a consistent best-fit model (CSR, TC, TC and TC/DTC respectively) on

each surface they are found on. Previous work has demonstrated that *Fractofusus* shows

consistently the same spatial distributions across multiple surfaces (18) in different geological

units (cf. H14 and 'E' surface, Fig. 1H). *Charniodiscus* also shows consistent spatial

distributions (Fig. 1I), even when the communities were in temporally separated localities in

different locations on Avalonia (e.g. *Charniodiscus* from the 'E' surface and Bed B). The

consistency of these results suggests that the small-spatial-scale ecological behaviour of these

taxa did not change over large spatial and temporal scales.

Two surfaces out of the five studied paleocommunities with more than one abundant taxon

present exhibited only CSR bivariate distributions (St. Shott's and the 'D' surface (22) Fig. 3,

Table S3). The 'E' surface (22,44), Spaniard's Bay and Bed B have exhibit non-CSR

bivariate distributions (Fig. 3) indicating shared habitat associations. On Bed B the non-CSR

bivariate distribution indicates shared habitat associations between the large *Charnia* and

Primocandelabrum specimens (Fig. 3B, Table S3), as do the three non-CSR bivariate

distributions on the 'E' surface (22, 44). For the *Primocandelabrum* 'E' surface and the Bed

B non-CSR habitat associations, the large specimens had a segregated spatial distribution

which corresponds to a reduced specimen density compared to CSR. This reduction in

specimen density indicates that there were not enough resources to sustain all of the

population and so is a niche process (44).   On the 'E' surface, inter-specific segregations

reduced specimen density by 25%, and aggregations increased specimen density by 56%

(Fig. 3C, D). In contrast, intra-specific dispersal processes had a large effect on specimen

density increasing their density between 250–600% (Fig. 1F). The two habitat associations on

the 'E' surface (Feather Dusters–*Fractofusus* and Feather Dusters–*Charniodiscus*) were

reflected in small bivariate aggregations (increase of 34% over distances under 0.2 m and

56% under 1.2 m respectively) with a reduction in their joint density at large spatial-scales

(11% over 1 m and 13% over 2.1 m respectively; Fig. 3C, D). Similarly, habitat association

between *Charnia* and *Primocandelabrum* on Bed B increased specimen density by 87%,

whereas segregations reduced taxon density by 10%. Univariate dispersal-generated

aggregations increased taxon density by 180–500% (Figs. 1A). The *Trepassia – Beothukis*

bivariate distribution on Spaniard's Bay is best modelled by the *Trepassia* best-fit model ITC

model (Table S3) which is a Thomas Cluster model fitted onto the heterogeneous Poisson

model background of *Beothukis* (Fig. 3A). This result demonstrates that there is a single

habitat heterogeneity impacting both taxa on scales above 40 cm, but influencing *Beothukis*

more strongly than *Trepassia* (Fig. 3G, although see *SI Appendix*). Across the

paleocommunities, the bivariate habitat associations are much weaker in PCF magnitude than

the univariate distributions (Figs 1 and 3), showing that the bivariate (niche) processes had

less impact on spatial distributions than univariate (neutral) processes. These spatial

distributions suggest that competition is rare, and that where it was present it was relatively

weak in magnitude (Fig. 3; cf. refs. 22, 44).

Discussion

Our results support combined theories of community assembly whereby niche and neutral

theories are not mutually exclusive, but instead act along a continuum or spectrum, with

differing extents of niche and neutral processes present in different circumstances (45, 46).

The dominance of neutral best-fit univariate models, repetition of best-fit univariate models

across different paleocommunities, and the rarity and weakness of bivariate niche best-fit

models, all combine to provide strong evidence that neutral processes dominated Avalonian

paleocommunities, with only limited niche-based influence. These neutral-dominated

community dynamics contrast with those observed in the modern marine realm, where

neutral processes are rare (47, 48).

The difference in dominance of niche versus neutral processes raises the question of whether

the nature of community dynamics of Ediacaran metazoan-dominated paleocommunities was

fundamentally different to those of the present day. The only other work on niche-neutral

influences on paleocommunity assembly, is from the Quaternary (2.58 – 0.01 Ma), where

fossil assemblages provides strong model and empirical support for environment-led (niche)

models of assembly (49). There are some notable differences between Avalonian

paleocommunities and extant marine communities. Avalonian paleocommunities appear to

differ from the majority of extant marine systems in the extent of their ecological maturity, in

that no more than three generations are seemingly preserved (22), though some

paleocommunities include rare survivors (23) and/or evidence of secondary community

succession (50). These characteristics suggest that the fossil communities are not always

mature, many having been curtailed by high frequency incursions of sediment, limiting their

maturity (22). Recent models show that community dynamics in small populations

periodically subjected to disturbance events are dominated by neutral processes, implying a

lack of small-spatial scale environmental control on their ecological dynamics (46).

