

Video Article

Experimental Manipulation of Body Size to Estimate Morphological Scaling Relationships in *Drosophila*

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Abstract

The scaling of body parts is a central feature of animal morphology¹⁻⁷. Within species, morphological traits need to be correctly proportioned to the body for the organism to function; larger individuals typically have larger body parts and smaller individuals generally have smaller body parts, such that overall body shape is maintained across a range of adult body sizes. The requirement for correct proportions means that individuals within species usually exhibit low variation in relative trait size. In contrast, relative trait size can vary dramatically among species and is a primary mechanism by which morphological diversity is produced. Over a century of comparative work has established these intra- and interspecific patterns^{3,4}.

Perhaps the most widely used approach to describe this variation is to calculate the scaling relationship between the size of two morphological traits using the allometric equation $y = bx^a$, where x and y are the size of the two traits, such as organ and body size^{8,9}. This equation describes the within-group (e.g., species, population) scaling relationship between two traits as both vary in size. Log-transformation of this equation produces a simple linear equation, $\log(y) = \log(b) + a\log(x)$ and log-log plots of the size of different traits among individuals of the same species typically reveal linear scaling with an intercept of $\log(b)$ and a slope of a , called the 'allometric coefficient'^{9,10}. Morphological variation among groups is described by differences in scaling relationship intercepts or slopes for a given trait pair. Consequently, variation in the parameters of the allometric equation (b and a) elegantly describes the shape variation captured in the relationship between organ and body size within and among biological groups (see^{11,12}).

Not all traits scale linearly with each other or with body size (e.g.,^{13,14}) Hence, morphological scaling relationships are most informative when the data are taken from the full range of trait sizes. Here we describe how simple experimental manipulation of diet can be used to produce the full range of body size in insects. This permits an estimation of the full scaling relationship for any given pair of traits, allowing a complete description of how shape covaries with size and a robust comparison of scaling relationship parameters among biological groups. Although we focus on *Drosophila*, our methodology should be applicable to nearly any fully metamorphic insect.

Video Link

The video component of this article can be found at <http://www.jove.com/video/3162/>

Protocol

1. Rearing and manipulation of diet to produce variation in body size and wing size

Rational and Overview. The expression of scaling relationships ultimately hinges on the developmental regulatory processes that produce variation in final organ and body size. These processes are best described from work on fully metamorphic (i.e., holometabolous) insects such as *D. melanogaster*, where organs grow as undifferentiated 'imaginal discs' within the grub-like larva^{11,15-17}. In *Drosophila*, final organ and body size are regulated by growth during the larval stages; larvae grow through the first and second larval instar until they reach a minimal viable weight for eclosion (MVW_E) towards the beginning of the third larval instar¹⁸. Starvation before attainment of MVW_E prevents adult eclosion. However, larvae that are starved at MVW_E generate viable adults, albeit at a very small size. This is because reduced nutrition during development reduces final body and organ size, not just in flies but in almost all animals¹⁹⁻²¹. By manipulating food availability after MVW_E is attained, but before pupation occurs, it is possible to induce extreme phenotypic plasticity in body size that greatly outweighs other sources of variation (e.g., individual genetic variation).

1. Eggs are collected from the population to be phenotyped, divided into lots of 50 and placed in vials containing 10ml of standard fly food. Because oviposition occurs continuously, eggs are collected every 24h for three days, producing three age cohorts.
2. When the oldest cohort reaches the wandering stage or has just begun to pupate, the diet manipulation is applied. At this point, the oldest cohort is well beyond the MVW_E, the youngest cohort is very close to the MVW_E and the remaining cohort is somewhere in between these extremes. Thus, removal of food at this point will produce very large adults from the oldest cohort, very small adults from the youngest cohort

and intermediate-sized adults from the remaining cohort. To remove the larvae from the vials, approximately 5ml of 40% sucrose solution is added to all vials. To free the larvae, the food is agitated for 15-20 minutes on a shaker table set to an intermediate speed.

3. Floating larvae are removed from the sucrose solution with a fine (00) paintbrush, placed into vials containing a wet cotton plug and allowed to pupate. Alternatively, larvae could be placed on a non-digestible food additive such as methyl cellulose²².

