



## OPEN

# Characterizations of how species mediate ecosystem properties require more comprehensive functional effect descriptors

R. Hale<sup>1,3</sup>, M. N. Mavrogordato<sup>2</sup>, T. J. Tolhurst<sup>1</sup> & M. Solan<sup>3</sup>

<sup>1</sup>University of East Anglia, School of Environmental Science, Norwich, NR4 7TJ, UK, <sup>2</sup>University of Southampton, Engineering and the Environment, Highfield, Southampton, SO17 1BJ, UK, <sup>3</sup>Ocean and Earth Science, National Oceanography Centre, Southampton, University of Southampton, Waterfront Campus, European Way, Southampton, SO14 3ZH, UK.

The importance of individual species in mediating ecosystem process and functioning is generally accepted, but categorical descriptors that summarize species-specific contributions to ecosystems tend to reference a limited number of biological traits and underestimate the importance of how organisms interact with their environment. Here, we show how three functionally contrasting sediment-dwelling marine invertebrates affect fluid and particle transport - important processes in mediating nutrient cycling - and use high-resolution reconstructions of burrow geometry to determine the extent and nature of biogenic modification. We find that individual functional effect descriptors fall short of being able to adequately characterize how species mediate the stocks and flows of important ecosystem properties and that, in contrary to common practice and understanding, they are not substitutable with one another because they emphasize different aspects of species activity and behavior. When information derived from these metrics is combined with knowledge of how species behave and modify their environment, however, detailed mechanistic information emerges that increases the likelihood that a species functional standing will be appropriately summarized. Our study provides evidence that more comprehensive functional effect descriptors are required if they are to be of value to those tasked with projecting how altered biodiversity will influence future ecosystems.

Changes in the structure and composition of communities driven by environmental variability and anthropogenic forcing have concomitant effects on biogeochemical cycles and other important ecosystem processes and functions of global importance<sup>1</sup>, but the direction and magnitude of response is highly dependent on the functional contributions of the modified extant community<sup>2</sup>. Changes to the portfolio of functional effect traits within an assemblage can be more important than alterations to species richness<sup>3</sup>, and has renewed interest in developing approaches<sup>4</sup> that relate observed changes in ecosystem properties to the functional attributes of species. This is appealing because consideration of functional effect traits can offer mechanistic explanations of how changes in the biotic community result in transformations in functioning at habitat or ecosystem scales<sup>5</sup>, which is not always possible when relating changes in function to more abstract or less directly connected concepts, such as species relatedness or altered phylogenetic diversity<sup>6</sup>.

When assigning individual species to functional effect groupings, a fundamental requirement is that the relevant aspects of how a species mediates specific ecosystem properties have been accounted for and characterized. In order to be useful, summary functional effect descriptors need to provide the most relevant representation possible and be sufficiently comprehensive so that they provide acceptable explanatory power for observed species contributions to ecosystem function<sup>7</sup>. However, at least for marine benthic communities, this requirement is seldom fulfilled or tested<sup>8-10</sup>, and complementary functional descriptors are rarely simultaneously presented<sup>11</sup> or combined to improve fit<sup>2</sup>. Furthermore, in practice, assigning individual species to specific functional effect groupings tends to be based on perceived contributions to functioning *per se*<sup>12</sup>, rather than on direct measurements of specific functions obtained in the presence of known species<sup>13</sup>. A fundamental problem is that the importance of organism-sediment interactions in determining how traits are expressed and mediate function is largely ignored<sup>11,14</sup>; in particular, the effects of biogenic structures, such as tubes, burrows and mounds, have not

## SUBJECT AREAS:

COMMUNITY ECOLOGY

ECOSYSTEM ECOLOGY

BIOGEOCHEMISTRY

CLIMATE-CHANGE ECOLOGY

Received

30 June 2014

Accepted

1 September 2014

Published

24 September 2014

Correspondence and requests for materials should be addressed to R.H. ([r.hale@soton.ac.uk](mailto:r.hale@soton.ac.uk))



been considered as useful descriptors when determining the full extent of a species influence on ecosystem properties. Instead, functional effect groupings have become synonymous with trophic feeding position, morphological or physiological characteristics, or descriptions of specific sets of behavior<sup>15</sup>, restricting interpretative capacity of how species and assemblages mediate ecosystem process and functioning to a subset of the total inventory of effect parameters<sup>16</sup>. Rather than diversify the number of available functional effect parameters to address this shortcoming, recent efforts have focused on the details of how species contributions alter in relation to changes in context<sup>5,17–19</sup>. Whilst these have been informative, the mainstay of these studies perpetuates a limited view of what constitutes the functional role of a species. Any derived indices or coefficients, therefore, typically offer low levels of explanatory power and are of limited value in explaining the mechanistic basis for biodiversity-function relations<sup>20</sup>.

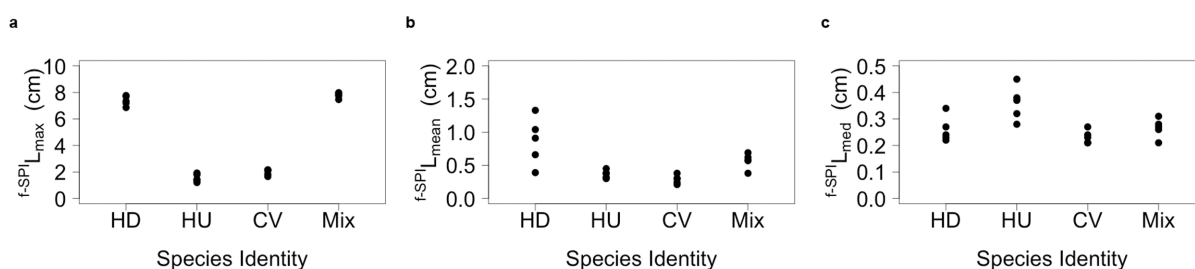
Despite evidence of their pivotal influence in shaping ecosystem properties<sup>21–23</sup>, the modifying effects of biogenic structures and species behaviour (e.g. burrows, mounds, pits, tubes, irrigation and/or feeding behaviour<sup>24</sup>) remain a less prominent feature of functional classifications<sup>15</sup>. Whilst context-dependent changes in species behaviour (e.g. in relation to food supply<sup>18</sup>, tidal periodicity<sup>25</sup>, temperature<sup>26</sup> or photoperiod<sup>27</sup>) and key features of biogenic structures, including length, diameter, volume and configuration, have been described in detail for individual species<sup>28–30</sup> and communities<sup>31</sup>, there is a paucity of information on whether such information is of utility in distinguishing the functional standing of individual species<sup>32–34</sup>. Here, we investigate the functional effect of three co-occurring species of benthic invertebrates, alone and in mixture, using a combination of high-resolution sediment profile imaging and computerized tomography that, together, allow multiple components of particle and fluid transport to be assessed alongside important aspects of burrow morphology at functionally relevant scales (microns to mm). Our focus is to determine whether commonly used alternative functional effect descriptors provide consistent information, and to investigate whether assessments of species contributions to benthic ecosystem process will benefit from the consideration of species behavior (bioirrigation) and/or various properties of associated biogenic structures.