The relative influence of niche versus neutral processes has been shown to be effected by the

dispersal ranges of taxa within communities (52). Wide dispersal ranges increase the

connectivity between populations, and so expand effective community size with the net effect

of enhancing ecological selection (competition) and thereby increasing the relative

importance of niche processes (52). The opposite is true when dispersal is limited, making

these communities neutral dominated (52). Within the Ediacaran, the global distributions of

some Avalonian taxa provide evidence that these taxa were capable of wide dispersal (51,

53). Further evidence of dispersal ranges is found in their spatial distributions (20, 44). For

example, we can see in the PCF plots that *Fractofusus, Charniodiscus* and

Primocandelabrum (Fig. 1A, H and I) have very short dispersal ranges, of <10 cm for

Fractofusus and *Primocandelabrum* and <20 cm for *Charniodiscus*. However, *Fractofusus*

was also capable of a waterborne propagule stage (20), and the global distribution of

Charniodiscus suggests that it was as well, but that the waterborne phrase resulted in a

minority of the population (20).   Wide dispersal ranges are suggested by the global

distribution of taxa such as *Charnia* (53) and also by the CSR and HP distributions of

Beothukis and *Bradgatia* (Table S3). Six of the seven studied paleocommunities were

dominated by taxa such as *Fractofusus* and *Charniodiscus* which predominantly exhibit

limited local dispersal (Fig 1H, I; 18, 22, 44). The studied Ediacaran paleocommunities have

comparatively small populations, experienced frequent disturbance events, and include many

taxa with short dispersal ranges, so within this framework we would expect neutral processes

to dominate. While the dominance of neutral-based processes within these paleocommunities

differs significantly to the majority of the modern marine realm, the underlying dynamics are

entirely consistent with models of assembly that include both niche and neutral processes,

and are similar to those of modern communities subject to the same conditions. Thus, it is

therefore likely that the fundamental mechanisms of metazoan community assembly were

already in place in the Ediacaran Period, and so may have existed unchanged for ~570

million years.

In a similar manner to ecological processes, evolutionary processes can be categorised as

niche (selection) or neutral (drift) processes (54). Selection (niche) processes are considered

deterministic because external factors, such as limited resources, lead to competition in a

predictable way: given a set of initial conditions, the organisms/communities will always

respond to these conditions (environment) in the same way (54). By contrast, drift (neutral)

processes are considered stochastic because they result from random fluctuations in

population demography, so, given a set of the same initial conditions, different

populations/communities may emerge. Hence, the observed dominance of neutral ecological

processes in the Ediacaran Avalonian paleocommunities establishes that they are inherently

stochastic/probabilistic with the possible implication that early metazoan diversification was

not a systematic adaption to optimise survival under prevailing environmental conditions

(which would be niche processes, and so deterministic). Instead, their existence under a

stochastic regime would mean that diversification could have been merely driven by

demographic differences resulting from random within-population. If this hypothesis is

correct, and early metazoan evolution was stochastic, then this stochasticity may help to

explain why neutral models of evolution can reproduce substantial macro-evolutionary trends

such as the Cambrian Explosion (cf. 55), despite the known importance of niche processes in

shaping evolution (e.g. 56).

Conclusions

We have shown that paleocommunities of early macroscopic metazoans were overwhelming dominated by neutral ecological processes, with only limited and weak evidence for niche processes. Our results strongly contrast with modern marine systems, but because our Ediacaran paleocommunities have traits (short dispersal ranges, small populations and frequent disturbances) that are associated with neutral ecological models our results suggest that the fundamental mechanisms of community assembly may have been in place since the early stages of metazoan evolution. The dominance of neutral processes in these paleocommunities suggests that systematic adaptation of the Ediacaran organisms to their local environment may not have been the underlying driver of early metazoan diversification. Instead, it is possible that late Neoproterozoic metazoan diversification may result from stochastic demographic differences, with only limited environmental influence.

