Shape shifting predicts ontogenetic changes in metabolic scaling in diverse aquatic invertebrates

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Metabolism fuels all biological activities, and thus understanding its variation is fundamentally important. Much of this variation is related to body size, which is commonly believed to follow a 3/4-power scaling law. However, during ontogeny many kinds of animals and plants show marked shifts in metabolic scaling that deviate from 3/4-power scaling predicted by general models. Here we show that in diverse aquatic invertebrates, ontogenetic shifts in the scaling of routine metabolic rate from near isometry (b_R = scaling exponent ~ 1) to negative allometry (b_R < 1), or the reverse, are associated with significant changes in body shape (indexed by b_L = the scaling exponent for body mass in relation to body length). The observed inverse correlations between b_R and b_L are predicted by metabolic scaling theory that emphasizes resource/waste fluxes across external body surfaces, but contradict theory that emphasizes outward-directed transport of resources through internal networks. Geometric estimates of the scaling of surface area with body mass (b_A) further show that ontogenetic shifts in b_R and b_A are positively

correlated. These results support new metabolic scaling theory based on surface-area influences that may be applied to ontogenetic shifts in b_R shown by many kinds of animals and plants.

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Subject Areas:

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- 39 surface area

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1. Introduction

- 44 All living activities depend on metabolism for energy and materials. Therefore, understanding
- variation in metabolic rate is of fundamental importance in biology. Much of this variation is
- related to body size, but how and why these relationships occur remain vexing questions.
- 47 General models assume that metabolic rate scales monotonically with body size according to the
- 48 simple power function

$$49 R = aM^b, (1.1)$$

- where R is metabolic rate, a is the scaling coefficient (antilog of the intercept in a log-log plot),
- 51 M is body mass, and b (henceforth b_R) is the scaling exponent (linear slope of a log-log plot that
- frequently approximates 3/4) [1, 2]. However, metabolic scaling often shows marked shifts
- during ontogeny in animals and plants (b_R varying mostly between 2/3 and 1, but also showing
- values outside this range) [3-7] that are not well understood. These metabolic shifts are

important because they appear to be fundamentally linked to other ontogenetic changes in the physiology, growth rate, cell size, body composition, behavior and ecology of a species [3-7].

Here we show that ontogenetic changes in body shape and associated surface-area-related resource supply predict frequently observed shifts from near isometric ($b_R \sim 1$) to negatively allometric ($b_R < 1$) intraspecific metabolic scaling (= type III scaling [4]) in diverse aquatic invertebrates. Crucially, we also show that shape shifting predicts more rarely observed changes in metabolic scaling that occur in the opposite direction (from shallow to steep scaling). Our results demonstrate that the prediction of metabolic scaling from the body-shape related scaling of surface area applies more widely than that recently described by Hirst et al. [8]. We show that this predictive power applies to marked variation in metabolic scaling seen not only among diverse pelagic (open-water) animal taxa [8], but also during the intraspecific ontogeny of both pelagic invertebrates and those that exhibit developmental shifts from pelagic to benthic (bottomdwelling) lifestyles. A critical assumption of our shape-shifting model is that the supply of oxygen and (or) nutrients scales with external body surface, thus implying that resource uptake is distributed over the body surface, which we evaluate herein. We also discuss potential implications of our findings for ontogenetic shifts in metabolic scaling observed in many other kinds of animals and plants.

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2. Theoretical background

Metabolic rate may be controlled by the supply of, or demand for, resources [9, 10]. Both should be considered in order to attain a comprehensive understanding of the scaling of metabolic rate with body size [4, 11, 12]. Here we test opposing predictions from influential metabolic scaling

77 theories that focus on transport of resources and waste products between the external environment and metabolizing cells. This transport may be influenced by two major steps: the 78 exchange of materials across body surfaces, and the transport of materials through internal 79 80 networks. These steps are the foci for two prominent theoretical approaches to understanding and predicting biological scaling: surface area (SA) theory [4, 8, 13, 14] and resource-transport 81 network (RTN) theory [1, 2, 15, 16], respectively. 82 Although both surface area and internal transport networks may be important in influencing 83 metabolic rate and its scaling with body size, SA theory predicts that body-shape changes should 84 85 have diametrically opposite effects on the scaling exponent b (see equation 1.1) than those predicted by existing RTN theory [8]. If an organism shows isomorphic growth (i.e., it grows 86 with equal proportions in all three dimensions, so as to maintain a constant shape), SA theory 87 (based on simple Euclidean geometry) predicts that $b_R = 2/3$, whereas RTN theory typically 88 predicts that $b_R = 2/3$ [2, 16] or 3/4 [1, 2, 15], depending on the physical properties of the 89 transport network [17, 18]. These predicted scaling exponents of 2/3 and 3/4 have received the 90 most attention by biologists since the seminal studies of Rubner [13] over a century ago. 91 However, if an organism displays nearly 2D growth (e.g., it grows in length and width without 92 any significant change in depth, thus appearing increasingly flat), SA theory predicts that $b_R \sim 1$ 93 [8, 14], whereas RTN models predict that $b_R \sim 1/2$ [16], 5/8 [2, 8] or 2/3 [15], depending on 94 network geometry and dynamics. In addition, if an organism exhibits nearly 1D growth (e.g., it 95 grows in length without any significant change in width or depth, thus showing an increasingly 96 elongated shape), SA theory predicts that $b_R \sim 1$ [8, 14], whereas RTN theory predicts that $b_R \sim 0$ 97 [16], 1/4 [2, 8] or 1/2 [15], again depending on network properties. Therefore, increased 98 99 elongation or flattening during ontogeny (trends toward 1D or 2D growth) should lead to an

increase in b_R , according to SA theory, whereas RTN theory predicts a decrease in b_R . Conversely, increased thickening during ontogeny (specifically trends away from increasingly elongated 1D or flattened 2D growth, but toward isomorphic 3D growth) reverses the changes in b_R predicted by the two theories: SA theory predicts decreasing, and RTN theory increasing b_R . Studying the effects of ontogenetic shape-shifting on metabolic scaling thus provides an excellent opportunity to test the relative validity of models based on two major competing theories of metabolic scaling, which is much needed for the field to advance [19, 20].

