Gliding towards an understanding of the origin of flight in bats

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Abstract

Bats are the only mammals capable of powered flight and have correspondingly specialized body plans, particularly in their limb morphology. The origin of bat flight is still not fully understood due to an uninformative fossil record but, from the perspective of a functional transition, it is widely hypothesized that bats evolved from gliding ancestors. Here, we test predictions of the gliding-to-flying hypothesis of the origin of bat flight by using phylogenetic comparative methods to model the evolution of forelimb and hindlimb traits on a dataset spanning four extinct bats and 231 extant mammals with diverse locomotor modes. Our results reveal that gliders exhibit adaptive trait optima (1) toward relatively elongate forelimbs that are intermediate between those of bats and non-gliding arborealists, and (2) toward relatively narrower but not longer hindlimbs that are intermediate between those of non-gliders and bats. We propose a hypothetical adaptive landscape based on limb length and width optimal trends derived from our modeling analyses. Our results support a hypothetical evolutionary pathway

wherein glider-like postcranial morphology precedes a bat-like morphology adapted to powered-

flight, setting a foundation for future developmental, biomechanical, and evolutionary research to

test this idea.

Introduction

Powered flight has evolved at least four times over the course of organismal evolution,

consistently serving as a gateway to ecomorphological and species diversification (Heard and

Hauser 1995; Hunter 1998; Jetz et al. 2012; Nicholson et al. 2014). As the only mammals

capable of powered flight, bats have correspondingly specialized body plans that include highly

derived limbs. These specialized morphologies are understood to be the result of adaptations for

the demands of flight; the skeletal elements of bat forelimbs are elongated to support flight

membranes and increase aerodynamic efficiency, whereas bat hindlimbs are not only specialized

for hanging and catching prey in flight (Norberg 1990; Gunnell and Simmons 2005; Giannini

2012), but also contribute to flight itself (Cheney et al. 2014).

The evolutionary origin of flight in bats remains a subject of ongoing research and debate

due to an uninformative fossil record linking bats to their non-flying ancestors. Nevertheless, bat

flight is widely hypothesized to be the result of an ancestral "trees-down" transition from non-

gliding tree-dweller (hereafter, 'arborealist') to glider to flyer (Norberg 1985B; Bishop 2008;

Simmons et al. 2008), although this idea has not been tested using phylogenetic comparative

methods. It should be noted that all phylogenomic evidence shows that bats are not closely

related to any extant glider (Upham et al., 2019; Álvarez-Carretero et al., 2022); bats are nested

within Laurasiatheria where neighboring clades are predominantly adapted to terrestrial

locomotion. Laurasiatheria is far from other placental gliders, including dermopterans and

rodents, located in the Eucarchontoglires superorder.

Following the arborealist-to-glider-to-flier hypothesis, the ancestors of bats–which

originated about 70-60 million years ago (Teeling et al. 2000; Upham et al. 2019; Álvarez-

Carretero et al. 2022)–are hypothesized to have evolved more elongated limbs, followed by

patagia (wing membranes) on the armwing to facilitate gliding, and finally membranes in the

handwing to enable powered flight. The armwing consists of stylopodal (i.e., humerus) and

zeugopodal (i.e., ulna and radius) bones, while the handwing consists of autopodal bones (i.e.,

the metacarpals and phalanges). The evolution of the handwing is thought to be the key

innovation that catalyzed bats' adaptive radiation (Vaughan 1970; Sears et al. 2006; Amador et

al. 2019), resulting in the evolution of over 1,500 extant species (Simmons and Cirranello

2022B).

Previous work has examined many biological features associated with the evolution of

bat flight, such as sensory adaptations (e.g., echolocation or vision), aerodynamics (e.g., aerofoil

and flight performance), ecology (e.g., diet, habitat, roosting), and development (e.g., patagia

development and digit elongation) (Bishop 2008; Cooper and Tabin 2008; Giannini 2012;

Anthwal et al., 2023; Feigin et al., 2023). Morphological approaches to understanding the origin

of bat flight further include analyses of skeletal dimensions in the context of development

(Adams 1992, 2008; Sears et al. 2006; Nojiri et al. 2023), biomechanics (Swartz et al. 1992;

Swartz 1997; Swartz and Middleton 2008), and functional morphology (Vaughan 1970). These

studies often focus on aspects of the bat forelimb (e.g., Amador et al. 2019) or hindlimb (e.g.,

Riskin et al. 2005), but seldom both in tandem in a phylogenetic context (but see Swartz and

Middleton 2008 and Nojiri et al. 2023). Here, we test predictions based on the gliding-to-flying

hypothesis of the origin of bat flight by modeling the evolution of forelimb and hindlimb traits

on a dataset spanning four extinct bats and 231 extant mammals with diverse locomotor modes

including ground-dwellers, arborealists, gliders, and flyers.

Following the arborealist-to-glider-to-bat hypothesis, we test two predictions regarding

the evolution of limb morphologies using an Ornstein-Uhlenbeck evolutionary model-fitting

approach. First, we predict that selection will lead the forelimb skeletal traits of arborealists,

gliders, and bats towards three separate adaptive trait optima/attractors (i.e., θ parameters of

fitted OU models, which are commonly interpreted as the locations of adaptive peaks in

morphospace; hereafter "optima"), where the glider forelimb optimum will be intermediate in

value to those of arborealists and bats. If supported, this demonstrates that glider morphology

may fall on an intermediate adaptive peak between bat and arborealist morphologies.

Specifically, we predict wider and longer skeletal element optima values progressing from

arborealists to gliders to bats. Relatively longer bones (especially in the handwing) and larger-

diameter bones (especially in combination with thinner, highly mineralized walls), may give bats

enough resistance to torsion during powered flight (Swartz et al. 1992) while still increasing their

patagia area. Glider optima with intermediate forelimb length and width optima between

arborealist and bat optima may represent an intermediate adaptive zone between the anatomical

adaptations required for tree-dwelling (Cartmill 1985) and those required for powered flight

(Norberg 1990). Support for this prediction would suggest selective pressures for flight

influenced the evolution of bat forelimb skeletal elements from glider-like forelimbs.