Acknowledgments: The Parks and Natural Areas Division (PNAD), Department of Environment and Conservation, Government of Newfoundland and Labrador provided permits to conduct research within the Mistaken Point Ecological Reserve (MPER) in 2010, 2016 and 2017. Readers are advised that access to MPER is by scientific research permit only. Contact PNAD for further information. Access to Bed B was kindly facilitated by Natural England and landowners in Charnwood Forest. This work has been supported by the Natural Environment Research Council [grant numbers NE/P002412/1 to EGM; NE/P002412/1 to CGK and PRW; and Independent Research Fellowship NE/L011409/2 to AGL], a Gibbs Travelling Fellowship from Newnham College, Cambridge and a Henslow Research Fellowship from Cambridge Philosophical Society to EGM. CGK also acknowledges a Research Studentship funded by the Cambridge Philosophical Society.

The authors declare no competing interests.

References

1. MacArthur, R. H. (1984). *Geographical ecology: patterns in the distribution of species.*

Princeton University Press.

2. Adler, P. B., Hille R, Lambers, J., & Levine, J. M. (2007). A niche for neutrality. *Ecology*

Letters, 10: 95-104.

3. Hubbell, S. P. (2001). The *unified neutral theory of biodiversity and biogeography* (MPB-

32) (monographs in population biology).

4. Condit, R., Pitman, N., Leigh, E. G., Chave, J., Terborgh, J., Foster, R. B., Núnez, P.,

Aguilar, S., Valencia, R., Villa, G. and Muller-Landau, H.C (2002). Beta-diversity in tropical

forest trees. *Science*, 295: 666-669.

5. Chisholm, R. A., & Pacala, S. W. (2010). Niche and neutral models predict asymptotically

equivalent species abundance distributions in high-diversity ecological communities.

Proceedings of the National Academy of Sciences, 107: 5821-15825..

6. Gravel, D., Canham, C. D., Beaudet, M., & Messier, C. (2006). Reconciling niche and

neutrality: the continuum hypothesis. *Ecology Letters*, 9: 399-409.

7. Droser, M. L., Tarhan, L. G., & Gehling, J. G. (2017). The rise of animals in a changing

environment: global ecological innovation in the late Ediacaran. *Annual Review of Earth and*

Planetary Sciences, 45: 593-617.

8. Budd, G. E., & Jensen, S. (2017). The origin of the animals and a 'Savannah' hypothesis

for early bilaterian evolution. *Biological Reviews*, 92: 446-473.

9. Wood, R., Liu, A. G., Bowyer, F., Wilby, P., Dunn, F. S., Kenchington, C. G., Hoyal

Cuthill, J., Mitchell, E. G., & Penny, A. (2019). Integrated records of environmental change

and evolution challenge the Cambrian Explosion. In press *Nature Ecology and Evolution*

10. Butterfield, N. J. (2007). Macroevolution and macroecology through deep time.

Paleontology, 50: 41-55.

11. Darroch, S. A., Laflamme, M., & Wagner, P. J. (2018). High ecological complexity in

benthic Ediacaran communities. *Nature Ecology Evolution*, 2: 1541.

12. Dunne, J. A., Williams, R. J., Martinez, N. D., Wood, R. A., & Erwin, D. H. (2008).

Compilation and network analyses of Cambrian food webs. *PLoS Biology*, 6: e102.

13. Waggoner, B. (2003). The Ediacaran biotas in space and time. *Integrative and*

Comparative Biology 43: 104–113.

14. Pu, J. P., Bowring, S. A., Ramezani, J., Myrow, P., Raub, T. D., Landing, E., Mills, A.,

Hodgin, E. and Macdonald, F. A., (2016). Dodging snowballs: Geochronology of the

Gaskiers glaciation and the first appearance of the Ediacaran biota. *Geology*, 44: 955-958.

15. Noble, S. R., Condon, D. J., Carney, J. N., Wilby, P. R., Pharaoh, T. C., & Ford, T. D.

(2015). U-Pb geochronology and global context of the Charnian Supergroup, UK: constraints

on the age of key Ediacaran fossil assemblages. *GSA Bulletin*, 127: 250-265.

16. Wood, D. A., Dalrymple, R. W., Narbonne, G. M., Gehling, J. G. & Clapham, M. E.

(2003). Paleoenvironmental analysis of the late Neoproterozoic Mistaken Point and

Trepassey formations, southeastern Newfoundland. *Can. J. Earth Sci.* 40: 1375–1391.

17. Narbonne, G. M. (2005). The Ediacara biota: Neoproterozoic origin of animals and their

ecosystems. *Annu. Rev. Earth Planet. Sci.*, 33: 421-442.

18. Clapham, M. E., Narbonne, G.M. & Gehling, J. G. (2003) Paleoecology of the oldest

known animal communities: Ediacaran assemblages at Mistaken Point, Newfoundland.

Paleobiology 29: 527–544.