2. Estimation of body size

Rational and Overview. Previous studies indicate that thorax length is a less than ideal proxy for overall body size²². Instead, we use pupal size as a measure of body size. Maximum body size is fixed at the cessation of feeding that precedes pupation. As a result, there is a tight correlation between pupal size and adult body size²³.

1. Pupae are arranged ventral side up on glass microscope slides and imaged using a digital camera attached to a dissecting microscope. Each image is captured, assigned a unique identifying code and transferred onto a computer via a live feed connecting the digital camera to the computer.
2. Imaged pupae are moved to individual 2 ml eptubes containing 1ml of fly food, labeled with the unique code for that individual, and punctured with airholes to allow gas exchange.
3. Pupal size is estimated as the number of pixels of the pupal silhouette. Other size metrics could also be used.

3. Trait (wing) size estimation

Rational and Overview. To image wings, we use a 'wing grabber'²⁴, which pulls and holds the wing of a live, anesthetized fly perpendicular to its body. Pressed between two pieces of glass, this allows the wing to be imaged as a two dimensional object.

1. Flies are anesthetized lightly with CO₂ and placed in position in the wing grabber on a dissecting microscope fitted with a digital camera that is connected to a computer.
2. The wing is imaged and saved under the unique code for that individual.
3. Wing size can be estimated using the minimum convex polygon that circumscribes marginal fixed landmarks or as the pixel count of the wing silhouette.

4. Scaling relationship estimation

1. Data are pooled across cohorts but separated by sex and log-transformed.
2. Scaling relationships are fit to the size data using Type II (i.e., Reduced Major Axis) regression (see review in ¹²) and the parameters estimated for each group.

5. Representative Results:

Our diet manipulation will produce a broad range of body sizes (Figure 1) and trait sizes (Figure 2). Plotting the log transformed traits against each other reveals their pattern of covariation within a given biological group and permits fitting the scaling relationship to the group (Figure 3). Extraction of the scaling relationship parameters that describe how shape scales with body size allows comparison of how shape covaries with size across groups.



Figure 1. Pupa from extremes of the size distribution produced via diet manipulation after individuals achieve the MVW_E . Actual pupal images are shown at left with their corresponding silhouettes for the measurement of pupal size at right. Scale bar is 1mm.

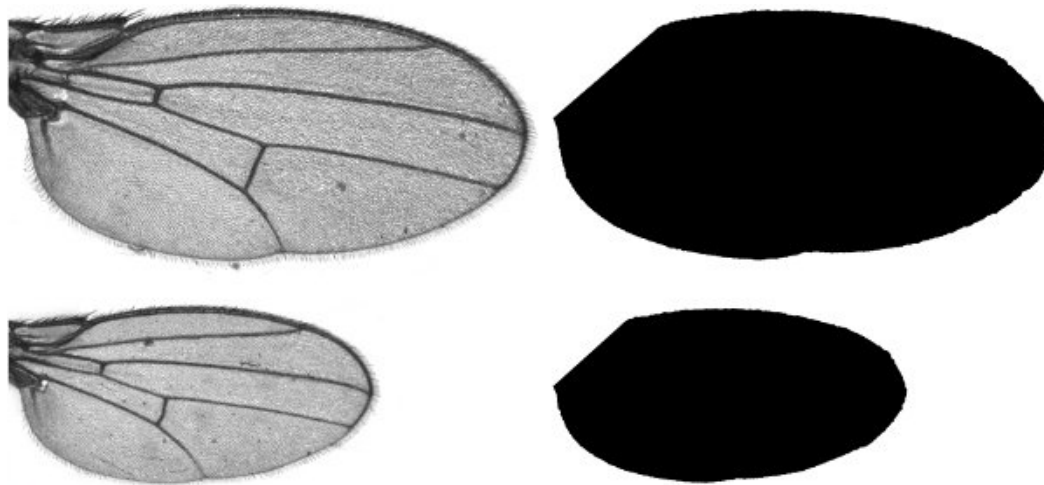


Figure 2. Wings taken from extremes of the body size distribution produced via diet manipulation after individuals achieve the MVW_E . Actual wing images are shown on the left with their corresponding silhouettes for the measurement of wing size shown on the right. Scale bar is 1mm.