## Results

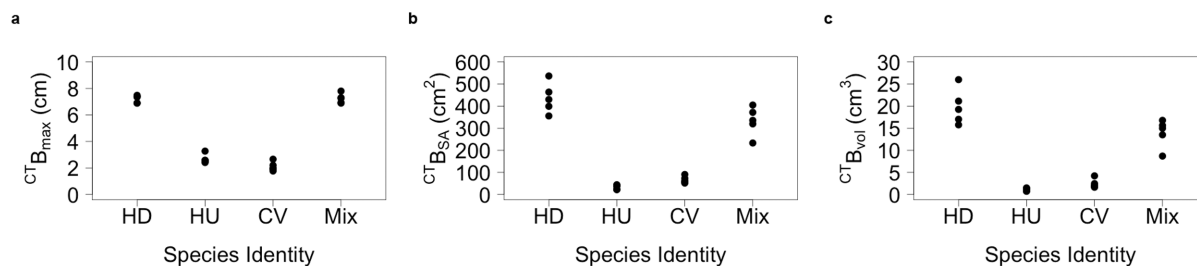
The two-dimensional imaging of redistributed tracer particles (Supplementary Figures S2–S5) revealed little intra-specific variation but consistent inter-specific contrasts in redistribution profiles (Supplementary Figure S6). The rank order of species contributions, however, reflected subtle differences in the functional emphasis of each response variable (compare panels in Figure 1). The species contributing the greatest effect on particle redistribution were *Hediste diversicolor* for  $f\text{-SPIL}_{\text{mean}}$ , *Hydrobia ulvae* for  $f\text{-SPIL}_{\text{med}}$  and the species mixture for  $f\text{-SPIL}_{\text{max}}$ . Overall, the maximum depth of vertical displacement ( $f\text{-SPIL}_{\text{max}}$ ,  $n = 5$ ,  $\pm 95\%$  CI; Figure 1a) ranged from  $1.546 \pm 0.374$  cm for *Hydrobia ulvae* to  $7.788 \pm 0.249$  cm for

species in mixture, whilst the mean depth ( $f\text{-SPIL}_{\text{mean}}$ ,  $n = 5$ ,  $\pm 95\%$  CI; Figure 1b) ranged from  $0.286 \pm 0.081$  cm for *Corophium volutator* to  $0.866 \pm 0.446$  cm for *H. diversicolor* and the median depth ( $f\text{-SPIL}_{\text{med}}$ ,  $n = 5$ ,  $\pm 95\%$  CI; Figure 1c) ranged from  $0.232 \pm 0.031$  cm for *C. volutator* to  $0.360 \pm 0.080$  cm for *H. ulvae*. Closer examination of the minimum adequate models revealed distinct inter-specific differences in the depth of  $f\text{-SPIL}_{\text{max}}$  (L-ratio = 68.592, d.f. = 3,  $p < 0.0001$ ; Supplementary Model S3),  $f\text{-SPIL}_{\text{mean}}$  (L-ratio = 19.182, d.f. = 3,  $p = 0.0003$ ; Supplementary Model S1), and  $f\text{-SPIL}_{\text{med}}$  (L-ratio = 11.505, d.f. = 3,  $p = 0.0093$ ; Supplementary Model S2), but despite such subsurface mixing we were unable to determine any notable effect on surface boundary roughness (intercept only model,  $F = 0.3446$ , d.f. = 3,  $p = 0.7935$ ; Supplementary Model S4). For  $f\text{-SPIL}_{\text{max}}$ , these inter-specific differences were distinct across all possible pairwise species permutations, with the effects of *C. volutator* (coefficient  $\pm$  s.e. =  $-5.399 \pm 0.195$ ,  $t = -27.66$ ,  $p < 0.0001$ ) and *H. ulvae* (coefficient  $\pm$  s.e. =  $-5.831 \pm 0.215$ ,  $t = -27.19$ ,  $p < 0.0001$ ) less than those of *H. diversicolor*, and the effects of the species mixture (coefficient  $\pm$  s.e. =  $0.411 \pm 0.189$ ,  $t = 2.17$ ,  $p = 0.0454$ ) greater than those of *H. diversicolor*. For  $f\text{-SPIL}_{\text{mean}}$  (Figure 1b), the activities of *H. diversicolor* and the species mixture were indistinguishable from one another (coefficient  $\pm$  s.e. =  $-0.298 \pm 0.169$ ,  $t = -1.766$ ,  $p = 0.097$ ) and resulted in deeper mean mixing levels than those of *C. volutator* and *H. ulvae*, which were also similar, albeit marginally, to one another (coefficient  $\pm$  s.e. =  $0.080 \pm 0.039$ ,  $t = 2.031$ ,  $p = 0.059$ ) but different to those of *H. diversicolor* and the species mixture (compare coefficients, Supplementary Model S1). The patterns of mixing shown by  $f\text{-SPIL}_{\text{med}}$  (Figure 1c) reveal that the activities of *H. ulvae* (mean depth,  $n = 5$ ,  $\pm 95\%$  CI,  $= 0.360 \pm 0.08$  cm) were greater than those of *H. diversicolor* (coefficient  $\pm$  s.e. =  $0.100 \pm 0.036$ ,  $t = 2.774$ ,  $p = 0.014$ ), *C. volutator* (coefficient  $\pm$  s.e. =  $0.128 \pm 0.031$ ,  $t = 4.144$ ,  $p < 0.001$ ) and the species mixture (coefficient  $\pm$  s.e. =  $0.094 \pm 0.033$ ,  $t = 2.839$ ,  $p = 0.012$ ), which were all insignificant from one another (compare coefficients, Supplementary Model S2). Interestingly, in the case of  $f\text{-SPIL}_{\text{med}}$ , the degree of mixing in the species mixture did not necessarily reflect that of the most dominant species (*H. ulvae*) contribution in monoculture. Indeed, both *H. diversicolor* and *C. volutator* were clearly the most influential species across all three measures of particle redistribution (compare panels, Figure 1).