19. Matthews, J.J., Liu, A.G. and McIlroy, D., 2017. Post-fossilization processes and their

implications for understanding Ediacaran macrofossil assemblages. *Geological Society,*

London, Special Publications, 448: 251-269.

20. Mitchell, E. G., Kenchington, C. G., Liu, A. G., Matthews, J. J., & Butterfield, N. J.

(2015). Reconstructing the reproductive mode of an Ediacaran macro-organism. *Nature*, 524:

343-346.

21. Liu, A. G., McIlroy, D., Antcliffe, J. B., & Brasier, M. D. (2011). Effaced preservation in

the Ediacara biota and its implications for the early macrofossil record. *Paleontology* 54:

607–630.

22. Mitchell, E. G., & Butterfield, N. J. (2018). Spatial analyses of Ediacaran communities at

Mistaken Point. *Paleobiology*, 44: 40-57.

23. Wilby, P. R., Kenchington, C. G., & Wilby, R. L. (2015). Role of low intensity

environmental disturbance in structuring the earliest (Ediacaran) macrobenthic tiered

communities. *Paleogeography, Paleoclimatology, Paleoecology,* 434: 14-27.

24. Liu, A. G. (2016). Framboidal pyrite shroud confirms the 'death mask' model for moldic

preservation of Ediacaran soft-bodied organisms. *Palaios*, 31: 259-274.

25. Illian, J., Penttinen, A., Stoyan, H., & Stoyan, D. (2008). *Statistical Analysis and*

Modelling of Spatial Point Patterns Vol. 70 (John Wiley).

26. Wiegand, T., Gunatilleke, S., & Gunatilleke, N. (2007). Species associations in a

heterogeneous Sri Lankan dipterocarp forest. *The American Naturalist*, 170: E77–E95.

27. Seidler, T. G., & Plotkin, J. B. (2006). Seed dispersal and spatial pattern in tropical trees.

PLoS Biology, 4: e344.

28. Getzin, S., Dean, C., He, F., Trofymow, J. A., Wiegand, K., & Wiegand, T. (2006).

Spatial patterns and competition of tree species in a Douglas fir chronosequence on

Vancouver Island. *Ecography,* 29: 671–682.

29. Lingua, E., Cherubini, P., Motta, R., & Nola, P. (2008). Spatial structure along an

altitudinal gradient in the Italian central Alps suggests competition and facilitation among

coniferous species. *Journal of Vegetation Science*, 19: 425-436.

30. Getzin, S., Wiegand T. , Wiegand, K., & He, F. (2008). Heterogeneity influences spatial

patterns and demographics in forest stands. *Journal of Ecology* 96: 807–820.

31. Lin, Y. C., Chang, L. W., Yang, K. C., Wang, H. H., & Sun, I. F. (2011). Point patterns of

tree distribution determined by habitat heterogeneity and dispersal limitation. *Oecologia*, 165:

175-184.

32. Diggle, P., Zheng, P., & Durr, P. (2005). Nonparametric estimation of spatial segregation

in a multivariate point process: Bovine tuberculosis in Cornwall, UK. *Journal of the Royal*

Statistical Society: Series C (Applied Statistics), 54: 645-658.

33. Law, R., Illian, J., Burslem, D. F., Gratzer, G., Gunatilleke, C. V. S., & Gunatilleke, I. A.

U. N.. (2009). Ecological information from spatial patterns of plants: insights from point

process theory. *Journal of Ecology,* 97: 616–628.

34. Comita, L. S., Condit, R., & Hubbell, S. P. (2007). Developmental changes in habitat

associations of tropical trees. *Journal of Ecology*, 95: 482-492.

35. Wiegand, T., Gunatilleke, S., Gunatilleke, N., & Okuda, T. (2007). Analyzing the spatial

structure of a Sri Lankan tree species with multiple scales of clustering. *Ecology,* 88: 3088–

3102.

36. Hubbell, S. P., Foster, R. B., O'Brien, S. T., Harms, K. E., Condit, R., Wechsler, B., &

De Lao, S. L. (1999). Light-gap disturbances, recruitment limitation, and tree diversity in a

neotropical forest. *Science*, 283: 554-557.

37. Harms, K. E., Wright, S. J., Calderón, O., Hernandez, A., & Herre, E. A. (2000).

Pervasive density-dependent recruitment enhances seedling diversity in a tropical

forest. *Nature*, 404: 493-495.

38. Kenkel, N. C. (1988). Pattern of self thinning in Jack pine: testing the random mortality

hypothesis. *Ecology*, 69: 1017-1024.