Second, we predict that bat hindlimb optima will exhibit a distinct trait optimum (i.e.,

adaptive peak) separate from ground-dwelling, arboreal, and gliding mammals. Previous work

has considered bat hindlimbs to be "reduced" in size (e.g., Farnum et al. 2008; Simmons et al.

2008; Louzada et al. 2019), whereas gliders exhibit relatively lengthened hindlimbs (Grossnickle

et al. 2020; Linden et al. 2023; Rickman et al. 2023). Therefore, the relative lengths and widths

of the hindlimb bones are not well characterized between these aerial groups as well as in

relation to arborealists. Here, we aim to identify where bat hindlimb adaptive optima lie relative

to other mammals. Specifically, we predict narrower and longer skeletal element optima values

progressing from arborealists to bats and gliders. We predict overlapping hindlimb optima

between gliders and bats because relatively longer and lower-diameter hindlimb bones in both

groups relative to arborealists and ground-dwellers may decrease weight while still generating

lift. Support for this prediction would suggest that shared selective pressures for less-

quadrupedal motion (i.e., as gliding and flying both incorporate hindlimbs into the wing

membrane) influenced the evolution of bat hindlimb skeletal elements from ancestors with

glider-like hindlimbs. Overall, this study offers the first phylogenetically informed test of

morphological data to investigate whether the adaptive optima of arborealists, gliders, and flyers

illustrate a possible route across adaptive peaks in the evolution of mammalian flight.

Materials & Methods

Morphological Data

We compiled linear measurements of 14 forelimb and 15 hindlimb traits (Fig. 1) from

231 extant mammals (1–2 specimens per species) across 21 orders (78% of mammalian orders)

and 73 families (47% of mammalian families). Of these, 26 representative species of bats (48

specimens) were measured for the flyer group. Linear measurements were obtained from

specimens held at the Burke Museum of Natural History and Culture or from previously

published work (Chen and Wilson 2015; Grossnickle et al. 2020; Weaver and Grossnickle 2020;

Pevsner et al. 2022). All specimens measured for this study were fully mature, determined by the

closure of exoccipital–basioccipital and basisphenoid–basioccipital sutures on the cranium and

ossification of all vertebrae and limb bones. Specimens were a mixture of male, female, and

unknown sexes. We size-corrected all linear measurements by calculating log-shape ratios of

each trait measurement (Mosimann 1970) by $\log_{10}$-transforming the mean trait value divided by

the cube root of body mass prior to running all statistical analyses. Unfortunately, most

specimens lacked accompanying body mass data; therefore we compiled species-level body mass

data from the PanTHERIA database (Jones et al. 2009) and the *Handbook of the Mammals of the*

World book series (Wilson and Mittermeier 2009).

We also collected the 29 measurements for four extinct bats (*Onychonycteris finneyi,*

Icaronycteris index, Hassianycteris messelensis, and Tachypteron franzeni) using ImageJ

(Schneider et al. 2012) on published full-body images that include scale bars (Storch et al. 2002;

Gunnell and Simmons 2005; Simmons et al. 2008). We estimated the body masses of extinct bats

using a linear regression between the body mass and geometric mean of all linear measurements

from the extant bats of our dataset and used those body mass estimations to size-correct our

fossil data again using log-shape ratios. We included fossil bats in our morphological dataset

because, although extinct bats have a high degree of morphological similarity to extant bats

(Norberg 1985B; Bishop 2008; Simmons et al. 2008), the fossil bat data may alter model

estimations of the adaptive zone between flyers and gliders in an informative way. For example,

the inclusion of fossils may make the flyer-glider trait optima more similar or more disparate by

adding more phylogenetic history information and thus inform the understanding of the adaptive

landscape underlying the evolution of mammalian flight.

To visualize the distribution of locomotor groups in the forelimb and hindlimb

morphospaces, we performed principal component analyses (PCA) of the morphometric data

using the *princomp* function in the *R* package *stats* (R Core Team 2021).

Locomotor Data

We categorized species into four locomotor groups: ground dwellers ($n = 117$),

arborealists ($n = 71$), gliders ($n = 16$), and flyers ($n = 27$) following previous work (Chen and

Wilson 2015; Grossnickle et al. 2020; Weaver and Grossnickle 2020; Pevsner et al. 2022). The

ground-dwelling mammals included diverse ambulatory, cursorial, saltatorial, and fossorial taxa.

This 'catch-all' category was intentional as we are interested in investigating the arboreal to

gliding to flying specialization trends only. Our arborealist category was also broad in that it

includes both fully arboreal (tree-dwelling) and scansorial taxa. Gliders included tree-dwelling

mammals with derived morphologies (i.e., patagia and elongated limbs) for gliding locomotion

(Grossnickle et al. 2020). Flyers included bats only.

Phylogenetic Comparative Analyses

We used evolutionary models (Hansen 1997; Butler and King 2004) to examine the

influence of locomotor modes on the evolution of forelimb and hindlimb morphologies in our

extant-only and extant+extinct datasets. We first tested the hypothesis that locomotor modes

influenced the evolution of the forelimb and hindlimb using multivariate evolutionary models on

our extant datasets (Butler & King, 2004; Hansen, 1997) in the R package mvMORPH (Clavel et

al., 2015). To reduce the dimensions of our data, we conducted a principal component analysis

(PCA) on our 14-trait forelimb dataset and 15-trait hindlimb dataset. We retained the first three

principal components (PCs) of our forelimb dataset (91.8% of the forelimb variation) and

hindlimb dataset (85.3% of hindlimb variation). We then fit six multivariate models to the scores

of these three PCs. We fit a multivariate Brownian motion model (mvBM), which models trait

variance in a random walk over time; a single-regime multivariate Ornstein-Uhlenbeck model