3. Testing theory using animals with mixed ontogenetic metabolic scaling

Several kinds of aquatic animals with complex life cycles exhibit ontogenetic shifts in metabolic scaling, with scaling exponents (b_R) most often changing from near 1 in larvae or young juveniles to < 1 in older juveniles or adults [3, 4, 21], but also more rarely showing reverse shifts [22, 23]. From the literature we collected data on aquatic invertebrates that have complete or partial pelagic (open water) life histories to test whether and how these ontogenetic shifts in metabolic scaling are related to changes in body shape.

Scaling exponents (b_L) of least squares regressions (LSR) of \log_{10} body mass in relation to \log_{10} major body length, separately calculated for larvae, juveniles and adults, were used to quantify differences in shape-related growth between these life-history stages [24]. Logarithmic transformation was used to permit easy detection of proportional changes [25]. If growth is occurring proportionally in three dimensions without any change in mass density, b_L should be 3, whereas if growth involves pure elongation in only one dimension (along the major length axis) or pure flattening because of size increases in only two dimensions, b_L should be 1 or 2, respectively [8]. For intermediate patterns of body-shape change, involving disproportionate

growth in one or two of the longest dimensions, $1 \le b_L \le 3$. Values of b_L may even be > 3, if growth in width and (or) depth is proportionately greater than that for length (i.e., the animal is becoming thicker and/or broader) [8].

Unfortunately, actual measurements of body surface area during ontogeny are rare [8]. Therefore, as a first-order approximation, we used Euclidean geometry for smooth surfaces to deduce scaling exponents (b_A) of \log_{10} body surface area in relation to \log_{10} body mass from b_L values. Values of b_A were inferred from b_L values that are ≤ 3 by using formulae for the extreme possibilities of different degrees of elongation (1D growth) and flattening (2D growth) [8]. The formula for different degrees of elongation is:

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$$b_A = \frac{1}{2} (1 + \frac{1}{b_L}),$$
 (3.1)

whereas the formula for different degrees of flattening is:

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$$b_A = 2(1/b_L)$$
. (3.2)

Equation 3.1 applies for $1 \le b_L \le 3$; and equation 3.2 applies for $2 \le b_L \le 3$. Ranges of potential b_A values were also inferred from b_L values that are > 3 by using formulae quantifying disproportionate thickening in one or two dimensions. The formula for thickening of just the shortest dimension is the same as equation 3.2, whereas the formula for thickening of just the two shortest dimensions is the same as equation 3.1. When data on the scaling of body width with body length were available, it was possible to predict a single value of b_A rather than a range (see Supplementary Information).

The empirical b_L values and inferred b_A values were then compared to scaling exponents (b_R) for regressions of \log_{10} routine metabolic (oxygen consumption) rate (RMR) in relation to \log_{10}

body mass of different ontogenetic stages both within and among several species of aquatic invertebrates.

Both b_L and b_R values were based on LSR, the method used for all of the literature scaling analyses included in our study. An alternative, often used method, reduced major axis (RMA) analysis, gave similar b values ($b_{RMA} = b_{LSR}/r$) to those from LSR, because reported correlation coefficients (r) were always high (≥ 0.8). The r values for b_L averaged 0.97±0.01 (\pm one standard error, n = 15), and those for b_R averaged 0.96±0.01 (n = 19) (calculated from r^2 values in the Supplementary Information for Table 1). As a result, b_L and b_R values based on RMA analyses averaged only ~3-4% higher than those based on LSR.

4. Results

We compared conspecific b_L , b_R and inferred b_A values for different ontogenetic phases of a ctenophore, a scyphozoan, two bivalves, two crustaceans, and two thaliaceans (table 1), as well as mean heterospecific values for the nauplii, copepodites and adults of several copepod crustaceans (table 2).

For seven of the eight species sampled, ontogenetic shifts in the metabolic scaling exponent (b_R) are accompanied by inverse shifts in the scaling exponent of body mass in relation to length (b_L) (table 1, figure 1a). The only exception is the pelagic tunicate *Salpa thompsoni*, which showed no significant difference in b_R or b_L between the solitary (oozoid) and aggregate (blastozoid) life-cycle stages (table 1, figure 1a). Remarkably, even this exception supports a link between b_R and b_L , because both life stages of *S. thompsoni* have relatively high b_R values associated with relatively low b_L values, as compared to the exponents exhibited by the other

species sampled (figure 1a). Furthermore, inverse shifts in b_R and b_L occurred in the other seven species sampled regardless of whether b_R showed an increase or decrease during ontogeny (table 1, figure 1a). Inverse relationships between b_R and b_L , also seen among all of the species averaged together (figures 1b, c), follow SA theory (figure 1b), but contradict all existing RTN models (figure 1c). Although ontogenetic stages with a mean $b_L \sim 3$ (2.97±0.21 95% CI) exhibited a mean b_R (0.72±0.08) not significantly different from 2/3 or 3/4, as predicted by both SA and RTN theory, stages with a mean $b_L \sim 2.3$ (2.30±0.12) exhibited a mean b_R (0.98±0.07) not significantly different from 1 and significantly greater than 2/3 and 3/4, as predicted by SA theory, but in contradiction to all RTN models, which predict b_R values < 2/3 or 3/4 (figures 1b, c).