39. Diggle, P. (2003). *Statistical Analysis of Spatial Point Patterns*. 2nd edn (Arnold).

40. Levin, S. A. (1995). In *Ecological Time Series* Vol.2 (eds Powell, T. M. & Steele, J. H.)

pp 277–326 (Springer).

41. Murrell, D. J., & Law, R., (2003). Heteromyopia and the spatial coexistence of similar

competitors. *Ecology Letters,* 6: 48-59.

42. McIntire, E. J., & Fajardo, A. (2009). Beyond description: the active and effective way to

infer processes from spatial patterns. *Ecology,* 90: 46–56.

43. Wiegand, T., & Moloney, K. A. (2013). *Handbook of Spatial Point-Pattern Analysis in*

Ecology (CRC press).

44. Mitchell, E. G., & Kenchington, C. G. (2018). The utility of height for the Ediacaran

organisms of Mistaken Point. *Nature Ecology and Evolution*, 2: 1218-1222.

45. Thomas, M. (1949) A generalization of Poisson's binomial limit for use in ecology.

Biometrika 36, 18–25.

46. Fisher, C. K., & Mehta, P. (2014). The transition between the niche and neutral regimes

in ecology. *Proceedings of the National Academy of Sciences*, 111: 13111-13116.

47. Dornelas, M., Connolly, S. R., & Hughes, T. P. (2006). Coral reef diversity refutes the

neutral theory of biodiversity. *Nature*, 440: 80-82.

48. Connolly, S. R., MacNeil, M. A., Caley, M. J., Knowlton, N., Cripps, E., Hisano, M., &

Brandt, A. (2014). Commonness and rarity in the marine biosphere. *Proceedings of the*

National Academy of Sciences, 111: 8524-8529.

49. Jackson, S. T., & Blois, J. L. (2015). Community ecology in a changing environment:

Perspectives from the Quaternary. *Proceedings of the National Academy of Sciences,* 112:

4915-4921.

50. Liu, A. G., McIlroy, D., Matthews, J. J., & Brasier, M. D. (2012). A new assemblage of

juvenile Ediacaran fronds from the Drook Formation, Newfoundland. *Journal of the*

Geological Society, 169: 395–403.

51. Boag, T. H., Darroch, S. A., & Laflamme, M. (2016). Ediacaran distributions in space

and time: testing assemblage concepts of earliest macroscopic body fossils. *Paleobiology*, 42:

574-594.

52. Ron, R., Fragman-Sapir, O., & Kadmon, R. (2018). Dispersal increases ecological

selection by increasing effective community size. *Proceedings of the National Academy of*

Sciences, 115: 11280-11285.

53. Darroch, S. A., Laflamme, M., & Clapham, M. E. (2013). Population structure of the

oldest known macroscopic communities from Mistaken Point,

Newfoundland. *Paleobiology*, 39: 591-608.

54. Chave, J. (2004). Neutral theory and community ecology. *Ecology Letters*, 7: 241-253.

55. Budd, G. E., & Mann, R. P. (2018). History is written by the victors: the effect of the

push of the past on the fossil record. *Evolution*, 72: 2276-2291.

56. Hutchinson, G. E. (1957). *A Treatise on Limnology*, Wiley and Sons.

57. Kenchington, C. G., Harris, S. J., Vixseboxse, P. B., Pickup, C., & Wilby, P. R. (2018).

The Ediacaran fossils of Charnwood Forest: shining new light on a major biological

revolution. *Proceedings of the Geologists' Association,* 129: 264-277.

58. Mitchell, E. G., Kenchington, C. G., Harris, S., & Wilby, P. R. (2018). Revealing

rangeomorph species characters using spatial analyses. *Canadian Journal of Earth*

Sciences, *55*(11), 1262-1270.

59. Jones, S. P. (Ed.). (2003). *Haskell 98 language and libraries: the revised report.*

Cambridge University Press.

60. Shen, B., Dong, L., Xiao, S. & Kowalewski, M. (2008). The Avalon explosion: evolution

of Ediacara morphospace. *Science,* 319: 81–84.