(mvOU1), in which scores of each PC evolve towards a single trait optimum (θ parameter of OU

models), giving a single mean optimum for all locomotor groups; and multivariate OU models

(mvOUM) with a given number of optima corresponding to selective regimes (based on

locomotor modes). To test our hypotheses on overall limb evolution, we fit a four-peak OU

model (mvOUM$_{loc4}$) for distinct optima among ground-dweller, arborealist, glider, and flyer

regimes. To test alternative hypotheses, we fit three three-peak models (mvOUM$_{loc3a-c}$) with

distinct combinations of the four locomotor groups. The mvOUM$_{loc3a}$ model tests for distinct

optima among ground-dweller, arborealist, and a combined glider-flyer regimes. The

mvOUM$_{loc3b}$ model tests for distinct optima among ground-dweller, combined arborealist-glider,

and flyer regimes. The mvOUM$_{loc3c}$ model tests for distinct optima among a combined ground-

dweller-arborealist, glider, and flyer regimes.

To account for uncertainty in phylogenetic topology and the ancestral character states

(Fig. 1), we used the *make.simmaps* function in the *R* package *phytools* (Revell 2012) to simulate

10 stochastic character maps across 1,000 tree topologies randomly drawn from the posterior

distribution of "completed" trees from Upham et al. (2019), resulting in 10,000 character maps

of locomotory regimes. We randomly sampled 250 trees and fit all six models to these trees. We

assessed relative model support using small sample corrected Akaike weights (AICcW). Models

with ΔAICc scores less than two were considered supported and interpreted as evidence of

locomotor groups influencing species' adaptive zones. We acknowledge that using a subset of

PCs instead of the full dataset could bias our results (D. C. Adams & Collyer, 2018; Uyeda et al.,

2015). However, our results remained consistent when we increased the number of PCs in our

model-fitting analyses to the first four or five PCs. Further, we used simulations to assess

whether we had adequate power to accurately distinguish between complex mvOU models from

Brownian motion (Boettiger et al., 2012).

If the mvOUM$_{loc4}$ model is best-supported, we interpreted this as each locomotor regime

having its own distinct adaptive optimum. This result alone does not provide evidence for the

evolutionary origin of bats. However, if gliders are located at an optimum between arborealists

and flyers (i.e., if arborealist < glider < flyer optima or arborealist > glider > flyer optima), there

is support for an evolutionary "pathway" between the two locomotor modes relative to an

ancestral arborealist or ground-dwelling state (i.e., one could theoretically bridge adaptive peaks

via these pathways). Therefore, if the optima gradually change in value (as opposed to random

changes in value) from arborealists to gliders to bats, our hypothesis that bats evolved from

gliding ancestors is supported.

If the mvOUM$_{loc3a}$ model is best-supported, we interpret this as gliders+flyers having a

shared adaptive optimum and arborealists and ground-dwellers having their own distinct

evolutionary optima. This model provides the clearest support for flyers evolving from gliding

ancestors because lineages of both groups are evolving towards the same shared adaptive optima.

Therefore, if this model best fit the data, our hypothesis is supported.

If the mvOUM$_{loc3b}$ model is best-supported, we interpret this as arborealists+gliders

having a shared adaptive optimum and flyers and ground-dwellers having their own distinct

evolutionary optima. There is phylogenetic justification for this grouping because all gliders

evolved from ancestrally arboreal groups (e.g., gliding squirrels evolved from tree squirrels; see

tree in Fig. 1). These groups are similar to the 4-group model in that if arborealists+gliders have

an optimum between those of ground-dwellers and flyers, there is support for an evolutionary

"pathway" between the two locomotor modes relative to an ancestral ground-dwelling state.

209   Finally, if mvOUM$_{loc3c}$ model is best-supported, we interpret this as ground-

dwellers+arborealists having a shared adaptive optimum and flyers and gliders having their own

distinct evolutionary optima. As with the mvOUM$_{loc4}$ and mvOUM$_{loc3b}$ models, if gliders have an

optimum between those of ground-dwellers+arborealists and flyers, there is support for an

adaptive "pathway" between the two locomotor modes relative to an ancestral arborealist or

ground-dwelling state, providing evidence for our hypothesis that flyers evolved from gliding

ancestors.

216   We also examined the influence of locomotor modes on the evolution of each individual

forelimb and hindlimb trait in our extant-only and extant+extinct datasets using univariate

evolutionary models. For each of the 29 traits, we fit a Brownian motion model (BM), a single-

optimum Ornstein-Uhlenbeck model (OU1), and four multiple-peak OU (OUM) models. The

OUM models included OUM$_{loc4,}$ OUM$_{loc3a,}$ OUM$_{loc3b}$, and OUM$_{loc3c}$, featuring the same

locomotor regimes as the multivariate mvMORPH models. All models were fit using the R

package $OUwie$ (Beaulieu et al. 2012). We applied a single selection parameter (α) and rate

parameter (σ^2) across all selective regimes while allowing optima (θ) to vary between regimes

(Hansen 1997; Butler and King 2004). For traits in which an OUM model was the best-supported

model, we computed 95% confidence intervals using parametric bootstrapping of the model

parameters with the $OUwie.boot$ function in the R package $OUwie$ (Beaulieu et al. 2012). We

randomly sampled 100 of our character maps and performed 10 bootstrap replicates per

stochastic character map sampled.

We repeated the above multivariate and univariate model fitting approaches on a merged

dataset that includes extant species and the four fossil species. All four models were fit across 50

stochastically mapped trees to account for uncertainty in the ancestral character states. We

manually placed the four extinct bats on the phylogeny as a polytomy with extant bats at 69.8

MYA.