Since the scaling exponent for body surface area (b_A), as inferred from Euclidean geometry, is inversely related to b_L (figure 1a; and equations 3.1 and 3.2), it follows that ontogenetic shifts in b_R and b_A should be positively correlated, as observed in seven of the eight species sampled (table 1, figure 2a). As predicted by SA theory, b_R and b_A are also positively correlated for all species averaged together (figure 2b), and when these exponents were compared pairwise among each of the ontogenetic phases of each species (figure 2c). The slope for the latter correlation (1.14) is not significantly different from 1, as expected if b_R varied in direct proportion to b_A (figure 2c).

Similar ontogenetic shifts are seen when b_R , b_L and b_A values are compared between nauplii and copepodites/adults averaged among several species of copepods (table 2, figure 1a). The heterospecific, inversely related shifts in b_R and b_L almost exactly parallel those observed for the single copepod species *Mesocyclops brasilianus* (figure 1a).

The spiny lobster Sagmariasus verreauxi nicely exemplifies how ontogenetic changes in metabolic rate and body shape correlate. The phyllosoma larvae are very thin and flat and show nearly 2D growth, until they metamorphose into adult-looking benthic juveniles that are much thicker and show 3D growth. This marked shift in growth pattern and body form is represented by an abrupt ontogenetic shift in b_L values: from 2.142 \pm 0.260 (95% CI) in the phyllosoma larvae to 2.991 \pm 0.037 in the juveniles (figure 3a, table 1). The phyllosoma b_L value is not significantly different from 2, whereas the juvenile b_L value is not significantly different from 3. As a result, the scaling exponents for surface area (b_A) can be inferred (see Supplementary Information) to be 0.91 for the phyllosomas and 0.67 for the juveniles (table 1). Like b_A , the b_R values for RMR are also higher in the phyllosomas (1.002 \pm 0.081) than in the juveniles (0.829 \pm 0.157) (figure 3b, table 1). Comparisons of the 95% CI [19, 28] reveal that the b_L and b_R values are both significantly different between the two life-history stages. The phyllosoma b_R value is not significantly different from 1, but is significantly greater than 1/2, 5/8, 2/3 and 3/4. By contrast, the juvenile b_R value is significantly less than 1, not significantly different from 3/4, and significantly greater than 1/2, 5/8 and 2/3 (just barely).

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5. Discussion

The parallel changes in the scaling exponents for surface area (SA) and routine metabolic rate (RMR) observed in seven of the eight species sampled supports the importance of SA changes in the observed ontogenetic shifts in metabolic scaling, but contradicts predictions of all current models emphasizing internal resource-transport networks. SA theory predicts positive correlations between the scaling exponents for SA (b_A) and RMR (b_R), as observed, whereas current RTN theory incorrectly predicts negative correlations between b_A and b_R . Although RTN

212 macroscopic multicellular eukaryotes [29], RTN theory may not apply to animals without circulatory systems or with open or incompletely closed circulatory systems [2, 4, 18], which are 213 214 far more common taxonomically than those with completely closed circulatory systems (including only vertebrates, cephalopod mollusks, and some annelid and nemertean worms [30, 215 31]. In fact, two of the animal species included in our study have no or very rudimentary 216 circulatory systems (a ctenophore and scyphozoan), whereas the other six have open or 217 incompletely closed circulatory systems (the bivalve, crustacean and thaliacean species) [31]. 218 219 Nevertheless, Hirst et al. [8] have shown that the similarly steep ontogenetic scaling of metabolic rate and surface area in squids (cephalopods), which have closed circulatory systems, is also 220 consistent with SA theory, but not with RTN theory. This additional deviation from RTN theory 221 222 may be because squids use both their skin and gills for gas exchange and also distribute resource-223 laden blood by means of multiple hearts found throughout the body [31], rather than from a single centralized heart or distribution center, as assumed by the theory [1, 2, 15, 16]. If current 224 225 RTN theory does not apply to squids for these reasons, then annelid worms that use multiple hearts to pump blood [31] and nemertean worms that have no heart at all [31] may also be 226 exceptions. Therefore, current RTN theory appears to apply to only a small subset of animals: 227 perhaps only vertebrates with single central hearts in a completely closed vascular system. Even 228 the application of RTN theory to vertebrates may be of limited use, because currently it cannot 229 explain large variation in the metabolic scaling exponent observed in various vertebrate classes 230 that appears to be related to physiological state, ecological lifestyle, or environmental conditions 231 [11, 32-35]. Perhaps next generation resource exchange and supply theory may resolve these 232 233 problems (also see section 6).

theory has been claimed to be universally applicable to all of life [1, 15] or at least to

In any case, the positive associations between b_A and b_R observed in this study are remarkable because they occur in animals with very different body designs from five different phyla, and furthermore, they occur regardless of whether b_A increases, or decreases, during ontogeny. Therefore, it is unlikely that these correlations are merely side-effects of b_A and b_R being independently related to other unmeasured factors associated with developmental maturation or ontogenetic age. Rather, it is more likely that shape shifting directly affects metabolic scaling via changes in surface area available for resource uptake and waste excretion.

This hypothetical mechanism for how shape-shifting may affect metabolic scaling seems especially applicable to the thin-skinned pelagic animals (larval or adult) that we have studied here and in a related paper [8]. Phyllosoma, veliger and other pelagic larvae of many marine animals that show biphasic or triphasic metabolic scaling appear to be 'skin-breathers', i.e., they can absorb oxygen and expel dissolved waste products such as CO₂ through their thin, permeable integuments [36, 37]. Some can even absorb nutrients through their body surfaces and metabolize them in their tissues [36, 38]. Remarkably, the body-mass scaling slope for metabolism in larvae of the Pacific oyster, *Crassostrea gigas*, is not significantly different from that for uptake of the amino acid alanine [36]. However, it is not known how much cutaneous absorption of nutrients contributes to the overall energy budget of these larvae and other pelagic animals under natural conditions [39]. Our findings point toward the importance of further research on integumentary energy and material exchange by aquatic animals, as a way to better understand variation in their metabolic scaling.