Reconstruction of biogenic structures using computer tomography techniques revealed species-specific burrow systems that differed markedly in burrow density, structural morphology and depth distribution (Figure 2, Supplementary Figures S7–S9 and Supplementary Movie Sequences 1–4). *Hediste diversicolor* constructed extensive branched burrow galleries throughout the sediment profile ( $CTB_{\text{max}}$  range, 6.89–7.48 cm), *Corophium volutator* constructed multiple U-shape burrows that were confined to the surficial layers ( $CTB_{\text{max}}$  range, 1.77–2.66 cm), and *Hydrobia ulvae* constructed vertical I-shape excavations, albeit infrequently, to depths in excess of 2.4 cm ( $CTB_{\text{max}}$  range, 2.41–3.27 cm). These clear distinctions in species burrowing behavior led to concomitant



**Figure 1 | Sediment reworking associated with benthic invertebrates.** Species-specific effects ( $n = 5$ ) on the particle reworking metrics (a)  $f\text{-SPIL}_{\text{max}}$ , (b)  $f\text{-SPIL}_{\text{mean}}$ , and (c)  $f\text{-SPIL}_{\text{med}}$ . HD = *Hediste diversicolor*, HU = *Hydrobia ulvae*, CV = *Corophium volutator*, and Mix = species mixture comprising equal proportions (biomass ratio, 1 : 1 : 1) of HD, HU and CV.



**Figure 2** | The extent of species-environment interaction associated with benthic invertebrates. Species-specific effects ( $n = 5$ ) on burrow geometry metrics (a)  $^{CT}B_{max}$ , (b)  $^{CT}B_{SA}$ , and (c)  $^{CT}B_{vol}$ . HD = *Hediste diversicolor*, HU = *Hydrobia ulvae*, CV = *Corophium volutator*, and Mix = species mixture comprising equal proportions (biomass ratio, 1 : 1 : 1) of HD, HU and CV.

changes in the maximum depth ( $^{CT}B_{max}$ , Figure 2a), surface area ( $^{CT}B_{SA}$ , Figure 2b) and lumen volume ( $^{CT}B_{vol}$ , Figure 2c) of the burrows but, when species were combined in mixture, the total was not necessarily additive (compare monoculture to mixture performance, Figure 2) and did not always match the maximum contributing species (comparison of mixture to *H. diversicolor* for:  $^{CT}B_{max}$ , coefficient  $\pm$  s.e. =  $0.034 \pm 0.213$ ,  $t = 0.160$ ,  $p = 0.875$ ;  $^{CT}B_{SA}$ , coefficient  $\pm$  s.e. =  $-103.98 \pm 42.17$ ,  $t = -2.466$ ,  $p = 0.025$ ;  $^{CT}B_{vol}$ , coefficient  $\pm$  s.e. =  $-5.914 \pm 2.283$ ,  $t = -2.591$ ,  $p = 0.020$ ; Supplementary Models S5–S7). For  $^{CT}B_{max}$ , the activities of *H. diversicolor* and the species mixture were indistinguishable from one another (coefficient  $\pm$  s.e. =  $0.034 \pm 0.213$ ,  $t = 0.160$ ,  $p = 0.875$ ) and were deeper than those of *H. ulvae* and they, in turn, were deeper than those of *C. volutator* (compare coefficients, Supplementary Model S5).  $^{CT}B_{SA}$  and  $^{CT}B_{vol}$  were, as expected, highly correlated ( $r = 0.99$ ) and showed the same sequence of species performance (HD  $\geq$  Mix > CV > HU; compare coefficients, Supplementary Models S6–S7).  $^{CT}B_{SA}$  and  $^{CT}B_{vol}$  in species mixture were intermediate relative to performance observed in respective monocultures (Figure 2).

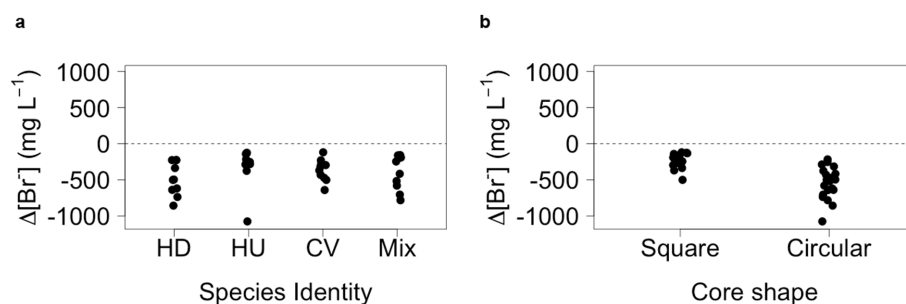
There were independent effects of species identity and core shape on bioirrigation behavior (Figure 3). For species, most bioirrigation activity (mean  $\Delta[Br^-]$ ,  $n = 10$ ,  $\pm 95\%$  CI =  $-486.79 \pm 162.92$  mg  $L^{-1}$ ) was performed by *H. diversicolor*, followed by the remaining species treatments (coefficient  $\pm$  s.e.: CV,  $103.37 \pm 53.16$ ,  $t = 1.944$ ,  $p = 0.060$ ; Mix,  $126.96 \pm 45.27$ ,  $t = 2.805$ ,  $p = 0.008$ ; HU,  $139.89 \pm 53.03$ ,  $t = 2.638$ ,  $p = 0.012$ ; Figure 3a), which were generally indistinguishable from one another (compare coefficients, Supplementary Model S8). In general, irrespective of species identity, there was a greater amount of bioirrigation (mean  $\Delta[Br^-]$ ,  $n = 20$ ,  $\pm 95\%$  CI) in the circular cores (circular,  $-546.15 \pm 102.11$  mg  $L^{-1}$ ; square,  $-234.11 \pm 43.79$  mg  $L^{-1}$ ) relative to the square cores (coefficient  $\pm$  s.e. =  $-325.20 \pm 41.88$ ,  $t = -7.765$ ,  $p < 0.0001$ ; Figure 3b).