Results

Forelimb morphospace and evolutionary model-fitting

PC1 explains 81.5% of the forelimb variation and separates flyers (both extant and

extinct) from the remaining three locomotor modes (Fig. 2a; Table S1a). PC1 also slightly

separates gliders from arborealists and ground-dwellers. This axis describes the relative lengths

of the armwing (humerus, radius, ulna) and the handwing (metacarpals and phalanges). PC1

varies across relatively shorter (-PC1) to elongate (+PC1) bones, revealing a trend within the

dataset in the elongation of multiple forelimb bone elements from arborealists to gliders to flyers.

PC2 explains 7.8% of the morphological variation and primarily describes the relative widths of

the long bones and hands. This variation of wide to narrow bones mostly occurs within

locomotor groups, especially within the diverse ground-dweller group. PC3 explains 2.9% of the

morphological variation with +PC3 dominated by metacarpal length, ulna olecranon length, and

scapula height and -PC3 dominated by intermediate phalanx length and humerus distal width.

Fossil bats overlap with extant bats in morphospace, which is consistent with the lack of

transitional forms in the bat fossil record.

The mvOUM$_{loc3b}$ model, which assigns arborealists and gliders to the same regime, was

the best-fitting model for the extant multivariate forelimb dataset (PCs 1-3) (AICcW = 0.99,

Table 1a). Distinct optima were supported for these three locomotor groups, with PC1 and PC2

optima scores increasing in value from ground-dweller to arborealist+glider to flyer hindlimbs

(Fig. 2c). Repeating the multivariate model-fitting with the addition of the four extinct flyer

species yielded a mvOUM$_{loc4}$ model as the best fit (mAICcW = 0.99; Table S6a). However, the

optima exhibited unrealistic magnitudes (Table S6b).

When we examined each forelimb trait individually using univariate models, we found

that the OUM$_{loc4}$ model was the best-fitting model for six of the 14 forelimb traits (AICcW =

0.51-1.00), OUM$_{loc3b}$ was the best-fitting model for five traits (AICcW = 0.36-0.65), OUM$_{loc3c}$

was the best fitting model for metacarpal length (AICcW = 0.63), and OUM$_{loc3a}$ was the best-

fitting model for ulna olecranon length (AICcW = 0.58; Table S2a). The OU1 model was the

best fit for the proximal phalanx width of the third manual digit (AICcW = 0.51). Armwing (i.e.,

humerus and radius) length traits have optima (from the best-fitting OUM$_{loc4}$ models) that

gradually increase from ground-dwellers to arborealists to gliders to flyers; parametric

bootstrapping revealed little to no overlap in the 95% confidence intervals of optima between

most locomotor groups (Fig. 3b-c). Optima of armwing widths (from the best-fitting OUM$_{loc3b}$

models) indicated that flyers exhibit relatively wider humeri at the distal end, proximal end, and

mid-shaft compared to all other locomotor groups (Fig. 3g-i; Table S3a). Flyers exhibit relatively

higher optima for handwing length traits (i.e., longer third manual metacarpal, proximal phalanx,

and intermediate phalanx) than all other locomotor groups (Fig. 3d-f; Table S3a). For each,

glider optima were intermediate between ground-dweller and arborealist optima. For handwing

width traits, flyers exhibit relatively higher OUM$_{loc3b}$ optima for metacarpal widths compared to

all other locomotor groups (Fig. 3j), but no differences in proximal phalanx width as evidenced

by OU1 support for this trait. An adaptive optimum for a flyer-glider combined group in the

OUM$_{loc3a}$ models was supported for ulna olecranon length only (Table S2a). In total, however,

glider optima (in combination with arborealists) were between the 95% confidence intervals of

ground-dweller and flyer optima for all traits except scapula length/width, humerus proximal

width, and metacarpal width (Fig. 3a, 3g, 3j; Table S3a). For these last trait optima,

glider+arborealist optima were statistically the same as that of ground-dwellers while flyers

exhibited higher optima than all other locomotor groups (Fig. 3a, 3g, 3j; Table S3a).

281   Adding fossil flyers into forelimb model-fitting analyses yielded overall slightly more

support for OUM$_{loc3a}$ models compared to extant-only modeling; scapula height and humerus

proximal width demonstrate support for a flyer-glider combined group (Table S4a). OUM$_{loc4}$ and

OUM$_{loc3b}$ models were still widely supported as the best-fitting model for six traits and five

traits, respectively; OU1 was still the best-fitting model for proximal phalanx width (Table S4a).

Overall, the length and width optima values for the forelimb were very similar between

extant+extinct and extant only-modeling (Table S5a).

Hindlimb morphospace and evolutionary model-fitting

290   PC1 explains 47.5% of hindlimb variation and more positive PC1 scores describe the

relative elongation (+PC1) of the fibula, metatarsal, and pedal proximal and intermediate

phalanges (Fig. 2b; Table S1b). PC1 slightly separates gliders and flyers, with gliders falling

towards +PC1 and flyers falling towards -PC1, indicating glider hindlimb autopodia are overall

more elongated than flyers'. Arborealists and ground-dwellers are spread across PC1. PC2

explains 17.8% of the variation, where +PC2 is associated with relatively lengthened pedal

proximal phalanges, intermediate phalanges, and ischium and -PC2 with relatively narrower

pedal proximal and intermediate phalanges. PC2 loosely separated ground-dwellers, arborealists,

and the glider-flyer group. PC3 explains 11.3% of the morphological variation with +PC3

dominated by ilium length, femur length, and fibula length and -PC3 dominated by metatarsal

width, intermediate phalanx width, and proximal phalanx width. These loadings show that

variation between the four locomotor groups mainly occurs throughout the hindlimb autopodia

region.

The mvOUM$_{loc4}$ model was the best-fitting model for the extant multivariate hindlimb

dataset (PCs 1-3) (AICcW = 0.92, Table 1b). Distinct optima were supported for the four

locomotor regimes, with PC1 and PC2 optima scores decreasing in value from ground-dweller to

arborealist to glider to flyer hindlimbs, albeit with overlap (Fig. 2d). Adding in the four extinct

flyer species yielded again the mvOUM$_{loc4}$ model as the best fit, and very little change in the

optima values (Table S6).