Although parallel shifts in b_A and b_R values have been typically seen in this study, possible exceptions invite further scrutiny. For example, although b_A and b_R decrease in tandem in the common mussel *Mytilus edulis*, as juveniles mature into adults, more research is needed to

257 determine whether shifts in SA and metabolic scaling also match as veliger larvae develop into juveniles. This is because, although the inferred b_A shift from 0.57 to 0.71-0.83 parallels the shift 258 of b_R from 0.77 (based on an average of four values) to 0.89, the individual b_R estimates for 259 260 veligers are highly variable, ranging from 0.59 to 0.90 (table 1). Some of this variation appears to be related to temperature, because as temperature increases from 6 to 18° C, b_R decreases from 261 0.90 to 0.59 (see Supplementary Information), as predicted by the metabolic-level boundaries 262 hypothesis [11]. Sampling error may also be important because the body-size range of growing 263 veligers is small (< 2 orders of magnitude), thus potentially increasing variation in b_R estimates 264 [32, 40, 41] relative to those of juveniles and adults with larger body-size ranges (> 3 and 4 265 orders of magnitude, respectively; see table 1). A similar explanation may apply to the variable 266 b_R estimates (0.35 and 1.01) for two ephyra samples of the scyphozoan Aurelia aurita that have 267 268 body-size ranges < 1.5 orders of magnitude, and that deviate markedly from the inferred b_A value of 0.64 (table 1). By contrast, the b_R estimate (0.63) for a third sample of ephyra larvae, with a 269 body-size range ≥ 2 orders of magnitude, is very close to the estimated b_A value. The b_R value 270 271 (0.65) for ephyra larvae of A. aurita, estimated from the metabolic data of several studies taken together, is also not significantly different from 0.64 [42]. It is also possible that our simple 272 Euclidean estimates of SA scaling (b_A) do not adequately represent the ontogenetic SA changes 273 occurring in a mussel veliger or an ephyra larva. A veliger's velum (foot), which has an 274 extensive, highly permeable surface [36], can extend far beyond the measured shell length used 275 in calculating b_L and b_A [43]. In addition, the ephyra larva shows complex changes in body 276 shape as it grows, thus making it difficult to accurately estimate the scaling of its surface area 277 (see footnote 6 in Supplementary Information for table 1). 278

Another major pattern evident in our results is that, although b_R is significantly correlated with b_A , it is usually greater than that predicted by b_A alone (see tables 1 and 2, and figures 1a, b and 2). Two major factors may help account for these upward deviations of b_R . First, metabolically costly growth may elevate b_R values, as has been observed in other animals and plants [3-7, 19]. These growth effects prompt the question: what are the relative influences of resource supply versus metabolic demand by growth and other biological processes on ontogenetic metabolic scaling [4, 9-12]? For example, does the steep scaling of SA (and presumably resource uptake) of many kinds of pelagic animals (including larvae) documented here and by Hirst et al. [8] permit or even drive the steep scaling of metabolism, or is steep SA scaling a secondary adjustment to steep metabolic scaling that is driven by high resource demand (e.g., high mass-specific growth rates that often occur in pelagic animals [4, 44, 45]). Attempting to answer this critical question brings into focus the importance of understanding the various factors influencing all of the steps of energy flow through an organism, and their integration, as a way to improve our knowledge of how and why metabolic rate varies with body size (also see [4]).

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A second major factor that may help to explain apparent upward deviations in b_R from that predicted by our SA model (figure 1a) is that we may have underestimated SA changes during ontogeny in our study species. Our geometric estimates of b_A based on b_L do not allow the detection of increases in SA related to increased convolutions (or fractal dimensionality) of the body surface, or to the development of special respiratory and nutrient-absorption structures (e.g., the velum in veligers, and the gills in other larval or juvenile forms). In addition, half of our inferred b_A values (see Table 1) may have been underestimated because they were based on the midpoint of a potential range of values, whereas in most cases the upper limit of this range,

which involves shape flattening, is probably closer to the actual b_A value than the lower limit, which involves shape elongation (also see section 3 and [8]). This claim is supported by two observations. First, when data on both b_L and the scaling of body width versus length were available (Supplementary Information), the inferred b_A value was almost always closer (in 9 of 10 cases) to the predicted upper limit involving mainly 2D growth (flattening) than the lower limit involving mainly 1D growth (elongation) (see table 1). Second, the mean empirical relationship between b_L and b_R (and by association b_A : see figures 2a, b, c) for all of our study species more closely parallels the upper predicted boundary involving flattening than the lower boundary involving elongation (figure 1b).

Other data consistent with effects of ontogenetic shifts in body shape and SA scaling on metabolic scaling include significant correlations between b_L and b_R values among diverse 'skinbreathing' pelagic animals, for which species-specific values also show a closer match to the upper boundary curve for predicted b_A than to the lower boundary curve [8], and parallel steep scaling of SA and metabolic rate in echinoid larvae [46] and epipelagic squid [47]. Steep (often near isometric) scaling of metabolic rate in larval fishes may not only be related to their rapid growth rates [4], but also to their ability to absorb oxygen and possibly nutrients through their entire body surface [48, 49]. In addition, shifts from isometric to negatively allometric metabolic scaling seen in plants may be related, at least in part, to decreases in their relative 'leafiness' (i.e., reductions in the relative contribution of high SA leaves to total biomass) as they grow [6].