61. R Core Team. (2013) *R: A Language and Environment for Statistical Computing.* R

Foundation for Statistical Computing Vienna, Austria

62. Baddeley, A., & Turner, R. (2005). Spatstat: an R package for analyzing spatial point

patterns. *J. of Statistical Software,* 12: 1–42

63. Berman, M. (1986) Testing for spatial association between a point process and another

stochastic process. *Applied Statistics,* 35: 54–62

64. Wiegand, T., & Moloney, K. (2004). Rings, circles, and null-models for point pattern

analysis in ecology. *Oikos,* 104: 209–229

65. Wiegand, T., Kissling, W., Cipriotti, P., & Aguiar, M. (2006). Extending point pattern

analysis for objects of finite size and irregular shape. *J. of Ecology,* 94: 825–837

66. Wiegand, T., Moloney, K., Naves, J., & Knauer, F. (1999). Finding the missing link

between landscape structure and population dynamics: a spatially explicit perspective. *The*

American Naturalist, 154: 605–627

67. Loosmore, N. B., & Ford, E. D. (2006). Statistical inference using the G or K point

pattern spatial statistics. *Ecology,* 87: 1925–1931

69. Wiegand, T., & Moloney, K. A. (2013). *Handbook of Spatial Point-pattern Analysis in*

Ecology. CRC Press

70. Levin, S. A. (1992). The problem of pattern and scale in ecology: the Robert H.

MacArthur award lecture. Ecology, 73(6), 1943-1967.

71. Besag, J. (1974). Spatial interaction and the statistical analysis of lattice systems. *J. of the*

Royal Statistical Society. Series B (Methodological), 36: 192–236

72. Thomas, M. A. (1949). A generalization of Poisson's binomial limit for use in ecology.

Biometrika, 36: 18–25.

73. Grabarnik, P., Myllymäki, M., & Stoyan, D. (2011). Correct testing of mark

independence for marked point patterns. *Ecological Modelling,* 222: 3888–3894

74. Fraley, C., & Raftery, A. E. (2006). *MCLUST version 3: an R package for normal*

mixture modeling and model-based clustering. Washington University Seattle Dept. of

Statistics.

75. Fraley, C., & Raftery, A. E. (2007). Bayesian regularization for normal mixture

estimation and model-based clustering. *Journal of Classification,* 24: 155–188

76. Pélissier, R., & Goreaud, F. (2001). A practical approach to the study of spatial structure

in simple cases of heterogeneous vegetation. *J. of Vegetation Science,* 12: 99–108

77. Stoyan, D., Kendall, W. S., & Mecke, J. (1995). *Stochastic geometry and its applications*.

2nd edition. 458 p. Springer Verlag.

78. Clapham, M. E., & Narbonne G. M., Gehling, J. G. Ediacaran epifaunal tiering.

Geology, 30, 627-630 (2002).

Figure 1. Univariate PCF for the seven Ediacaran fossil surfaces. A) Bed B, Charnwood

Forest, B) Bristy Cove/X-Ray surface, C) 'D' surface, D) H14/Johnson surface E) St. Shott's

surface, F) 'E' surface, G) Spaniard's Bay surface. The univariate PCFs of H) Fractofusus

and I) Charniodiscus *from multiple surfaces are shown to demonstrate the similarity of their*

spatial distributions between localities. Where the best-fit model for the distribution

represents a niche process (Table S3), it is drawn as a dashed line. Models indicating neutral

processes are drawn as solid lines. Black line represents the random model. The grey area is

the simulation envelope for 999 Monte Carlo simulations. The x-axis is the inter-point

distance between organisms in metres. On the y-axis PCF=1 indicates complete spatial

randomness (CSR), <1 indicates segregation, and >1 indicates aggregation. Different colors

indicate different taxa as follows: Thectardis *navy;* Fractofusus *light blue;* Charnia *bright*

yellow; Charniodiscus *dark red;* Aspidella *light green;* Bradgatia *dark green;* 'Feather

Duster', light orange; Primocandelabrum *dark orange;* Trepassia *dark purple;* Beothukis

bright pink; Pectinifrons *dark blue;* 'Brushes' *brown;* Avalofractus *dark blue;* Hylaecullulus

light yellow.

Figure 2. *Proportion of best-fit univariate models by surface. The percentage of taxa with*

univariate spatial distributions that are best described by CSR, HP, TC (or DTC) and ITC

models. CSR and TC are considered neutral models and shown in blue. HP and ITC are

niche models, shown in red.

Figure. 3. *Bivariate PCF for the taxa which had non-random spatial distributions. The grey*

area is the simulation envelope for 999 Monte Carlo simulations. The x-axis is the inter-

point distance between organisms in metres. On the y-axis, PCF=1 indicates complete

spatial randomness (CSR) and is indicated by a black line, <1 indicates segregation, and >1

indicates aggregation. Red line is the modelled distribution. A) Trepassia *and* Beothukis

from Spaniard's Bay. The red line is the heterogeneous Poisson model. B) The non-CSR

bivariate distribution of Charnia *and* Primocandelabrum *from Bed B, Charnwood Forest*

shows randomly distributed small specimens (< 5.0 cm)and segregated large specimens(>

10.0 cm). C) Feather Dusters and Fractofusus *and D) Feather Dusters and* Charniodiscus

from Mistaken Point 'E' Surface show aggregated small (< 3.0 cm) specimens and

segregated large specimens (> 5.5cm). C and D reproduced from ref. 22.