For the components of particle redistribution, properties of biogenic structure and bioirrigation behavior that were measured, we found no evidence of overyielding (Figure 4). With the exception of

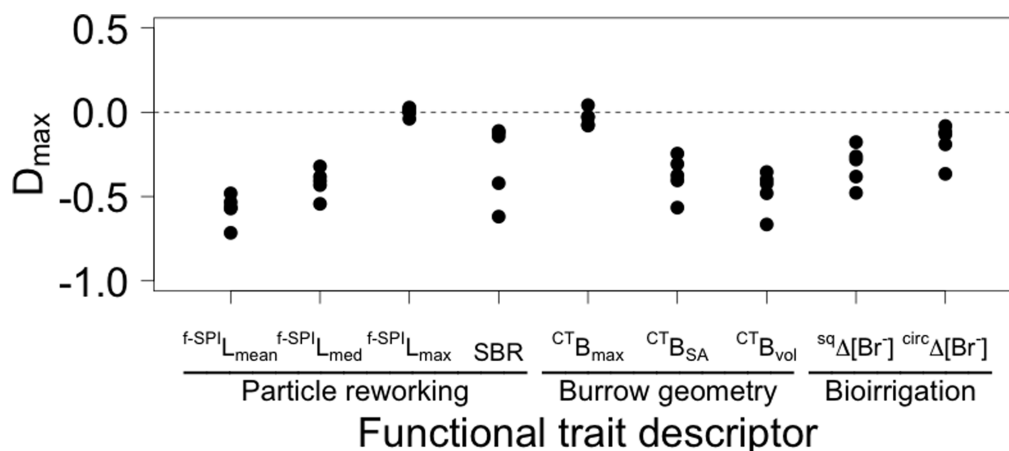
$^{f-SPI}L_{max}$  and  $^{CT}B_{max}$ , which maintain the equivalent level of process to *H. diversicolor* in monoculture, all other response variables exhibit a change in the mean ( $\pm 95\%$  CI,  $n = 5$ ) level of process when in species mixture (ranging from  $-0.177 \pm 0.139$  for  $^{circ}\Delta[Br^-]$  to  $-0.573 \pm 0.108$  for  $^{f-SPI}L_{mean}$ ; Figure 4).

## Discussion

We have demonstrated distinct species-specific differences in the redistribution of sediment particles, fluid transport, and the relative geometric properties of biogenic structures, three major mechanisms by which species mediate important ecosystem functions in the marine benthos<sup>15</sup>. Our data show, however, that evaluations of the contributions that species make to each of these categories can be misleading when based on a single, or a limited set of, functional effect descriptors. We find that different properties of routinely used data on the vertical redistribution of sediment particles reflect either short-term infaunal activity in the uppermost region of the sediment profile ( $^{f-SPI}L_{med}$ ), less frequently occurring infaunal activity at depth ( $^{f-SPI}L_{max}$ ), or the relative balance between the minimum and maximum depth of mixing ( $^{f-SPI}L_{mean}$ ). Similarly, bioirrigation ( $\Delta[Br^-]$ ) poorly related to burrow geometry, suggesting that the propensity to irrigate burrows was more important than the dimensions of burrow structures<sup>35</sup>. Surface boundary roughness was unable to discriminate between species, despite contrasts in the lifestyle and residency depth of the species under study, as all species were capable of redistributing surficial sediment. Importantly, these differences in emphasis across closely related functional effect descriptors lead to contradictory interpretations of the functional standing of a species because each descriptor integrates aspects of species behavior that are not universally attributable. Moreover, it is clear that routinely used proxies for various ecosystem processes only provide partial coverage of the mechanisms that contribute to functioning. Our analyses show, however, that these interpretative inconsistencies can be reconciled, and poor levels of discriminatory power can be improved, when elements of species behavior and/or properties of biogenic



**Figure 3** | Independent effects of (a) species identity and (b) core shape on bioirrigation activity ( $\Delta[Br^-]$ , mg  $L^{-1}$ ). Negative values indicate increased bioirrigation activity. For clarity, jitter ( $\alpha = 0.05$ ) has been applied to the  $x =$  argument of the plot function to avoid over-plotting. HD = *Hediste diversicolor*, HU = *Hydrobia ulvae*, CV = *Corophium volutator*, and Mix = species mixture comprising equal proportions (biomass ratio, 1 : 1 : 1) of HD, HU and CV.



**Figure 4** | Observed yield ( $D_{\max}$ ) of different components of particle reworking, burrow geometry and bioirrigation behaviour for species mixtures relative to species monocultures with equivalent biomass. The observed yield indicates whether a species mixture outperforms (overyielding,  $D_{\max} > 0$ ) or underperforms (underyielding,  $D_{\max} < 0$ ) relative to the maximally performing species in monoculture, and provides an indication of how species interactions may influence species contributions to ecosystem process.

structures are considered in concert with other ‘biological’ derived traits because the former are critical in determining the functional expression of traits<sup>11,14</sup>.

Here, we used three-dimensional reconstructions of burrow geometry ( $CT_{B_{SA}}$ ,  $CT_{B_{vol}}$ ,  $CT_{B_{max}}$ ) and two dimensional assessments of surface boundary roughness and bioirrigation activity ( $\Delta[Br^-]$ ) to gain important insights about the relative importance of the interaction between species behavior and biogenic structures. We found that there were distinct inter-specific differences in species behavior and burrow morphology (*H. diversicolor*, extensive inter-weaving galleries extending to ~7.5 cm depth with multiple openings at the sediment-water interface; *C. volutator*, U-shaped burrows extending to ~2.6 cm depth; *H. ulvae*, I- and J-shaped vertical shafts to ~3.3 cm) that dictate whether fluid transport is primarily diffusive or advective<sup>36</sup>. Our methodology did not allow us to determine the temporal pattern of bioirrigation activity, but we were able to identify distinct species-specific differences in relative performance. Whilst some of this information is known, at least qualitatively, it is important to emphasize that our approach of using multiple functional effect descriptors was instrumental in providing novel functionally relevant information. For instance, the behavior<sup>37,38</sup> and effects of *H. ulvae* on ecosystem process and functioning have been extensively described<sup>5,33,39,40</sup>, but the spatial distribution, morphology and significance of their burrow system has not previously been recognized or quantified. Prior summaries of the functional role of *H. ulvae* have emphasized surficial sediment destabilisation through surface browsing and disruption<sup>41</sup>, but the data presented here suggests that *H. ulvae* significantly contributes to much deeper sediment destabilisation, especially in areas of high sediment moisture content<sup>38</sup> or patchy resource supply<sup>5</sup> where activity levels are elevated. Such information provides mechanistic explanations for otherwise inconsistent observations of species contributions to ecosystem functioning, and reinforces the view that the functional role of a species is not static and cannot be adequately summarized without reference to variation in species behavior and use of multiple functional parameters<sup>42</sup>.