6. Conclusions

Ontogenetic shifts in metabolic scaling may often be linked to developmental changes in body shape and SA scaling. Our SA model predicts that this should happen in any organism that relies heavily on body surfaces for resource uptake, such as many kinds of thin-skinned pelagic animals and leafy plants. Ontogenetic shifts in metabolic scaling in other organisms with impermeable integuments may be related to shifts in the SA scaling of specialized respiratory structures (as occur for gills in developing fish [48] and for tracheae during molting in insect larvae [12]) or to other metabolically demanding developmental changes unrelated to body shape (e.g., shifts from ectothermy to endothermy in developing mammals [4]). Nevertheless, evidence reported here and elsewhere suggests that the effect of external surface area on metabolic scaling may often outweigh or modify the influence of resource-transport networks [4, 8, 11, 46, 47], contrary to currently influential general models [1, 2]. New resource transport theory is needed that can accommodate organisms that distribute resources from external surfaces rather than from (or in addition to) internal central "pumps" (also see [8]). Therefore, reports of the demise of surface-area theory in the field of metabolic scaling are incorrect [50, 51]. We argue that surface-area theory is alive and strong, and is essential to a comprehensive explanation of metabolic scaling.

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References

- West GB, Brown JH, Enquist BJ. 1997. A general model for the origin of allometric scaling laws in biology. Science 276, 122-126. (doi:10.1126/science.276.5309.122)
- Banavar JR, Moses ME, Brown JH, Damuth J, Rinaldo A, Sibly RM, Maritan A. 2010. A general basis for quarter-power scaling in animals. *Proc. Natl. Acad. Sci. U.S.A.* 107, 15816-15820.
 (doi:10.1073/pnas.1009974107)
- 3. Riisgård HU. 1998 No foundation of a '3/4 power scaling law' for respiration in biology. *Ecol. Lett.* **1**, 71-73. (doi:10.1046/j.1461-0248.1998.00020.x)
- 4. Glazier DS. 2005 Beyond the '3/4-power law': variation in the intra- and interspecific scaling of metabolic rate in animals. *Biol. Rev.* **80**, 611-662. (doi:10.1017/S1464793105006834)

- 351 5. Czarnołeski M, Kozłowski J, Dumiot G, Bonnet J-C, Mallard J, DuPont-Nivet M. 2008 Scaling of
- metabolism in *Helix aspersa* snails: changes through ontogeny and response to selection for
- increased size. *J. Exp. Biol.* **211**, 391-399. (doi:10.1242/jeb.013169)
- 354 6. Peng Y, Niklas KJ, Reich PB, Sun S. 2010 Ontogenetic shift in the scaling of dark respiration with
- 355 whole-plant mass in seven shrub species. Funct. Ecol. 24. 502-512. (doi:10.1111/j.1365-
- 356 2435.2009.01667.x)
- 7. Kutschera U, Niklas KJ. 2011 Ontogenetic changes in the scaling of cellular respiration with respect to
- size among sunflower seedlings. *Plant Signal. Behav.* **6**, 72-76. (doi:10.4161/psb.6.1.14001)
- 8. Hirst AG, Glazier DS, Atkinson D. 2014 Body shape shifting during growth permits tests that
- distinguish between competing geometric theories of metabolic scaling. Ecol. Lett.
- 361 (doi:10.1111/ele.12334).
- 9. Darveau C-A, Suarez RK, Andrews RD, Hochachka PW. 2002 Allometric cascade as a unifying
- principle of body mass effects on metabolism. *Nature* **417**, 166-170. (doi:10.1038/417166a)
- 10. Hofmeyr JHS, Rohwer JM. 2011. Supply-demand analysis: a framework for exploring the regulatory
- design of metabolism. *Methods Enzymol.* **500**, 533-554. (doi: 10.1016/B978-0-12-385118-5.00025-6)
- 11. Glazier DS. 2010 A unifying explanation for diverse metabolic scaling in animals and plants. *Biol. Rev.*
- **85**, 111-138. (doi:10.1111/j.1469-185X.2009.00095.x)
- 12. Callier V, Nijhout HF. 2012 Supply-side constraints are insufficient to explain the ontogenetic scaling
- of metabolic rate in the Tobacco Hornworm, *Manduca sexta. PLoS ONE* **7**, e45455.
- 370 (doi:10.1371/journal.pone.0045455)
- 13. Rubner M. 1883 Über den Einfluss der Körpergrösse auf Stoff- und Kraftwechsel. Zeit. Biol. 19, 535-
- 372 562.
- 373 14. Okie JG. 2013 General models for the spectra of surface area scaling strategies of cells and
- 374 organisms: fractality, geometric dissimilitude, and internalization. Am. Nat. **181**, 421-439.
- 375 (doi:10.1086/669150)
- 15. West GB, Brown JH, Enquist BJ. 1999 The fourth dimension of life: fractal geometry and allometric
- 377 scaling of organisms. Science 284, 1677-1679. (doi:10.1126/science.284.5420.1677)
- 16. Dodds PS. 2010 Optimal form of branching supply and collection networks. *Phys. Rev. Lett.* **104**,
- 379 048702. (doi:10.1103/PhysRevLett.104.048702)
- 17. Savage VM, Deeds EJ, Fontana W. 2008. Sizing up allometric scaling theory. *PLoS Comp. Biol.* **4**,
- 381 e1000171 (doi:10.1371/journal.pcbi.1000171)
- 18. Price CA, Weitz JS, Savage VM, Stegen J, Clarke A, Coomes DA, Dodds PS, Etienne RS, Kerkhoff
- AJ, McCulloh K, Niklas KJ, Olff H, Swenson NG. 2012 Testing the metabolic theory of ecology. *Ecol.*
- 384 Lett. **15**, 1465-1474. (doi:10.1111/j.1461-0248.2012.01860.x)
- 19. Glazier DS, Butler EM, Lombardi SA, Deptola TJ, Reese AJ, Satterthwaite EV. 2011 Ecological
- effects on metabolic scaling: amphipod responses to fish predators in freshwater springs. *Ecol.*
- 387 *Monogr.* **81**, 599-618. (doi.org/10.1890/11-0264.1)