When we examined each hindlimb trait individually using univariate models, we found

that the OUM$_{loc4}$ model was the best-fitting model for three of 15 traits, all of which were

hindlimb long bone lengths (AICcW = 0.33-0.60; Table S2b). Specifically, gliders have

relatively longer tibia and femoral optima compared to flyers and the other locomotor groups

(Fig. 4b-c, Table S3b). Fibula length was the third hindlimb trait supported by OUM$_{loc4}$, but

fibulae are rudimentary in bats. The OUM$_{loc3a}$ model best fit three traits (AICcW = 0.36-0.52),

all of which were widths: femur shaft width, metatarsal width, and pedal proximal phalanx width

(Table S3b). Parametric bootstrapping of these OUM$_{loc3a}$-supported models revealed that the

glider-flyer group had relatively narrower femur shafts, metatarsals, and pedal proximal

phalanges optima than the other locomotor groups (Fig. 4f, i, j; Table S3b). The OUM$_{loc3b}$ model

was the best-fitting model for four of the 15 traits (AICcW = 0.44-0.71): ilium, ischium,

metatarsal, and intermediate proximal phalanx length (Table S2b). Flyers had significantly

smaller relative ilium and ischium length optima than other locomotor groups.

Gliders+arborealists had metatarsal and intermediate proximal phalanx optima intermediate

between flyers and ground-dwellers (Fig. 4d-e; Table S3b). The OUM$_{loc3c}$ model best fit three

traits (AICcW = 0.42-0.48): pelvis length, femur distal width, and tibia proximal width (Table

S2b). In the latter two traits, gliders exhibited intermediate width optima between flyers (which

were smaller) and ground-dwellers+arborealists (where were larger). For pelvis length, like ilium

and ischium length, there was no clear trend in the direction of the optima values across

locomotor groups (Table S3b).

Adding fossil flyers into hindlimb model-fitting analyses yielded only one difference

compared to extant-only modeling: pelvis length was now best fit with a OUM$_{loc3b}$ instead of

OUM$_{loc3c}$ model (Table S4b). The length and width optima for the hindlimb were very similar

between extant+extinct and extant-only modeling (Table S5b).

Discussion

Forelimb trait evolution

We found evidence that mammalian glider forelimb morphologies are evolving toward

adaptive peaks that are intermediate between those of arborealists and flyers. Our multivariate

forelimb results and univariate length trait results provide support for the hypothesis that

selective pressures for flight influenced the evolution of flyer forelimbs from glider-like

forelimbs. Using multivariate evolutionary models, we found a three optima model with distinct

selective regimes for ground-dwellers, arborealists+gliders, and flyers as the best fit to the first

three PCs of our data (Table 1a). The arborealist+glider optima are intermediate between those

of flyers and ground-dwellers (Fig. 2c). PC1 is mostly explained by the variation in the forelimb

bone lengths and PC2 by their widths, creating a coarse picture that forelimb bones likely

become relatively longer and possibly wider from ground-dwellers to arborealists+gliders to

flyers. Results from our univariate evolutionary models are consistent with these findings,

revealing that glider forelimb optima are intermediate between flyer and arborealist optima for

length traits (Fig. 3a-f), but only partially for width traits (Fig. 3g-j) across both the armwing and

handwing (See Table S7 for a breakdown of whether each univariate model supports our

hypothesis).

Flyers exhibit trait optima indicating evolution of relatively longer forelimb bones

compared to all other locomotor groups (Fig. 2a, Fig. 3a-f) particularly in the handwing, which

exhibited rapid evolution as a necessary adaptation for bats (Vaughan 1970; Sears et al. 2006;

Amador et al. 2019). Increased bone lengths may enhance the surface area of the patagium for

greater lift (Norberg 1985b). Flyers also displayed evidence of evolution toward relatively wider

bones in the armwing compared to all other locomotor groups (Fig. 3g-j) consistent with findings

that bat humeri and radii exhibit significantly larger diameters compared to nonvolant mammals

(Swartz and Middleton 2008). These relatively wider proximal forelimb bones combined with

thinner yet highly mineralized walls provide resistance to bending and torsion during flapping

(Swartz et al. 1992; Papadimitrious et al. 1996). Furthermore, the transition from relatively more

robust proximal ends of the humerus to relatively less robust distal ends may also reflect greater

muscle mass closer to the center of rotation of the wing joints, which would greatly decrease

flight inertial power and facilitate economical flight (Norberg 1985A; Swartz 1997). In the

handwing, we found flyers exhibit relatively thicker third metacarpals (Fig. 3j) that corresponded

with its relatively increased length (Fig. 3d) compared to nonvolant mammals. In contrast, the

proximal phalanx length of the third manual digit was relatively longer compared to nonvolant

mammals whereas its width was best supported with an OU1 model, consistent with previous

work showing that the proximal phalanx diameter is not significantly smaller in bats compared to

nonvolant mammals (Swartz and Middleton 2008). As a result, bats exhibit relatively thick bone

cortices to combat greater bending moments associated with this relatively increased elongation

(Swartz et al. 1992; Papadimitriou et al. 1996; Swartz 1997). Increased cortical thickness along

with the reduction of mineralization of the handwing work together to resist bending failure

while simultaneously maintaining flexibility during flying (Papadimitriou et al. 1996).