Supplementary Information

Figure S1. Locality map of study sites, showing: A, the relative location of sites with respect

to the micro-continent of Avalonia (grey shading), highlighting the Mistaken Point

Ecological Reserve (MP) and the Bonavista Peninsula (BP) in Newfoundland, and Bed B in

Charnwood Forest, UK. B, the Newfoundland sites of the 'D', 'E', and Bristy Cove (X-Ray)

surfaces, all in the Mistaken Point Ecological Reserve; the St. Shott's (Sword Point) surface;

and the H14/Johnson surface, Bonavista Peninsula (modified from ref. 22). Associated spatial

maps for each locality show the positions of the fossil specimens, where the size of the circle

indicates the vertical height. Black scale bar = 1 m, grey scale bar = 0.1 m. Different colors

indicate different taxa as follows: Thectardis navy; Fractofusus light blue; Charnia bright

yellow; Charniodiscus dark red; Aspidella light green; Bradgatia dark green ; Feather Dusters

light orange; Primocandelabrum dark orange; Trepassia dark purple; Beothukis bright pink;

Pectinifrons dark blue; 'Brushes' brown; Avalofractus dark blue; Hylaecullulus light yellow.

Surface	Area mapped (m²)	Total number of specimens	Density (ind/m²)	Number of abundant (>30) taxa	Minimum size of consistently preserved features (cm)
Bed B	115.16	761	6.61	3	1.20
MP E	82.44	2977	36.83	6	0.70
MP D	70.89	1402	22.29	3	1.20
H14	82.4	4235	52.55	1	1.50
BC	0.77	76	133.25	1	0.70
St. Shott's	50.93	480	7.20	2	1.20
*Spaniard's Bay**	16.40	68	2.61	3	0.20

Table S1: Summary data for each surface. Area is total area mapped. Total number of

specimens includes non-abundant and taphomorph taxa. Density is of abundant taxa (>30

individuals) only. Minimum size consistently preserved is the modal height of small

specimens, because specimens beneath this threshold exhibit discontinuous distributions (cf

e.g. 79) so it is likely that specimens beneath this threshold may not have been

preserved/mapped. *Note due to low specimen numbers on the Spaniard's Bay surface, we

included >15 specimens as abundant taxa (see SI appendix).

Figure S2: Model density maps of surfaces showing erosion biases.

For each surface darker colors indicate higher modelled fossil density, and therefore lower

presumed erosional rate, normalised for the density on each surface. Note that Bed B has a

coarser pattern due to a relatively lower fossil density difference across the surface and that

the full spatial map is not provided in full due to concerns about fossil theft.

Surface	X	y	$\sqrt{[(x-x_1)^2 + (y-y_1)^2]}$	x_1	y_1
Bed B	3.26	**6.94**	1.54	89	112
Bristy Cove	-1.93	-1.93	-1.77	84	91
Mistaken Point D	-3.37	-12.29	-12.51	52	939
Mistaken Point E	-75.64	-5.06	-72.29	309	283
H14	43.26	37.69	**87.34**	98	70
St. Shott's	-0.36	14.17	**23.82**	93	113
Spaniard's Bay	0.14	-1.96	-1.78	294	143

672 **Table S2: ΔAIC values for density models used to investigate erosional biases.** *x* is

673 parallel to strike, *y* parallel to dip, $\sqrt{[(x-x_1)^2 + (y-y_1)^2]}$ is the distance from the point of least

674 erosion, and x_1 and y_1 are the co-ordinates of that point. These ΔAIC were used to determine the

675 best-fit models. ΔAIC > 0 indicates that the model has a better fit to the data than completely

676 spatially random model. Units are the centimetre co-ordinates of the spatial maps.

Figure S3. Univariate PCF analyses for Fractofusus on the 'D' surface (left) and Beothukis

on the 'E' surface (right) under four different spatial models: CSR, HP, DTC, ITC (see text

for model discussion). The model lines are black, the grey area represents the simulation

envelope of 999 Monte Carlo simulations and the coloured lines are the observed spatial

683   distributions For Fractofusus (left) the best-fit model is TC, whereas for Beothukis (right) it is

684   HP because they follow the model best as evidenced by the Monte-Carlo simulations and p_d

685   value (Table S3).