Whilst we were able to identify inter-specific differences in functional effects across our monocultures, our results also highlight the importance of considering whether the functional role of species changes when in the presence of other species. We found that particle reworking, bioirrigation, burrow surface area and burrow volume all fell below expected values when species were in mixture (underyielding,  $D_{\max} < 0$ ), an outcome consistent with findings elsewhere<sup>1,43</sup> that highlights the importance of the modifying effects (positive

and negative) of species interaction<sup>1</sup>. We interpret these effects to be in relation to competition for space and/or resources<sup>44,45</sup>, rather than being due to density effects<sup>46,47</sup>, lending support to the view that the functional role of a species may reflect a behavioral response to the biotic setting rather than the presence or absence of particular biological traits<sup>48</sup>; the exploratory and prospecting behavior of *H. diversicolor*, for example, is known to disrupt *C. volutator* burrows, resulting in the relocation of individuals and additional burrow construction activity<sup>5,17,39</sup> that alters the relative amount of time allocated to other functionally important behaviors, such as bioirrigation<sup>49</sup>. Such context-dependent behavioural responses constitute activity that is additional to, and distinct from, activity associated with routine behavior<sup>17</sup> and may fundamentally change the functional role of a species over extended timescales. An immediate challenge, therefore, is to distinguish the functional effect of transient changes in behavior<sup>5,18,25–27</sup> from those associated with acclimation or adaptation to novel abiotic and biotic circumstances<sup>50</sup>. This matters because the structural properties of a community may well be conserved after a perturbation, or following a period of directional forcing, but the functional role of individual species may fundamentally change relative to the pre-disturbance standard<sup>50,51</sup>. Failure to collate this information dramatically reduces the accuracy and predictive capabilities of next generation ecosystem models but, as the effects of core shape reveal in the present study, it will also be important to ensure that efforts towards this goal standardize experimental protocols and account for differences in experimental settings<sup>52,53</sup>. Nevertheless, a substantial revision of previously described functional effect groupings is required before the true value of biodiversity for humanity can be ascertained.

## Methods

**Surficial sediment** (less than 3 cm depth; mean particle size, 54.80  $\mu\text{m}$ ; mud content, 55.93%) and three co-occurring functionally contrasting inter-tidal invertebrates (the polychaete *Hediste diversicolor*, the gastropod *Hydrobia ulvae* and mud shrimp *Corophium volutator*) were collected from the mid-shore at Breydon water, Great Yarmouth, UK (N52° 37.030', E01° 41.390') and returned to the *Biodiversity and Ecosystem Futures Facility* at the University of Southampton to acclimatise to laboratory conditions (5 days). Sediment was sieved (500  $\mu\text{m}$  mesh) in a seawater (sand filtered, UV sterilized and salinity 33) bath to remove macrofauna, allowed to settle for 48 h to retain the fine fraction (less than 63  $\mu\text{m}$ ) and homogenized. Each aquarium was filled to a depth of 8 cm with sediment homogenate overlain by 4 cm seawater. Fauna were not added until the lower regions of the cores showed evidence of reducing conditions (visible anoxic microniche formation). Overlying seawater was replaced after 24 h to remove excess nutrients associated with assembly. Aquaria were maintained at  $12 \pm 0.1^\circ\text{C}$  under a 12:12 h light (Aqualine T5 Reef White 10 K fluorescent light tubes, Aqua Medic) cycle and were continually aerated. Although inter-tidal species exhibit a rhythm of activity associated with the tidal cycle, it was not



necessary to incorporate an immersion-emersion cycle into our experimental design because the biological rhythms in the species under study are known to be endogenous, i.e. they persist long after an organism is isolated from periodic tidal cues that alter their behavior.

Replicate ( $n = 5$ ) invertebrate communities were assembled in monoculture (*Hediste diversicolor*, HD; *Hydrobia ulvae*, HU; or *Corophium volutator*, CV) and in a mixture (Mix) of all three species (biomass fixed at 1 g wet weight aquaria<sup>-1</sup>;  $\sim 127 \text{ g m}^{-2}$ , equivalent to a monoculture density of  $\sim 850$ ,  $\sim 22,000$  or  $\sim 25,000 \text{ ind. m}^{-2}$  for HD, HU and CV respectively) in each of two types of aquaria (square vs circular) that shared an identical volume of habitat ( $1178 \text{ cm}^3$ ). Square aquaria (internal dimensions  $8.86 \times 8.86 \text{ cm} \times 15.0 \text{ cm}$ ,  $n = 20$ ) were required to facilitate quantification of particle reworking using 2-dimensional photographic imaging, whilst circular aquaria (internal diameter = 10 cm, 15 cm tall,  $n = 20$ ) were required to facilitate rotational quantification of biogenic structures using 3-dimensional computer tomography (CT) imaging.

Faunal mediated sediment particle reworking in the square aquaria was estimated non-invasively using a sediment profile imaging camera (Canon 400D set to 10 s exposure, aperture f5.6 and ISO 400;  $3888 \times 2592$  pixels, i.e. 10.1 megapixels, effective resolution of sediment  $56 \times 56 \text{ } \mu\text{m per pixel}$ ), optically modified to allow preferential imaging of fluorescently labeled sand-based particulate tracers under UV light (f-SPI<sup>54</sup>). These tracers were evenly distributed over the entire sediment surface ( $\sim 3 \text{ mm depth}$ ) 24 h after faunal addition to ensure that elevated levels of activity associated with initial burrow construction were not incorporated into estimates of infaunal mixing. The redistribution of luminophores (15 g aquarium<sup>-1</sup>, pink colour, size class less than 125  $\mu\text{m}$ , mean particle size  $\sim 80 \text{ } \mu\text{m}$ ; Brian Clegg Ltd., UK) was determined from stitched composite images (RGB colour, JPEG compression) of all four sides of each aquarium obtained in a UV illuminated imaging box<sup>55</sup> after 6 days using a custom-made semi-automated macro that runs within *ImageJ* (Version 1.47), a java-based public domain program developed at the US National Institutes of Health (<http://rsb.info.nih.gov/ij/index.html>, Rasband, W., ImageJ, (1997), Date of access 10/07/2013). From these data, the median ( $f\text{-SPI}_{\text{L}_{\text{med}}}$ , typical short-term depth of mixing), maximum ( $f\text{-SPI}_{\text{L}_{\text{max}}}$ , maximum extent of mixing over the long-term) and mean ( $f\text{-SPI}_{\text{L}_{\text{mean}}}$ , time dependent indication of mixing) mixed depth of particle redistribution were calculated. In addition, the maximum vertical deviation of the sediment-water interface (upper – lower limit = surface boundary roughness, SBR) provided an indication of surficial activity.