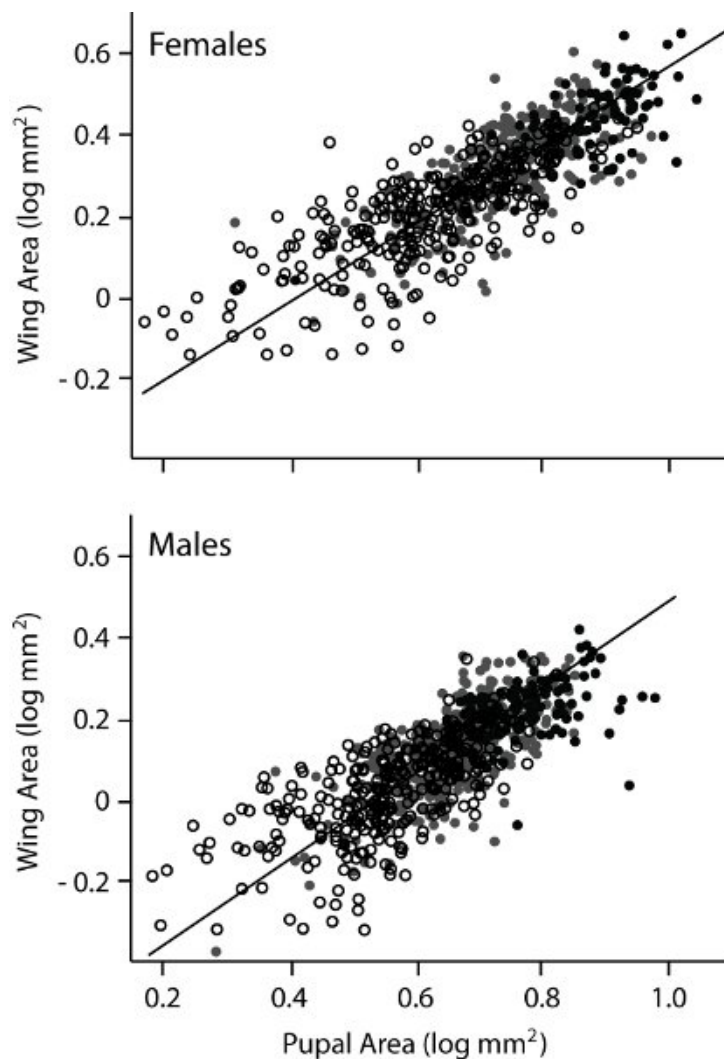


Figure 3. Wing-body size scaling for *Drosophila melanogaster* by sex. Log-transformed data for individuals of each sex that were fully-fed (dark circles), starved one (grey circles) or two days (open circles) as larvae. Type II regression line is shown for each sex. Between each age cohort,

declines in average wing area of ~0.1mm (~7%) and ~0.2mm (~8%) occurs, increasing the total range of phenotypes observed. The regressions reveal that females are slightly hypoallometric for this scaling relationship (slope of 0.96) whereas males are slightly hyperallometric (slope 1.06).

Discussion

Morphological scaling has been the subject of intense study in evolutionary biology for nearly a century. More recently, the proximate basis of the regulation of growth and scaling has become increasingly important in development, physiology and genetics (e.g., see reviews in ^{12,15,23-25}). Our methodology permits estimation of morphological scaling relationships across the entire range of trait and body sizes. This can be important, as trait size may not scale linearly with body size (e.g., ^{13,14}). However, this may not be apparent if animals are reared only under ideal lab conditions.

Although our protocol was developed explicitly to allow estimation of the wing-body size scaling relationship in *Drosophila*, it can be used with trait-specific modification to estimate the scaling relationship for any morphological trait in flies. Similarly, the diet manipulation can likely be used to produce the full range of trait and body sizes in other fully metamorphic insects; such applications simply require knowing when the MVW_E is achieved and applying the manipulation at that point in ontogeny.

Disclosures

No conflicts of interest declared.

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