Surface	Taxon	Model fit			P_d values			
		σ	100 ρ	Mean number in cluster	CSR	HP	TC	ITC
Bed B	*Charniodiscus*	**2.906**	**0.7789**	**6**	0.001	0.002	0.892	0.005
	Primocandelabrum	21.47	0.1156	10	0.124	0.002	0.647	0.137
	Charnia	12.027	0.4141	2	0.294	0.266	0.972	0.272
Bristy Cove	*Fractofusus*	NA	NA	NA	0.523	NA	NA	NA
Mistaken Point D	*Fractofusus*	4.312	0.638	3	0.001	0.51	0.77	0.21
	Pectinifrons	NA	NA	NA	0.575	NA	NA	NA
	Bradgatia	NA	NA	NA	0.664	NA	NA	NA
Mistaken Point E	*Bradgatia*	NA	NA	NA	0.44	NA	NA	NA
	Charniodiscus	2.133	1.667	10	0.01	0.13	0.41	0.14
	Beothukis	6.841	0.11	14	0.01	0.90	0.60	0.11
	Feather Dusters	5.616	0.329	11	0.01	0.09	0.28	0.18
	Thectardis	1.835	2.003	4	0.02	0.44	0.91	0.07
	Fractofusus	11.842	0.39	25	0.01	0.03	0.32	0.35
H14	*Fractofusus*	7.906	0.192	2	0.001	0.003	0.760	0.004
St. Shott's	*Aspidella*	9.226	0.4032	4	0.011	0.016	0.963	0.015
	Charnia	10.97	0.2443	2	0.257	0.225	0.331	0.271
Spaniard's Bay	*Avalofractus*	5.607	0.2151	2	0.087	0.142	0.665	0.307
	Beothukis	36.70	0.834	8	0.401	0.810	0.530	0.308
	Trepassia	7.148	0.1941	3	0.005	0.022	0.980	0. 850

Table S3: Summary Table of Univariate PCF analyses. For the heterogeneous

backgrounds, the moving window radius is 0.5 m, using the same taxon density as the taxon

being modelled. $p_d = 1$ corresponds to a perfect fit of the model to the data, while $p_d = 0$

corresponds to no fit. Where observed data did not fall outside CSR Monte-Carlo simulation

envelopes, no further analyses were performed, which is indicated by NA. σ: cluster radius,

ρ: density of specimens, CSR: Complete spatial randomness, HP: Heterogeneous Poisson

model, TC: Thomas Cluster model and ITC: inhomogeneous Thomas Cluster model. Note

that if the cluster model is not a good fit, the *mean number in cluster* will not necessarily be

appropriate.

Surface	Taxon 1	Taxon 2	Model	Output values		
				σ (m)	Cluster Size	p-value
Spaniard's Bay	*Beothukis*	*Trepassia*	**LCM**	**36.70**	**8**	**0.797**
	Bradgatia	***Beothukis***	SS	7.43	14	0.166
Bed B	*Charnia*	*Primocandelabrum*	LCM	**21.47**	**13**	**0.006**
	Charnia	*Primocandelabrum*	SS	**8.571**	**17**	**0.786**

Table S4. Bivariate parameters for the best-fit models for the aggregated distributions.

The parameters used for the shared source models (SS), linked cluster models (LCM) and

linked double cluster models (LDCM). $p_d = 1$ corresponds to a perfect fit of the model to the

data, while $p_d = 0$ corresponds to no fit at all.

Supplementary Discussion

Erosional analyses

Figure S4: H14 spatial map showing the area mapped for Mitchell et al. 2015 study (red

outline). Spatial maps show the positions of the fossil specimens, with the size of the circle

indicated the length of the fossils (*Fractofusus*) or height (fronds) (indicated by a circle).

Black scale bar = 1 m. Different colors indicate different taxa as follows: *Thectardis* navy;

Fractofusus light blue; *Charnia* bright yellow; *Charniodiscus* dark red; Ivesheadiomorphs

dark grey.

Supplementary References

S1.   Mitchell, E. G., Kenchington, C. G., Liu, A. G., Matthews, J. J., & Butterfield, N. J. (2015). Reconstructing the reproductive mode of an Ediacaran macro-organism. *Nature*, 524: 343-346.

S2.   Narbonne, G. M. (2004). Modular construction of early Ediacaran complex life forms. *Science*, 305:1141-1144.

S3.   Brasier, M. D., Liu, A. G., Menon, L., Matthews, J. J., McIlroy, D., & Wacey, D. (2013). Explaining the exceptional preservation of Ediacaran rangeomorphs from Spaniard's Bay, Newfoundland: a hydraulic model. *Precambrian Research*, 231: 122-135.