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# THE ROYAL SOCIETY

## **Physiology**

# Extreme and variable torpor among highelevation Andean hummingbird species

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Torpor is thought to be particularly important for small endotherms occupying cold environments and with limited fat reserves to fuel metabolism, yet among birds deep torpor is both rare and variable in extent. We investigated torpor in hummingbirds at approximately 3800 m.a.s.l. in the tropical Andes by monitoring body temperature ( $T_{\rm b}$ ) in 26 individuals of six species held captive overnight and experiencing natural air temperature ( $T_{\rm a}$ ) patterns. All species used pronounced torpor, with one *Metallura phoebe* reaching a minimum  $T_{\rm b}$  of 3.26°C, the lowest yet reported for any bird or non-hibernating mammal. The extent and duration of torpor varied among species, with overnight body mass ( $M_{\rm b}$ ) loss negatively correlated with both minimum  $T_{\rm b}$  and bout duration. We found a significant phylogenetic signal for minimum  $T_{\rm b}$  and overnight  $M_{\rm b}$  loss, consistent with evolutionarily conserved thermoregulatory traits. Our findings suggest deep torpor is routine for high Andean hummingbirds, but evolved species differences affect its depth.

### 1. Introduction

Hummingbirds (Apodiformes: Trochilidae) occupy elevations up to approximately 5000 m.a.s.l. in the Andes Mountains, providing one of the most spectacular examples of avian adaptation to extreme environments. The challenges of living in these cold, wet and hypoxic environments are compounded by hummingbirds being among the smallest of endotherms and possessing the highest mass-specific metabolic rates of any vertebrates [1–3]. Pronounced thermoregulatory costs are combined with very high costs of hovering flight at high elevations [4,5] and a diet of flower nectar requiring daily intake rates sometimes exceeding hummingbirds' own body masses ( $M_b$ ) [6,7]. These energetic challenges have focused long-standing interest in physiological and behavioural processes that facilitate hummingbird occupancy of high elevations [8–10]. The major hummingbird clades vary in the extent to which they have occupied montane and cold regions [11,12], but the basis for these evolutionarily conserved environmental niches is not fully understood.

A suite of physiological and behavioural adaptations facilitates hummingbird occupation of high elevations. In addition to roosting in thermally buffered caves and bouts of intense feeding before dark to maximize fat reserves for overnight metabolism, hummingbirds in the high Andes are thought to make extensive

use of nocturnal torpor [8,13]. Torpor, or daily heterothermy, is characterized by facultative hypometabolism and reductions of body temperature (T<sub>b</sub>), typically by 10–30°C below normothermic values that, unlike hibernation, are restricted to a single circadian cycle [14,15]. Torpor is widespread among hummingbirds [13,16-19], with variation in frequency or depth attributed to factors including nutritional status [20], migratory status [21], weather [19], typical thermal environment [18], seasonal acclimatization [22] and foraging behaviour [23]. However, the role of phylogenetic structure as a source of interspecific variation in setpoint T<sub>b</sub> and related variables among co-occurring species has received little attention.

We investigated torpor in six hummingbird species experiencing natural cycles of air temperature (Ta) at 3800 m.a.s.l. in the Peruvian Andes, with the goal of quantifying interspecific variation among free-ranging populations with different evolutionary histories. We tested four predictions: first, all species in a high-elevation community routinely use torpor at night, with torpid  $T_b$  closely approaching  $T_a$ ; second, overnight  $M_b$  losses are directly related to torpor bout duration, with longer bouts associated with smaller overnight  $M_b$  losses [24]; third, variation in torpor  $T_b$  and overnight  $M_{\rm b}$  loss is at least partly explained by phylogeny; and fourth, lower T<sub>b</sub> and longer torpor bouts characterize species in the 'coquette' clade [11], a group particularly diverse and abundant in high-elevation, cold habitats.