Quantification of biogenic structures in the circular aquaria was achieved using a 225/450 kV Nikon/Metris custom designed micro-focus computer tomography scanner housed within the  $\mu$ -VIS imaging centre, University of Southampton. Batches of 5 aquaria were stacked and secured in a custom-made holding brace to ensure stability during scanning. During each acquisition, the aquaria were rotated through  $360^\circ$  whilst collecting 3142 projections averaging over 8 frames per 250 ms projection. X-ray conditions were set to 300 kV and 326  $\mu\text{A}$  with a 3 mm Cu filter, and an XRD 1621 CN3 H5 PerkinElmer flat panel detector was used to collect the images. In the resulting images, levels of grey scale reflect the level of x-ray attenuation caused by variation in bulk density<sup>56</sup>. Hence, brighter pixels represent denser material (sediment) and darker pixels represent less dense material (burrow voids). Raw image slices ( $n = 2000$  aquarium<sup>-1</sup>, voxel resolution = 81  $\mu\text{m}$ ) were processed as follows: First, the projection data was reconstructed using *CTPro3D* (v. XT 2.2 service pack 10, Nikon Metrology, UK) and *CTAgent* (v. XT 2.2 service pack 10, Nikon Metrology, UK). The reconstructed volumes were converted to 8 bit format using *FIJI*<sup>57</sup> in order to reduce file sizes and computational loading. Finally, these images were opened as a 3D project in *VGStudio* (v. 2.1 Volume Graphics GmbH, Germany). From these data, regions of interest were segmented using a threshold based seed point growing algorithm from which the surface area ( $CT_{\text{BSA}}$ , an important determinant of microbial-mediated biogeochemical cycling), volume ( $CT_{\text{Vol}}$ , an indication of the extent of bioengineering and bioirrigation) and maximum depth of any biogenic features ( $CT_{\text{B}_{\text{max}}}$ ) were calculated.

Bioirrigation was estimated in both square and circular aquaria from absolute changes in the concentration of the inert tracer sodium bromide ( $\Delta[\text{Br}^-]$ ,  $\text{mg l}^{-1}$ ; negative values indicate increased bioirrigation activity<sup>58</sup>) over an 8 h period (day-time, in square and circular cores) on day 6, determined using a Tecator flow injection auto-analyser (FIA Star 5010 series).

In order to assess whether there were any effects of species interactions on our dependent variables, we compared performance in monoculture to performance in mixture<sup>59</sup>. Overyielding ( $D_{\text{max}} > 0$ ) occurs when a mixture outperforms expectations based on the corresponding monocultures.

Linear regression models were developed for the dependent variables ( $f\text{-SPI}_{\text{L}_{\text{med}}}$ ,  $f\text{-SPI}_{\text{L}_{\text{max}}}$ ,  $f\text{-SPI}_{\text{L}_{\text{mean}}}$ , SBR,  $CT_{\text{BSA}}$ ,  $CT_{\text{Vol}}$ ,  $CT_{\text{B}_{\text{max}}}$ ,  $\Delta[\text{Br}^-]$ ) with the independent nominal variable species identity (SPID) or, for bioirrigation, the nominal variables SPID and core shape (square versus circular). In all cases, we found evidence for a violation of homogeneity of variance so a *VarIdent* variance-covariate structure and a generalised least squares (GLS) estimation procedure<sup>60</sup> that allows the residual spread to vary with individual explanatory variables was applied. For all of the bioturbation analyses, the residual spread varied with species identity whilst, for the bioirrigation analysis, the residual spread varied with both core shape and species identity. We determined the optimal fixed-effects structure for each model using backward selection informed by Akaike Information Criteria (AIC) and inspection of model residual patterns. The optimal variance covariate structure was determined by comparing the initial ANOVA model without variance structure to the equivalent GLS model incorporating specific variance structures using AIC and visualisation of model residuals obtained by restricted maximum likelihood (REML) estimation. The optimal fixed

structure was then determined by applying backward selection using the likelihood ratio test obtained using maximum likelihood (ML) estimation. All analyses were performed in R (v. 2.15.2) using the *nlme* package<sup>61</sup>. All minimal adequate models and their associated coefficient tables are presented in supplementary material.