In gliders, we found evidence that forelimb bone length optima are largely intermediate

between ground-dwellers and flyers. Gliders tend to exhibit relatively greater length optima in

most forelimb bones compared to ground-dwelling and arboreal mammals (Thorington and

Heaney 1981; Grossnickle et al. 2020; Linden et al. 2023; Rickman et al. 2023) but do not reach

the lengths of flyer forelimb bones. Nevertheless, like flyers, the longer bones expand the surface

area of the patagium (Thorington and Heaney 1981) and enable further adjustment of their

patagium during flight, leading to more efficient aerodynamics, increased agility, and

maneuverability (Thorington et al. 1998; Zhao et al. 2019). Forelimb bone widths in gliders,

while clearly intermediate between those of ground-dwellers and flyers for some traits (Fig. 3h,

i) share an optimum with arborealists (shown by the OUM_{loc3b} support for these traits; Fig. 3g-j)

and overlap with ground-dwelling optima mammals for other traits (Fig. 3g, j). Together, these

results suggest gliders may maintain high relative widths while continuing to elongate forelimb

bones, supporting previous findings that the evolutionary elongation of glider forelimbs is not

also accompanied by slimming of the bones (Grossnickle et al. 2020).

Hindlimb trait evolution

We found mixed evidence that mammal glider hindlimb morphologies are intermediate to

those of arborealists and flyers. Using our multivariate modeling approach, we found distinct

optima for ground-dwellers, arborealists, gliders, and flyers that best fit the first three PCs of our

hindlimb data (Table 1b). The trait loadings indicate that the autopodial bones dictate the

distribution of PC scores in PCs 1 and 2 (Table S1b). However, there is much overlap among the

optima of ground-dwellers, arborealists, and gliders in morphospace (Fig. 2d).

The univariate models uncovered further patterns: 1) width optima of hindlimb bones

were consistently narrower in flyers and gliders compared to arboreal mammals (Fig. 4f-j), and

2) there was no pattern from arboreal to glider to flyer in the hindlimb length optima. Univariate

modeling revealed that the widths of hindlimb bones tend not to differ between flyers and gliders

(Fig. 4f-j) because the widths of hindlimb bones in flyers and gliders exhibited either a single

optimum (i.e., OUM$_{loc3a}$) separate from ground-dwelling and arboreal mammals (Fig. 4 f, i, j) or

an overlap in confidence intervals of optima between flyers and gliders (Fig. 4 g, h). This shared

flyer-glider adaptive zone suggests that flyers and gliders exhibit narrower hindlimb optima,

especially in the autopod region (Fig. 4f-j). In contrast, univariate modeling revealed distinct

optima in long bone lengths of the hindlimb among the four locomotor regimes, in which gliders

exhibited relatively longer femoral and tibial optimal lengths compared to flyers, arborealists,

and ground-dwellers (Fig. 4b-c). Unlike bat forelimbs, bat long bones in the hindlimb (i.e., the

femur and tibia) are not the longest compared to the other locomotor regimes; in fact, the

confidence intervals of optimal lengths largely overlap between flyers and arborealists (Table

S3b). Furthermore, flyers exhibit relatively shorter metatarsals but relatively longer pedal

intermediate phalanges compared to all other locomotor regimes, whereas these autopodial

lengths do not differ between gliders and arborealists (Fig. 4d-e). The different evolutionary

patterns of hindlimb bones demonstrate the unique locomotor diversity among the four regimes.

Although gliders and flyers are both aerial, these behaviors require different functional

morphologies. In gliders, relatively longer hindlimbs are needed to increase the surface area of

the patagium in tandem with relatively longer forelimbs to generate lift during gliding

(Thorington and Heaney 1981). In contrast, in bats, the relative lengthening of the forelimb

supports the majority of the wing membranes whereas the hindlimb, specifically the ankle,

serves as an anchor to the wing membrane. Bat hindlimbs are still used for flight by helping

modulate lift, drag, or pitch (Cheney et al. 2014).

Overall, gliders appear to be evolving relatively lengthened hindlimb bones while bats

are evolving more gracile hindlimb bones. In light of these results, we reject our prediction that

flyer and glider hindlimbs overlap in their adaptive optima. This finding may point to the fact

that modern gliders are completely distinct evolutionarily and potentially even morphologically

from the gliding ancestors of bats. These results may highlight the pitfalls of modeling modern

gliders as proxies for ancestral ones instead of invalidating our hypothesis that bats evolved from

ancestral gliders.

Incorporating fossil bats

The morphospace positions of fossil bats in the forelimb PCA confirm extinct bats'

morphological similarity to extant bats in the degree of forelimb specialization (Fig. 2a). Amador

et al. (2019) modeled aerodynamic traits instead of skeletal traits and found that the Eocene bat

Onchonycteris finneyi occupies an area of aerofoil morphospace between extant gliders and bats.

In our morphospace, *O. finneyi* is the closest fossil bat to gliders across PC1 (Fig. S1a),

congruent with Amador et al.'s (2019) findings.

Fossil flyers occupy the same region of hindlimb morphospace as extant flyers, but three

of four species overlap clearly with the region of morphospace of non-flyers, a higher proportion

compared to that of extant bats (Fig. 2b). *Tachypteron franzeni* most clearly overlaps with extant

flyers and is also the only fossil we included from an extant bat family, Emballonuridae (Storch

et al. 2002). Simmons et al. (2008) suggested that, based on its intermediate limb traits,

Onychonycteris finneyi was likely more proficient at climbing than extant bats and may have

employed quadrupedal locomotion and under-branch hanging. Interestingly, *O. finneyi* and

Desmodus rotundus (the common vampire bat) are almost indistinguishable in the hindlimb

morphospace (Fig. S1b). Vampire bats are known for their adept quadrupedal locomotion and

climbing (e.g., Riskin et al. 2005), so *O. finneyi's* similar morphology may lend support to more

walking-capable early bats. However, the linear measurements we used here may not reflect the

complex adaptations required for these behaviors (e.g., ninety-degree joint rotations for under-

branch hanging, Riskin et al. 2005), so the placement of these species in morphospace leaves

many possibilities open for ancestral locomotor behaviors.