- 388 20. Kearney MR, White CR 2012 Testing metabolic theories. Am. Nat. 180, 546-565.
- 389 (doi:10.1086/667860)
- 21. Zeuthen E. 1953 Oxygen uptake as related to body size in organisms. Q. Rev. Biol. 28, 1-12.
- 391 (http://www.jstor.org/stable/2810299)
- 392 22. Kinoshita J, Hiromi J, Kadota S. 1997 Do respiratory metabolic rates of the scyphomedusa *Aurelia*
- 393 aurita scale isometrically throughout ontogeny in a sexual generation? Hydrobiologia 347, 51-55.
- 394 (doi:10.1023/A:1002942806113)
- 395 23. Svetlichny LS, Abolmasova GI, Hubareva ES, Finenko GA, Bat, L, Kideys AE. 2004 Respiration rates
- of *Beroe ovata* in the Black Sea. *Mar. Biol.* **145**, 585-593. (doi:10.1007/s00227-004-1336-4)
- 397 24. Hirst AG. 2012 Intra-specific scaling of mass to length in pelagic animals: Ontogenetic shape change
- and its implications. *Limnol. Oceanogr.* **57**, 1579-1590. (doi:10.4319/lo.2012.57.5.1579)
- 399 25. Glazier DS. 2013 Log-transformation is useful for examining proportional relationships in allometric
- 400 scaling. *J. Theor. Biol.* **334**, 200-203. (doi.org/10.1016/j.jtbi.2013.06.017)
- 26. Jensen MA, Fitzgibbon QP, Carter CG, Adams LR. 2013a Effect of body mass and activity on the
- metabolic rate and ammonia-N excretion of the spiny lobster *Sagmariasus verreauxi* during ontogeny.
- 403 Comp. Biochem. Physiol., Part A 166, 191-198. (doi.org/10.1016/j.cbpa.2013.06.003)
- 404 27. Jensen MA, Fitzgibbon QP, Carter CG, Adams LR. 2013b The effect of stocking density on growth,
- 405 metabolism and ammonia-N excretion during larval ontogeny of the spiny lobster Sagmariasus
- 406 *verreauxi. Aquaculture* **376-379**, 45-53. (doi.org/10.1016/j.aquaculture.2012.10.033)
- 407 28. Cumming G. 2008 Inference by eye: reading the overlap of independent confidence intervals. Stat.
- 408 *Med.* **28**, 205-220. (doi:10.1002/sim.3471)
- 409 29. DeLong JP, Okie JG, Moses ME, Sibly RM, Brown JH. 2010 Shifts in metabolic scaling, production
- 410 and efficiency across major evolutionary transitions of life. *Proc. Natl. Acad. Sci. U.S.A.* **107**, 12941-
- 411 12945. (doi:10.1073/pnas.1007783107)
- 412 30. Reiber CL, McGaw IJ. 2009 A review of the "open" and "closed" circulatory systems: new terminology
- 413 for complex invertebrate circulatory systems in light of current findings. *Int. J. Zool.* **2009** (Article ID
- 414 301284), 1-8. (doi:10.1155/2009/301284)
- 31. McMahon BR, Wilkens JL, Smith PJS. 2011 Invertebrate circulatory systems. Compr. Physiol., 931–
- 416 1008. (doi:10.1002/cphy.cp130213)
- 417 32. Bokma F. 2004 Evidence against universal metabolic allometry. *Funct. Ecol.* **18**, 184-187.
- 418 (doi:10.1111/j.0269-8463.2004.00817.x)
- 419 33. White CR, Phillips NF, Seymour RS. 2006 The scaling and temperature dependence of vertebrate
- 420 metabolism. *Biol. Lett.* **2**, 125-127. (doi:10.1098/rsbl.2005.0378)
- 421 34. Glazier DS. 2008 Effects of metabolic level on the body-size scaling of metabolic rate in birds and
- 422 mammals. *Proc. Roy. Soc. Lond B* **275**, 1405-1410. (doi:10.1098/rspb.2008.0118)