### 2. Methods

Detailed methods are presented in the electronic supplementary material. In brief, we caught hummingbirds representing six species (figure 1) between 7 and 18 March 2015 at Bosque Japani, Peru (approx. 3800 m.a.s.l.; S11° 39′ 41″ W76° 26′ 48″). Night length this time of year (around the autumnal equinox) was approximately 12 h. After capture in mist nets, birds were temporarily held in tents adapted as aviaries. Food was withheld from 30 min before dark, at which time birds were transferred into individual roosting enclosures for overnight measurements of cloacal T<sub>b</sub> using 36-gauge Teflon-coated thermocouples, inserted 1–2 cm and secured to retrices using small pieces of laboratory tape. Total  $M_{\rm b}$  loss was taken as the difference between evening and morning measurements and bout duration as the period with  $T_b$ less than 30°C, a value often, albeit somewhat arbitrarily, used in studies of avian heterothermy [e.g. 25,26].

We analysed effects of bout duration on minimum  $T_b$  and  $M_b$ loss using generalized linear multilevel models (GLMMs) using the brms [27,28] and stan [29] packages in R [30]. We estimated the phylogenetic signal by calculating Pagel's  $\lambda$  and Bloomberg's K and quantified phylogenetic signal from GLMMs by estimating the proportion of total variance attributed to phylogeny or species random effects. In addition, we visualized minimum  $T_{\rm b}$  and overnight  $M_b$  loss across the phylogeny of our study species using a published hummingbird phylogeny [11] and the contMap() function in the R package phytools [31]. Detailed analytical methods and comparisons of alternative statistical models are provided in the electronic supplementary materials.

### 3. Results

All six species and 24 of 26 individual hummingbirds entered torpor, but bout duration and minimum T<sub>b</sub> varied within and among species (figures 1 and 2). Normothermic  $T_b$  in individuals that remained normothermic for part or all of a night varied from 35.8°C in P. gigas to 37.0°C in A. cupripennis (figure 1). Night-time  $T_a$  minima remained between 2.4°C and 5.9°C throughout the study.

The gradient between minimum  $T_b$  and  $T_a$  varied among species; for instance, Colibri coruscans appeared to defend a setpoint of approximately 8°C, whereas Metallura phoebe thermoconformed over the entire  $T_a$  range (figure 2). The mean minimum  $T_b$  of M. phoebe was  $5.13 \pm 1.18$ °C, with individual minima on the coldest nights of 3.80°C and 3.26°C. Moreover, M. phoebe was the only species with no indication of defending a  $T_b$  setpoint, maintaining  $T_b - T_a$  gradients of just  $0.87 \pm 0.53$ °C (figure 2). The  $T_{\rm b}$  of Oreotrochilus melanogaster tracked Ta closely at Ta greater than 3.7°C but increased to 2-4°C above  $T_a$  at lower  $T_a$  values (figure 2). Maximum cooling rates during torpor entry were approximately 0.6°C min<sup>-1</sup> in four species and peak rewarming rates ranged from approximately 1°C min-1 in P. gigas to approximately 1.5°C min<sup>-1</sup> in A. cupripennis (figure 1). Hummingbirds generally rewarmed while  $T_a$  was low and stable, but in a few instances 'hitch-hiked' increasing  $T_a$ and thereafter warmed endogenously (e.g. figure 2 – P. gigas).

Bout duration varied from 2.3 h in one P. gigas to 12.9 h in a M. phoebe (figure 2) with species means of 5.7–10.6 h (figure 1). In all models, minimum  $T_{\rm b}$  and overnight mass loss were negatively related to bout duration (table 1, figure 2). Among models of minimum  $T_b$ , but not models of overnight  $M_b$  loss, incorporating a species random effect, phylogenetic random effect or both improved fit compared to models with no random effect (table 1).

Phylogenetic signal was greater for minimum T<sub>b</sub> (Pagel's  $\lambda = 0.620$  (95% highest posterior density (HPD) 0.074–0.998); Bloomberg's K = 1.643, p = 0.007) than overnight  $M_b$  loss (Pagel's  $\lambda = 0.562$  (95% HPD 0.055–0.999); Bloomberg's K = 1.223, p = 0.048). Phylogenetic signal was important for all GLMMs with phylogenetic random effects, and 95% HPD did not overlap zero (table 1). Species random effects were also important, with 95% HPD not overlapping zero (table 1). Furthermore, both phylogeny and species explained a considerable proportion of total variation when included in models (tables 1, electronic supplementary material, tables S1 and S2).

