

Energetic basis of colonial living in social insects

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Understanding the ecology and evolution of insect societies requires greater knowledge of how sociality affects the performance of whole colonies. Metabolic scaling theory, based largely on the body mass scaling of metabolic rate, has successfully predicted many aspects of the physiology and life history of individual (or unitary) organisms. Here we show, using a diverse set of social insect species, that this same theory predicts the size dependence of basic features of the physiology (i.e., metabolic rate, reproductive allocation) and life history (i.e., survival, growth, and reproduction) of whole colonies. The similarity in the size dependence of these features in unitary organisms and whole colonies points to commonalities in functional organization. Thus, it raises an important question of how such evolutionary convergence could arise through the process of natural selection.

allometry | colony | scaling | metabolism | metabolic theory of ecology

Multicellularity and sociality represent two of life's major evolutionary innovations (1). Both are examples of how individual modules—cells and individuals—can cooperate to enhance evolutionary fitness. In the case of multicellularity, the emergence of such cooperation is relatively easy to explain in the context of natural selection given that all cells are governed by a single, shared genome. Sociality in general, and altruism in particular, is more difficult to explain in this context given that social groups often consist of multiple genotypes with varying degrees of genetic relatedness. In particular, understanding the cooperation observed in eusocial insects—ants, bees, wasps, and termites whose workers forgo reproduction to care for the young of their queens—has presented a paradox given the potential for genetic conflicts to cooperation. However, combining Hamilton's concept of inclusive fitness (2) with the genetics of haplodiploidy has gone some way toward resolving this apparent paradox in the evolution of eusociality (3).

However, simply because cooperation among multicellular individuals or members of eusocial colonies can arise through natural selection does not mean it will endure in nature. For this, we must better understand how cooperation affects the collective performance of those in the group. One key metric of performance is an organism's ability to harvest, store, and transform energy to produce offspring (4). As Boltzmann (1905, cited in ref. 5, p. 6) pointed out, "[The] struggle for existence is a struggle for free energy available for work," and Lotka (6) wrote, "In the struggle for existence, the advantage must go to those organisms whose energy-capturing devices are most efficient in directing available energy into channels favorable to the preservation of the species." For multicellular organisms, metabolic scaling theory has helped to quantify how changes in body size affect the energy use of species (7, 8). The theory and empirical work on this subject have shown that there are economies of scale related to energy use such that cells in larger, more complex animals require less energy per capita. For eusocial colonies, it has long been posited that these "superorganisms" experience similar relationships with colony size (3, 9–12), perhaps owing to shared constraints on the delivery of energy and materials (e.g., branching distribution networks, space-filling surface area to volume constraints) (e.g., see refs. 13–16). But empirical evidence for these relationships is scarce (but see refs. 17 and 18). This hypothesis deserves further

attention because, if unitary organisms and eusocial colonies show the same size-dependent allometries with respect to energy use, this may suggest that selection acts on colonies much as it acts on individuals (9).

Thus, here we assess how basic attributes of the physiology and life history of colonies vary as a function of whole-colony mass and then compare these findings to the equivalent relationships in unitary organisms. In doing so, we quantitatively compare the functional organization of colonies and unitary organisms. Specifically, we evaluate the hypothesis that the 1/4 power scaling of metabolic rate (a measure of energy flux) and associated life history traits—successfully predicted from metabolic scaling theory in unitary organisms (19–22)—can be extended to predict the following traits of whole colonies: (i) rates of energy uptake and utilization (i.e., metabolic rates); (ii) rates of survival, growth, and reproduction; and (iii) reproductive allocation (i.e., gonad-to-soma mass ratio). We test model predictions using a large data set compiled on social insect life history. In doing so, we do not distinguish between, or advocate for, any model or models that predict 1/4 power scaling in unitary organisms.

We use metabolic scaling theory to generate and test five quantitative predictions with respect to colonies. First, we predict whole-colony metabolic rate will scale with whole-colony mass as $B = B_0M^{3/4}$, where B_0 is a taxon-specific, body temperature-dependent normalization constant that is independent of body size. Second, we predict whole-colony biomass production (P) will scale with colony mass as $P = P_0M^{3/4}$, where P_0 is a normalization constant that represents the fraction of total metabolism allocated to biomass production (20–23). Third, under this same assumption, we extend metabolic scaling theory to predict that gonad mass will scale with somatic tissue mass as $G = G_0M^{3/4}$, where G_0 is a normalization constant. Fourth, we predict that the ontogenetic growth of colonies, which is fueled by metabolism, will be described by the equation $dn/dt = am^{3/4} - bm$ (7, 8, 24) (*SI Text*). Equivalently, assuming worker mass does not change appreciably over the ontogeny of the colony and that colonies are composed only of workers, this equation can be expressed in terms of worker number as

$$dn/dt = an^{3/4} - bn. \quad [1]$$

Eq. 1 assumes that the metabolic energy of a colony is partitioned between the maintenance of existing workers and the creation of new workers: $B = B_w n + (E_w)dn/dt$, where B is whole-colony metabolic rate, E_w is the energy required to create a worker, and B_w is the metabolic rate of a single worker. As such, the coefficients in Eq. 1 are defined in biological terms such that $a = B_0 m_w^{3/4}/E_w$, and $b = B_w/E_w$, which represents an

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maximum life span value by 1.45, as used by Keller and Genoud (36). Additionally, life span data from colonies held in the laboratory were converted to average field life span by dividing values by 2.5, the ratio estimated and used for endotherms and for social and solitary insects by other authors (19, 28, 39–43). Data from termites were not included in this analysis because no reasonable estimates of queen life span as a function of colony size were available.

To compare life span data for colonies to those for unitary organisms, we assumed life span, LS , is inversely proportional to mass-specific metabolic rate, such that $LS \sim \mu B_0^{-1} M^{1/4}$, after normalizing for temperature and B_0 . Again, as noted above for the other analyses, we first corrected or normalized the temperature of all nonflying species to 31 °C. The nest tem-

peratures for bees, wasps, and ants were roughly constant as estimated on the basis of the data shown in *SI Text*. Then, to correct for B_0 , we used the equation $LS_{B_0\text{-corrected}} = LS_{31^\circ\text{C}} \times B_{0,F/NF} / B_{0,\text{insect}}$, where $B_{0,F/NF}$ is the average metabolic normalization constant for flying ($0.00692 \text{ W/g}^{0.75}$) or nonflying ($0.00189 \text{ W/g}^{0.75}$) insects at 31 °C (from ref 30; corrected to 31 °C), and $B_{0,\text{insect}}$ is the average constant ($0.00260 \text{ W/g}^{0.75}$) from the combined insect data (26). For some species, worker wet mass data were estimated from data on head width or body length using the equations and values listed in *SI Text*.

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