- Cardinale, B. J. *et al.* Biodiversity loss and its impact on humanity. *Nature* **486**, 59–67 (2012).
- Solan, M. *et al.* Extinction and ecosystem function in the marine benthos. *Science* **306**, 1177–1180 (2004).
- Van Colen, C. *et al.* Organism-sediment interactions govern post-hypoxia recovery of ecosystem functioning. *PLOS One* **7**, e49795 (2012).
- Blaum, N., Mosner, E., Schwager, M. & Jeltsch, F. How functional is functional? Ecological groupings in terrestrial animal ecology: towards an animal functional type approach. *Biodiversity and Conservation* **20**, 2333–2345 (2011).
- Godbold, J. A., Bulling, M. T. & Solan, M. Habitat structure mediates biodiversity effects on ecosystem properties. *Proc. R. Soc. B.* **278**, 2510–2518 (2011).
- Cadotte, M. W., Carscadden, K. & Mirotnick, N. Beyond species: functional diversity and the maintenance of ecological processes and services. *J. Appl. Ecol.* **48**, 1079–1087 (2011).
- Stuart-Smith, R. D. *et al.* Integrating abundance and functional traits reveals new global hotspots of fish diversity. *Nature* **501**, 539–542 (2013).
- Teal, L. R., Parker, E. R. & Solan, M. Sediment mixed layer as a proxy for benthic ecosystem process and function. *Mar. Ecol. Prog. Ser.* **414**, 27–40 (2010).
- Birchenough, S. N. R., Parker, R. E., McManus, E. & Barry, J. Combining bioturbation and redox metrics: Potential tools for assessing seabed function. *Ecological Indicators* **12**, 8–16 (2012).
- Séguin, A., Harvey, É., Archambault, P., Nozais, C. & Gravel, D. Body size as a predictor of species loss effect on ecosystem functioning. *Sci. Rep.* **4**, 4616. DOI: 10.1038/srep046162014 (2014).
- Needham, H. R., Pilditch, C. A., Lohrer, A. M. & Thrush, S. F. Context-specific bioturbation mediates changes to ecosystem functioning. *Ecosystems* **14**, 1096–1109 (2011).
- Queirós, A. M. *et al.* A bioturbation classification of European marine infaunal invertebrates. *Ecology and Evolution* **3**, 3958–3985 (2013).
- Murray, F., Douglas, A. & Solan, M. Species that share similar traits do not necessarily form distinct and universally applicable functional effect groups. *Mar. Ecol. Prog. Ser.* doi:10.3354/meps11020 (In press).
- Needham, H. R., Pilditch, C. A., Lohrer, A. M. & Thrush, S. F. Habitat dependence in the functional traits of *Austrohelice crassa*, a key bioturbating species. *Mar. Ecol. Prog. Ser.* **414**, 179–193 (2010).
- Pearson, T. H. Functional group ecology in soft-sediment marine benthos: The role of bioturbation. *Oceanogr. Mar. Biol. Ann. Rev.* **39**, 233–267 (2001).
- Teal, L. R., Parker, E. R. & Solan, M. Coupling bioturbation activity to metal (Fe and Mn) profiles *in situ*. *Biogeochemistry* **10**, 2365–2378 (2013).
- Bulling, M. *et al.* Species effects on ecosystem processes are modified by faunal responses to habitat composition. *Oecologia* **158**, 511–520 (2008).
- Nogaro, G. *et al.* Food supply impacts sediment reworking by *Nereis diversicolor*. *Hydrobiologia* **598**, 403–408 (2008).
- Maire, O. *et al.* Indirect effects of non-lethal predation on bivalve activity and sediment reworking. *J. Exp. Mar. Biol. Ecol.* **395**, 30–36 (2010).
- Solan, M., Batty, P., Bulling, M. T. & Godbold, J. A. How biodiversity affects ecosystem processes: implications for ecological revolutions and benthic ecosystem function. *Aquatic Biology* **2**, 289–301 (2008).
- Aller, R. C. The importance of relict burrow structures and burrow irrigation in controlling sedimentary solute distributions. *Geochim. Cosmochim. Acta* **48**, 1929–1934 (1984).
- Davey, J. T. The architecture of the burrow of *Nereis diversicolor* and its quantification in relation to sediment water exchange. *J. Exp. Mar. Biol. Ecol.* **179**, 115–129 (1994).
- Kinoshita, K., Wada, M., Kogure, K. & Furota, T. Mud shrimp burrows as dynamic traps and processors of tidal-flat materials. *Mar. Ecol. Prog. Ser.* **247**, 159–164 (2003).
- Nilsson, H. C. & Rosenberg, R. Benthic habitat quality assessment of an oxygen stressed fjord by surface and sediment profile images. *J. Mar. Syst.* **11**, 249–264 (1997).
- Solan, M. & Kennedy, R. Observation and quantification of *in situ* animal-sediment relations using time-lapse sediment profile imagery (t-SPI). *Mar. Ecol. Prog. Ser.* **228**, 179–191 (2002).
- Ouellette, D. *et al.* Effects of temperature on *in vitro* sediment reworking processes by a gallery bioinfiltrator, the polychaete *Neanthes virens*. *Mar. Ecol. Prog. Ser.* **266**, 185–193 (2004).
- Rosenberg, R. & Lundberg, L. Photoperiodic activity pattern in the brittle star *Amphiura filiformis*. *Mar. Biol.* **145**, 651–656 (2004).
- Coelho, V. R., Cooper, R. A. & Rodrigues, S. D. Burrow morphology and behavior of the mud shrimp *Upogebia omisssa* (Decapoda: Thalassinidea: Upogebiidae). *Mar. Ecol. Prog. Ser.* **200**, 229–240 (2000).
- Yunusa, I. A. M., Braun, M. & Lawrie, R. Amendment of soil with coal fly ash modified the burrowing habits of two earthworm species. *Appl. Soil Ecol.* **42**, 63–68 (2009).