- 423 35. Killen SS, Atkinson D, Glazier DS. 2010 The intraspecific scaling of metabolic rate with body mass in
- fishes depends on lifestyle and temperature. Ecol. Lett. 13, 184-193. (doi:10.1111/j.1461-
- 425 0248.2009.01415.x)
- 426 36. Manahan DT. 1990 Adaptations by invertebrate larvae for nutrient acquisition from seawater. Am.
- 427 Zool. **30**, 147-160. (doi:10.1093/icb/30.1.147)
- 428 37. Rodriguez Souza JC, Strüssmann CA, Takashima F, Satoh H, Sekine S, Shima Y, Matsuda H. 2010
- Oral and integumental uptake of free exogenous glycine by the Japanese spiny lobster *Panulirus*
- 430 *japonicas* phyllosoma larvae. *J. Exp. Biol.* **213**, 1859-1867. (doi: 10.1242/jeb.040030)
- 431 38. Haond C, Charmantier G, Flik G, Bonga SE. 2001 Identification of respiratory and ion-transporting
- epithelia in the phyllosoma larvae of the slipper lobster Scyllarus arctus. Cell Tissue Res. 305, 445-
- 433 455. (doi:10.1007/s004410100405)
- 434 39. Gomme, J. 2001 Transport of exogenous organic substances by invertebrate integuments: the field
- 435 revisited." J. Exp. Zool. 289, 254-265. (doi:10.1002/1097-010X(20010401/30)289:4<254::AID-
- 436 JEZ6>3.0.CO;2-F)
- 437 40. White CR, Seymour RS. 2005 Sample size and mass range effects on the allometric exponent of
- 438 basal metabolic rate. *Comp. Biochem. Physiol. A* **142**, 74-78. (doi.org/10.1016/j.cbpa.2005.07.013)
- 439 41. Moses ME, Hou C, Woodruff WH, West GB, Nekola JC, Zuo W, Brown JH. 2008 Revisiting a model of
- ontogenetic growth: estimating model parameters from theory and data. Am. Nat. 171, 632-645.
- 441 (http://www.jstor.org/stable/10.1086/587073)
- 42. Gambill M, Peck MA. 2014 Respiration rates of the polyps of four jellyfish species: Potential thermal
- triggers and limits. *J. Exp. Mar. Biol. Ecol.* **459**, 17-22. (doi:10.1016/j.jembe.2014.05.005)
- 43. Hamburger K, Møhlenberg F, Randløv A, Riisgård HU. 1983 Size, oxygen consumption and growth in
- the mussel *Mytilus edulis. Mar. Biol.* **75**, 303-306. (doi:10.1007/BF00406016)
- 446 44. Glazier DS. 2006 The 3/4-power law is not universal: evolution of isometric, ontogenetic metabolic
- scaling in pelagic animals. *Bioscience* **56**, 325-332. (doi:10.1641/0006-
- 448 3568(2006)56[325:TPLINU]2.0.CO;2)
- 449 45. Hirst AG, Forster J. 2013 When growth models are not universal: evidence from marine invertebrates.
- 450 *Proc. R. Soc. B* 280: 20131546. (doi.org/10.1098/rspb.2013.1546)
- 451 46. McEdward LR. 1984 Morphometric and metabolic analysis of the growth and form of an
- 452 echinopluteus. J. Exp. Mar. Biol. Ecol. 82, 259-287. (doi.org/10.1016/0022-0981(84)90109-6)
- 47. Seibel BA. 2007 On the depth and scale of metabolic rate variation: scaling of oxygen consumption
- rates and enzymatic activity in the Class Cephalopoda (Mollusca). J. Exp. Biol. 210, 1-11.
- 455 (doi:10.1242/jeb.02588)
- 48. Post JR, Lee JA. 1996 Metabolic ontogeny of teleost fishes. Can. J. Fish. Aquat. Sci. 53, 910-923.
- 457 (doi:10.1139/f95-278)
- 458 49. Glover CN, Bucking C, Wood, CM. 2013 The skin of fish as a transport epithelium: a review. J. Comp.
- 459 *Physiol. B* 183, 877-891. (doi: 10.1007/s00360-013-0761-4)

Press. 51. Savage, VM, Gillooly JF, Woodruff WH, West GB, Allen AP, Enquist BJ, Brown JH. 2004 The predominance of quarter-power scaling in biology. Funct. Ecol. 18, 257-282. (doi:10.1111/j.0269-8463.2004.00856.x)

50. Peters RH. 1983 The ecological implications of body size. Cambridge, UK: Cambridge University

Table 1. Ontogenetic scaling exponents from least squares regressions of \log_{10} body mass in relation to \log_{10} body length (b_L) and \log_{10} routine metabolic rate in relation to \log_{10} body mass (b_R) for larvae, juveniles, and (or) adults of the Atlantic ctenophore $Beroe\ ovata$, moon jellyfish $Aurelia\ aurita$, Pacific oyster $Crassostrea\ gigas$, common mussel $Mytilus\ edulis\ and\ spiny\ lobster\ Sagmariasus\ verreauxi$; for nauplii, copepodites and adults of the copepod $Mesocyclops\ brasilianus$; and for solitary and aggregate life-cycle stages of the salps (pelagic tunicates) $Salpa\ fusiformis\ and\ S.\ thompsoni$. Values of b_L and b_R were taken or calculated from data in sources listed in the Supplementary Information, where additional data and methodological information can be found. Values of b_A for \log_{10} body surface area in relation to \log_{10} body mass were estimated from b_L values and scaling exponents of body width versus length (Supplementary Information). When data for scaling of width versus length were not available, ranges of potential b_A values are given (based on equations 3.1 and 3.2). Note the parallel changes in b_A and b_R values (shown in bold), regardless of whether decreases or increases in b_R were observed during ontogeny (except possibly for the transition from veliger larvae to juveniles in M. edulis; also see text). Statistically significant ontogenetic shifts in b_L and b_R values occur in all species, except for between solitary and aggregate life stages of S. thompsoni.

Species	Stage	b_L	b_A	b_R
Beroe ovata	juveniles	2.92	0.67 (0.67-0.68)	0.62
Ovata	adults	2.47	0.80 (0.70-0.81)	0.99
	addito	2.23	0.87 (0.72-0.90)	1.04
		1.78	1.09 (0.78-1.00)	0.86
 Aurelia	ephyra larvae	3.14	0.64 (0.64-0.66)	0.63
aurita				0.35
				1.01
	medusae	2.50	0.80 (0.70-0.80)	0.93
		2.72	0.74 (0.68-0.74)	1.11
				1.01
Crassostrea gigas	veliger larvae	2.12	0.95 (0.74-0.94)	0.96
	adults	2.79	0.68-0.72	0.77
Mytilus	veliger larvae	3.49	0.57 (0.57-0.71)	0.90
edulis				0.90
				0.70
				0.59
	juveniles	2.42	0.71-0.83	0.89

541 542 543		adults	3.17	0.63-0.68	0.66 0.68
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546 547	Sagmariasus verreauxi	phyllosoma larvae	2.14	0.91 (0.73-0.93)	1.00
547 548 549		juveniles	2.99	0.67	0.83
550 551 552 553	Mesocyclops brasilianus	nauplii	2.15	0.73-0.93	1.08
		copepodites & adults	3.12	0.64-0.68	0.56
554	Salpa fusiformis	solitary zooids	2.40	0.71-0.83	1.15
		aggregate zooids	2.78	0.68-0.72	0.68
	Salpa thompsoni	solitary zooids	2.28	0.72-0.88	0.84
560 561	·	aggregate zooids	2.41	0.71-0.83	0.92
562					
563					
564					

Table 2. Mean ontogenetic scaling exponents from least squares regressions of \log_{10} body mass in relation to \log_{10} body length (b_L) and \log_{10} routine metabolic rate in relation to \log_{10} body mass (b_R) for nauplii, copepodites and adults of several copepod species (b_L and b_R values were taken from the indicated sources, whereas the ranges of potential b_A values for \log_{10} body surface area in relation to \log_{10} body mass were estimated from b_L values using equations 3.1 and 3.2 in the text). Mean scaling exponents were calculated by averaging mean conspecific values among species. The 95% confidence intervals (CI) and number of species sampled (n) are given for each scaling exponent. Note the parallel changes in b_A and b_R values (shown in bold). Data sources are given in the Supplemental Information.