Including fossil flyers in our model-fitting analyses as the closest morphologies we have

to flyer-glider intermediaries did not significantly alter estimations of the adaptive zone between

flyers and gliders; the inclusion of fossils appeared to make the flyer-glider trait optima

marginally more similar for the forelimbs and had no change for the hindlimbs (Table S7). Fossil

addition to the multivariate model-fitting analysis shifted support from mvOUM$_{loc3b}$ to

mvOUM$_{loc4}$, suggesting that the glider optimum is potentially distancing itself from arborealists'

and towards flyers' (Table S6a). Unfortunately, the extant+extinct mvOUM$_{loc4}$ model predicted

extremely negative optima values, which represent unrealistic morphologies that are potentially

artifactual, making it difficult to gain more insight into this result (Table S6b). Fossil addition to

univariate forelimb model-fitting analyses yielded overall slightly more support for OUM$_{loc3a}$

models compared to extant-only modeling (Table S4a). Fossil addition to hindlimb model-fitting

analyses yielded overall no change in support for the mvOUM$_{loc4}$ model or OUM$_{loc3a}$ models

compared to extant-only modeling (Table S6, S4b). Regardless, overall the length and width

trends are the same as above between locomotor groups across the fore- and hindlimb (Table

S5).

Limb trait optima and the adaptive landscape

Our results provide insight on the potential adaptive landscape connecting arboreal,

gliding, and flying locomotion (Fig. 5). Overall, we found that glider forelimb optima fall

intermediate between those of arborealists and flyers in their lengths but only partially in their

widths. Except for long bone lengths, glider hindlimb optima are intermediate between ground-

dweller/arborealist and flyer optima or even overlap with flyer optima. Together, these results

emphasize that selection on key length/width traits may be "pulling" glider traits towards a flying

adaptive zone yet that certain glider traits (e.g., femur and tibia lengths) are derived specifically

for gliding morphology. Bishop (2008) proposed that the high degree of convergence among

gliders suggests functional constraints on the evolution of powered flight from gliding; however,

more recent work found that gliding lineages display incomplete convergence (species retain

unique morphologies even while evolving some similarities; Leal et al. 2002; Stayton 2006). For

gliders, there is little evidence that gliding is under strong selective pressures toward a single-

optimum gliding morphology (Grossnickle et al. 2020). The clustering of bat species in forelimb

morphospace (Fig. 2a) may represent a narrow adaptive peak within which forelimb

morphologies are highly conserved. In contrast, the broad glider morphospace may represent a

gradually inclined selection plane, with gliders still far from a peak (Grossnickle et al. 2020).

This broad adaptive slope could represent the foothill of a steep flyer peak such that gliders are

very slowly evolving towards a flyer peak (Fig. 5a). The evolutionarily oldest extant glider

lineages, dermopterans (colugos) and anomalurids (scaly-tailed squirrels), have evolved farthest

from ancestral arborealists (Grossnickle et al. 2020; although some extinct gliders may have

experienced relatively greater change from ancestral arborealists, e.g., Meng et al. 2017; Luo et

al. 2017). Here, the dermopteran species *Cynocephalus volans* and *Galeopteras variegatus* are

the gliders closest to the flyer forelimb region of morphospace (Fig. 3a; Fig. S1a), suggesting a

trajectory in morphospace from gliding to flying. Further, colugos are known as the "mitten

gliders" because of their interdigital membranes, which, along with their common carpal

morphology with bats (Simmons and Geisler 1998), reflects hypotheses that interdigital webbing

was key to evolving powered flight (Anderson and Ruxton 2020 but see Gardner and Dececchi

2022 for an alternate view).

Our results also suggest that both flyers and gliders exhibit broad adaptive optima (i.e.,

having large confidence intervals) in the hindlimb due to gliders sharing an optimum or

overlapping optima with flyers, or being intermediate between non-flyers and flyers for many

traits (Fig. 2d, Fig. 4). These broad adaptive optima could represent overlapping peaks for gliders

and flyers separated only by a shallow fitness valley driven by the gliders' elongated femur and

tibia lengths (Fig. 5b). The hypothetical adaptive landscape is therefore valuable because the

transition from quadrupedalism to hindlimb suspension has been hypothesized to enable

forelimbs to evolve more freely since they would no longer be weight-bearing structures (Bishop

2008). Indeed, there is both quadrupedal suspension and hindlimb suspension in dermopterans,

resulting in a tradeoff in the ability to walk quadrupedally upright (Fujiwara et al. 2011;

Granatosky 2018). In the PC1 and PC2 hindlimb morphospace, *G. variegatus* is the only glider

that overlaps with the flyer morphologies (Fig. 2b; Fig. S1b), reflecting a hindlimb adaptive

optimum potentially more easily attained than that of flyers' specialized forelimbs (Fig. 5).

Our modeling approach offers a first step towards support for the hypothesis that

selection may more strongly "pull" certain limb traits towards a flyer adaptive zone. In

particular, gliders' intermediate forelimb length and overlapping hindlimb width optima between

that of arborealists and flyers could be the traits driving lineages across fitness valleys due to

their greater effect in, for example, the functionality of the gliding/flying apparatus. However, to

complete the tentative adaptive landscape presented here (Fig. 5), these results could be

strengthened by more comprehensive coverage of the lengths/widths across more mammal

species and biomechanical studies tying these traits to specific functions and selective pressures.

Conclusions

Our findings complement earlier work investigating the evolution of bat powered flight

by testing the selective regimes acting on bone dimensions across the mammalian forelimb and

hindlimb. Our results reveal that the relatively lengthened forelimbs in gliders are intermediate to

those of bats and arborealists, whereas bats and gliders exhibit similarly relatively narrower

hindlimbs compared to arborealists. Thus, solely based on morphometrics, the traditional highly

specialized view of glider/bat forelimbs and the "reduced" view of bat hindlimbs may represent

more malleable adaptive zones than previously thought. There is a need for future studies to test

the biomechanical consequences of such trait optima alongside more synergistic studies taking

into account bat and gliders' complex physiology and development to test these hypotheses

regarding the evolution of mammalian powered flight.