30. Salvo, F., Dufour, S. C., Archambault, P., Stora, G. & Desrosiers, G. Spatial distribution of *Alitta virens* burrows in intertidal sediments studied by axial tomodesitometry. *J. Mar. Biol. Assoc. U.K.* **93**, 1543–1552 (2013).
31. Rosenberg, R., Gremare, A., Duchene, J. C., Davey, E. & Frank, M. 3D visualization and quantification of marine benthic biogenic structures and particle transport utilizing computer-aided tomography. *Mar. Ecol. Prog. Ser.* **363**, 171–182 (2008).
32. Bouchet, V. M. P. *et al.* Influence of the mode of macrofauna-mediated bioturbation on the vertical distribution of living benthic foraminifera: First insight from axial tomodesitometry. *J. Exp. Mar. Biol. Ecol.* **371**, 20–33 (2009).
33. Gilbertson, W. W., Solan, M. & Prosser, J. I. Differential effects of microorganism-invertebrate interactions on benthic nitrogen cycling. *FEMS Microbiology Ecology* **82**, 11–22 (2012).
34. Passarelli, C., Olivier, F., Paterson, D. M. & Hubas, C. Impacts of biogenic structures on benthic assemblages: microbes, meiofauna, macrofauna and related ecosystem functions. *Mar. Ecol. Prog. Ser.* **465**, 85–97 (2013).
35. Renz, J. R. & Forster, S. Effects of bioirrigation by the three sibling species of *Marenzelleria* spp. on solute fluxes and porewater nutrient profiles. *Mar. Ecol. Prog. Ser.* **505**, 145–159 (2014).
36. Meysman, F. J. R., Galaktionov, E. S., Gribsholt, B. & Middelburg, J. J. Bioirrigation in permeable sediments: Advective pore-water transport induced by burrow ventilation. *Limnol. Oceanogr.* **51**, 142–156 (2006).
37. Little, C. & Nix, W. The burrowing and floating behaviour of the gastropod *Hydrobia ulvae*. *Estuar. Coast. Mar. Sci.* **4**, 537–544 (1976).
38. Orvain, F. & Sauriau, P. G. Environmental and behavioural factors affecting activity in the intertidal gastropod *Hydrobia ulvae*. *J. Exp. Mar. Biol. Ecol.* **272**, 191–216 (2002).
39. Dyson, K. E. *et al.* Influence of macrofaunal assemblages and environmental heterogeneity on microphytobenthic production in experimental systems. *Proc. R. Soc. B.* **274**, 2547–2554 (2007).
40. Hicks, N. *et al.* Impact of biodiversity-climate futures on primary production and metabolism in a model benthic estuarine system. *BMC Ecology* **11**, 7. DOI:10.1186/1472-6785-11-7 (2011).
41. Orvain, F., Sauriau, P. G., Bacher, C. & Prineau, M. The influence of sediment cohesiveness on bioturbation effects due to *Hydrobia ulvae* on the initial erosion of intertidal sediments: A study combining flume and model approaches. *J. Sea Res.* **55**, 54–73 (2006).
42. Hector, A. & Bagchi, R. Biodiversity and ecosystem multifunctionality. *Nature* **448**, 188–190 (2007).
43. Mermillod-Blondin, F., Francois-Carcaillet, F. & Rosenberg, R. Biodiversity of benthic invertebrates and organic matter processing in shallow marine sediments: an experimental study. *J. Exp. Mar. Biol. Ecol.* **315**, 187–209 (2005).
44. Miron, G., Desrosiers, G., Retiere, C. & Masson, S. Variations in time budget of the polychaete *Nereis virens* as a function of density and acclimation after introduction to a new burrow. *Mar. Biol.* **114**, 41–48 (1992).
45. van Nugteren, P. *et al.* Spatial distribution of detrital resources determines the outcome of competition between bacteria and a facultative detritivorous worm. *Limnol. Oceanogr.* **54**, 1413–1419 (2009).
46. Dupont, E., Stora, G., Tremblay, P. & Gilbert, F. Effects of population density on the sediment mixing induced by the gallery-diffuser *Hediste (Nereis) diversicolor* O.F. Müller, 1776. *J. Exp. Mar. Biol. Ecol.* **336**, 33–41 (2006).
47. De Backer, A. *et al.* Bioturbation effects of *Corophium volutator*: Importance of density and behavioural activity. *Estuar. Coast. Shelf Sci.* **91**, 306–313 (2011).
48. Caliman, A., Carneiro, L. S., Bozelli, R. L., Farjalla, V. F. & Esteves, F. A. Bioturbating space enhances the effects of non-additive interactions among benthic ecosystem engineers on cross-habitat nutrient regeneration. *Oikos* **120**, 1639–1648 (2011).
49. De Backer, A., van Ael, E., Vincx, M. & Degraer, S. Behaviour and time allocation of the mud shrimp, *Corophium volutator*, during the tidal cycle: a laboratory study. *Helgol. Mar. Res.* **64**, 63–67 (2010).
50. Godbold, J. A. & Solan, M. Long-term effects of warming and ocean acidification are modified by seasonal variation in species responses and environmental conditions. *Phil. Trans. R. Soc. B.* **368**, UNSP 20130186 (2013).
51. Nielsen, A. & Totland, O. Structural properties of mutualistic networks withstand habitat degradation while species functional roles might change. *Oikos* **123**, 323–333 (2014).
52. Du Clos, K. T., Lindsay, S. M. & Jumars, P. A. Spatial distribution of *Alitta virens* and *Clymenella torquata* with respect to rigid boundaries in mud and sand. *J. Mar. Res.* **71**, 211–226 (2013).
53. Lindqvist, S., Gilbert, F., Eriksson, S. P. & Hulth, S. Activities by *Hediste diversicolor* under different light regimes: Experimental quantification of particle reworking using time-resolved imaging. *J. Exp. Mar. Biol. Ecol.* **448**, 240–249 (2013).
54. Solan, M. *et al.* *In situ* quantification of bioturbation using time-lapse fluorescent sediment profile imaging (f-SPI), luminophore tracers and model simulation. *Mar. Ecol. Prog. Ser.* **271**, 1–12 (2004).
55. Schiffrs, K., Teal, L. R., Travis, J. M. J. & Solan, M. An open source simulation model for soil and sediment bioturbation. *PLoS ONE* **6**, e28028. doi:10.1371/journal.pone.0028028 (2011).
56. Wellington, S. L. & Vinegar, H. J. X-ray computerized tomography. *J. Petroleum Technol.* **39**, 885–898 (1987).
57. Schindelin, J. *et al.* Fiji: an open-source platform for biological-image analysis. *Nature Methods* **9**, 676–682 (2012).
58. Forster, S., Glud, R. N., Gundersen, J. K. & Huettel, M. *In situ* study of bromide tracer and oxygen flux in coastal sediments. *Estuar. Coast. Shelf Sci.* **49**, 813–827 (1999).
59. Loreau, M. Separating sampling and other effects in biodiversity experiments. *Oikos* **82**, 600–602 (1998).
60. Pinheiro, J. C. & Bates, D. M. *Mixed-effects models in S and S-plus*. New York, NY: Springer (2000).
61. Pinheiro, J. C. *et al.* *nlme: Linear and Nonlinear Mixed Effects Models*. R package version 3.1-111. <http://cran.r-project.org/package=nlme> (2013). (Date of access: 10/09/2013).

## Acknowledgments

We thank J.A. Godbold (University of Southampton) for assistance with setting up the experiments and A.C. Raffan (University of Aberdeen) for bromide analysis. Supported by a University of East Anglia studentship (R.H.) and the University of Southampton (M.S.).

## Author contributions

R.H. and M.S. conducted the experiments, analysed the data and wrote the manuscript. M.N.M. and R.H. performed the computer tomography scanning and image construction, and T.J.T. commented on the manuscript.

## Additional information

**Supplementary information** accompanies this paper at <http://www.nature.com/scientificreports>

**Competing financial interests:** The authors declare no competing financial interests.

**How to cite this article:** Hale, R., Mavrogordato, M.N., Tolhurst, T.J. & Solan, M. Characterizations of how species mediate ecosystem properties require more comprehensive functional effect descriptors. *Sci. Rep.* **4**, 6463; DOI:10.1038/srep06463 (2014).



This work is licensed under a Creative Commons Attribution 4.0 International License. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in the credit line; if the material is not included under the Creative Commons license, users will need to obtain permission from the license holder in order to reproduce the material. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>