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Stage	b _L ±95% CI (n)	b _A	<i>b</i> _R ±95% CI (n)
Nauplii	2.28±0.27 ¹ (17)	0.72-0.88	1.00 ±0.16 ² (4)
Copepodites and adults	2.84±0.20 ¹ (20)	0.68-0.70	0.72 ±0.22 ² (7)

¹ Based on dry mass, carbon mass or nitrogen mass.

² Based on dry body mass

Figure legends

Figure 1. Ontogenetic shifts of b_R in relation to b_L of several species of aquatic invertebrates in five different phyla (a) and for all species averaged together (b, c), where b_R is the scaling exponent for \log_{10} routine metabolic rate in relation to \log_{10} body mass, and b_L is the scaling exponent for \log_{10} body mass in relation to \log_{10} total body length (data from tables 1 and 2). Multiple b values for the life-history stage of a species or group of species (copepods) were averaged. Arrows indicate the direction of ontogenetic change. In (b, c) the upper left and lower right points were calculated by averaging all of the paired b_R and b_L values that occurred in ontogenetic stages with the higher versus lower b_R , respectively. The 95% confidence limits are shown for each mean value of b_R and b_L . Also shown in (b) are the bounded range of values of b_A (\log_{10} body surface area in relation to \log_{10} body mass) in relation to b_L (depicted as light purple lines) calculated using equations 3.1 and 3.2 based on Euclidean geometry (also see [8]); and in (c) the predicted effects of body-shape changes (b_L) on b_R (depicted as dashed colored lines) according to the resource-transport network models of West et al. (1999)[15], Banavar et al. (2010)[2] and Dodds (2010)[16] (also see [8]). Note that in 7 of 8 species sampled, ontogenetic shifts in b_L are accompanied by inverse shifts in b_R , as predicted by surface-area scaling theory (b), but in contradiction to resource-transport network scaling theory (c).

Figure 2. Ontogenetic shifts of b_R in relation to b_A of several species of aquatic invertebrates (a) and for all species averaged together (b), where b_R is the scaling exponent for \log_{10} routine metabolic rate in relation to \log_{10} body mass, and b_A is the scaling exponent for \log_{10} body mass in relation to \log_{10} total body length (data from table 1). Values of b_A were calculated using b_L values and additional data in the Supplementary Information. Multiple b_R values for the life-history stage of a species were averaged, whereas the midpoint was used when only a range of b_A values was available. In (b) the lower left and upper right points were calculated by averaging all of the paired b_R and b_A values that occurred in ontogenetic stages with the lower versus higher b_R , respectively. The 95% confidence limits are shown for each mean value of b_R and b_A . As predicted by surface-area scaling theory (also see [8]), in 7 of 8 species sampled, ontogenetic shifts in b_A are accompanied by positively correlated shifts in b_R . Also shown in (c) is a significant positive correlation between b_R and b_A for each ontogenetic stage of each species. The equation for the least squares regression line is: $b_R = 1.14 \pm 0.64$ (95% CI) (b_A) – 0.0092 ($r^2 = 0.454$; P = 0.0018, n = 17). In (a, b, c), the dotted diagonal lines represent $b_R = b_A$.

Figure 3. Scaling of \log_{10} wet body mass in relation to \log_{10} total body length (*a*) and \log_{10} routine metabolic rate (RMR) in relation to \log_{10} wet body mass (*b*) in phyllosoma larvae (open points) and juveniles (solid points) of the spiny lobster *Sagmariasus verreauxi* maintained at 21-23° C. Each point in (*a*) and (*b*) is based on 4-11 replicate measurements, respectively [26, 27]. The phyllosoma points

657 represent instars 1, 3, 6, 9, 12, 15 and 17. The standard errors for each RMR value are all less than 20% 658 of the mean. The least squares regression equations for the scaling lines and their coefficients of determination (r^2) and significance levels (p) for phyllosoma larvae and juveniles are: (a) Y = -3.931 + 659 2.142 (X), $r^2 = 0.988$, p = 0.00001; Y = -4.574 + 2.991 (X), $r^2 = 1.000$, p < 0.00001; and (b) Y = -0.773 + 660 1.002 (X), $r^2 = 0.995$, p < 0.00001; Y = -0.673 + 0.829 (X), $r^2 = 0.993$, p = 0.00353, respectively. The 661 number by each line is the scaling slope (exponent). Linear extrapolations of the empirical scaling lines 662 663 for the phyllosoma larvae are shown as dotted lines to highlight the ontogenetic scaling shifts for both 664 body shape (a) and metabolic rate (b).

— Beroe ovata (larvae - adults) CTENOPHORA

— Aurelia aurita (ephyrae - medusae) CNIDARIA

— Crassostrea gigas (larvae - adults) MOLLUSCA

— Mytilus edulis (larvae - juveniles - adults) MOLLUSCA

— Sagmariasus verreauxi (larvae - juveniles) ARTHROPODA

— Mesocyclops brasilianus (nauplii - copepodites & adults) ARTHROPODA

— Means of copepod species (nauplii - copepodites & adults) ARTHROPODA

— Salpa fusiformis (solitary - aggregate) CHORDATA

— Salpa thompsoni (solitary - aggregate) CHORDATA