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Figure Captions

Fig. 1. Morphological measurements and phylogeny. We conducted analyses based on this a)

phylogeny (Upham et al. 2019) across b) 14 forelimb linear measurements for each specimen of

scapula height and length (sh, sl), humerus length (hl), distal width (hdw), proximal width (hpw),

and mid-shaft width (hsw), radius length (rl), ulna length and olecranon length (ul, uol), and third

manual digit metacarpal length and width (mcl, mcw), proximal phalanx length and width (ppl,

ppw), and intermediate phalanx length (ipl) and we collected c) 15 hind limb measurements

consisted of pelvis, ilium, and ischium length (pel, il, isl); femur length (fl), distal width (fdw),

and midshaft width (fsw); tibia length (tl), proximal length (tpw), and midshaft width (tmw);

fibula length (fbl), and third pedal digit metatarsal length and width (mtl, mtw), proximal

phalanx length and width (ppxl, ppxw), and intermediate phalanx length (ipxl). We measured

only the third digit in this dataset (and not, for example, the fifth even though that might be of

biological interest) in order to build on previous datasets in the literature to generate a large

enough dataset for evolutionary modeling.

Fig. 2. Forelimb and hind limb PCAs. Morphospaces consist of PC1 and PC2 of 14 forelimb

traits (left panels) and 15 hind limb traits (right panels). The extant+extinct morphospace is

shown for the a) forelimbs and b) hind limbs. The extant-only morphospace of c) the

mvOUMloc3b optima (large icons) overlaid on the forelimb morphospace and the d) the

mvOUMloc4 optima (large icons) overlaid on the hind limb morphospace. Cartoons of a e)

quadrupedal mammal skeleton (Rattus norvegicus) versus a f) bat skeleton (Pipistrellus abramus

are shown where dark gray shaded bones on the cartoon skeletons correspond to those measured

to generate the morphospace.

Fig. 3. Forelimb trait optima from model-fitting of univariate traits. Forelimb OUM optima (θ) as

a size-corrected trait value (log-shape ratio) with 95% confidence intervals (Tables S3a) for the

best-fitting models (ΔAIC = 0) for 10 traits. The AICcW of the model is on top of each panel

(Table S2a).

Fig. 4. Hind limb trait optima from model-fitting of univariate traits. Hind limb OUM optima (θ)

as a size-corrected trait value (log-shape ratio) with 95% confidence intervals (Tables S3b) for

the best-fitting models (ΔAIC = 0) for 10 traits. The AICcW of the model is on top of each panel

(Table S2b).

Fig. 5. Hypothetical adaptive landscapes of forelimb and hind limb skeletal relative length and

width evolution. Hypothetical fitness peaks are shown for arborealists, gliders, and bats for a

broad summary of our conclusions from the skeletal trait optima results discussed. The "length"

and "width" in the figure refer to the lengths and widths of individual limb elements. For the a)

forelimb, we depict arborealists on a relatively steep adaptive peak at relatively lower

length/width optima, gliders on a broad, shallow peak/incline at relatively higher length but not

width optima, and bats on a very steep peak at relatively higher length and width optima. For the

b) hind limb, we depict arborealists again on a fairly steep adaptive peak at relatively higher

width but not length optima, bats on a shallow steep peak at relatively higher length and lower

width optima, and gliders on a peak similar to and overlapping with bats' at relatively higher

length but similar width optima.

| | ground-dwellers | | arborealists | | arborealists+gliders | | gliders | | extant bats | | extinct bats |

armwing length traits

a) OUM$_{loc3b}$ AICcW = 0.49
scapula length optima

b) OUM$_{loc4}$ AICcW = 0.92
humerus length optima

c) OUM$_{loc4}$ AICcW = 1.00
radius length optima

handwing length traits

d) OUM$_{loc3c}$ AICcW = 0.63
metacarpal length optima

e) OUM$_{loc4}$ AICcW = 0.99
int. phalanx length optima

f) OUM$_{loc4}$ AICcW = 0.84
proximal phalanx length optima

armwing width traits

g) OUM$_{loc3b}$ AICcW = 0.40
humerus proximal width optima

h) OUM$_{loc3b}$ AICcW = 0.65
humerus shaft width optima

i) OUM$_{loc3b}$ AICcW = 0.36
humerus distal width optima

handwing width trait

j) OUM$_{loc3b}$ AICcW = 0.44
metacarpal width optima

armwing   handwing

ground-dwellers   ground & arb.   arborealists   arb. & gliders   gliders   gliders & bats   bats

hind limb length traits

hind limb width traits

Legend:
- ground-dwellers
- ground & arb.
- arborealists
- arb. & gliders
- gliders
- gliders & bats
- bats

a)

summation of optima

width

length

b)

summation of optima

width

length

Table 1: AICcW scores from multivariate model-fitting analyses (via mvMORPH).
Comparisons of the fits of multivariate models to the first three principal components of the a)
forelimb dataset and b) hindlimb dataset.

a) forelimb	AIC	ΔAICc	mAICcW	b) hindlimb	AIC	ΔAICc	mAICcW
mvBM1	-94.42	225.50	0.00	mvBM1	-134.64	230.05	0.00
mvOU1	-248.73	71.20	0.00	mvOU1	-330.72	33.97	0.00
mvOUM$_{loc3a}$	-306.82	13.10	0.00	mvOUM$_{loc3a}$	-358.41	6.28	0.04
mvOUM$_{loc3b}$	**-319.93**	**0.00**	**0.99**	mvOUM$_{loc3b}$	-353.51	11.17	0.00
mvOUM$_{loc3c}$	-308.77	11.16	0.00	mvOUM$_{loc3c}$	-358.10	6.59	0.03
mvOUM$_{loc4}$	-302.35	17.57	0.00	**mvOUM$_{loc4}$**	**-364.69**	**0.00**	**0.92**