### 4. Discussion

Frequent use of torpor and accompanying low  $T_{\rm b}$  values support our prediction that heterothermy is a routine component of thermoregulation in high-elevation hummingbirds. Although torpor use is responsive to proximate organismal and environmental variables [18-23], the significant phylogenetic signal in minimum  $T_b$  and overnight  $M_b$  loss reveals that phylogenetically conserved evolution explains significant portions of variation in torpor performance among our study species. In particular, the tendency for lower  $T_b$  and longer torpor bouts among species in the coquette clade (O. melanogaster, P. caroli, M. phoebe), together with traits such as haemoglobin oxygen-binding affinity [10], may help to explain the over-representation of this clade in high-elevation Andean assemblages.

The minimum torpor  $T_b$  of O. melanogaster and M. phoebe during torpor are the lowest yet documented in hummingbirds; Calder and Booser [19] recorded a temperature of 6.5°C in an artificial egg under an incubating female Selasphorus platycercus at 2900 m.a.s.l., and Carpenter [13] documented cloacal T<sub>b</sub> of approximately 6.5°C (5.0°C in one

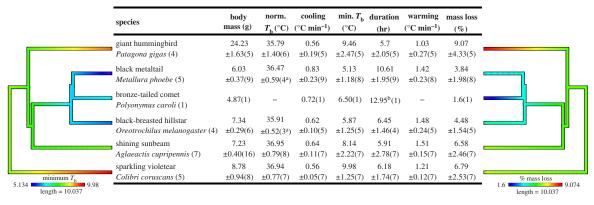
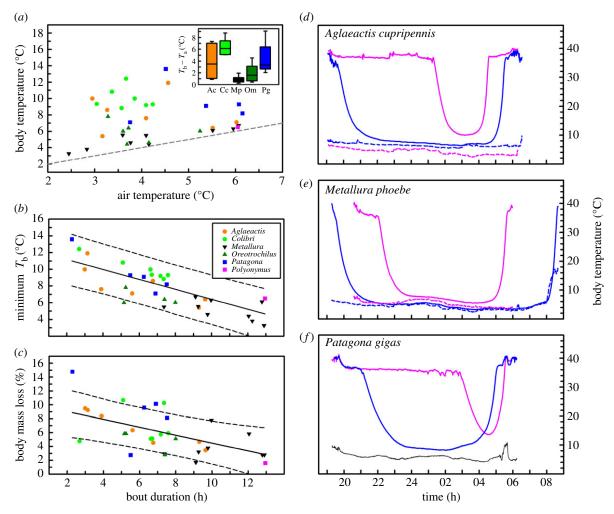


Figure 1. Torpor-related parameters for hummingbirds at approximately 3,800 m.a.s.l. in the Peruvian Andes: normothermic body temperature (norm. T<sub>b</sub>), maximum cooling rate during torpor entry, minimum torpor body temperature (min.  $I_h$ ), bout duration and maximum rewarming rate during arousal. Values are means  $\pm$  s.d., with sample sizes in parentheses. Phylogenetic reconstructions of minimum  $T_b$  and overnight body mass loss are at left and right, respectively. Superscripts: a = fewerdata because some individuals entered torpor immediately after thermocouple insertion, and dislodged thermocouple upon rewarming; b = did not rewarm until placed in sun.



**Figure 2.** Relationships between torpor variables among six species of hummingbirds at 3,800 m.a.s.l in the Peruvian Andes (a,b,c), and traces of body temperature  $(T_h)$  illustrating individual variation in bout duration (d,e,f). Minimum body temperatures  $(T_h)$  varied among species (a; dashed line indicates equality) and the gradient between minimum  $T_b$  and  $T_a$  (inset) varied significantly. Minimum  $T_b$  (b) and overnight body mass loss (c) were significantly related to bout duration, defined as the period with  $T_h$  less than 30°C. Solid lines are best-fit models (table 1), and dashed lines 95% highest posterior density intervals. In d, e and f, the solid pink and blue lines show  $T_b$  during the shortest and longest bouts, respectively, for each species. Dashed lines show corresponding  $T_a$  (both P. gigas traces obtained on the same night).

individual) in O. estella. In the present study, M. phoebe showed no evidence of maintaining a setpoint  $T_b$  at even the lowest  $T_a$ encountered (figure 2), raising the possibility that it may reach even lower  $T_b$  during colder conditions.

The minimum  $T_b$  values of 3.3°C and 3.8°C in two M. phoebe individuals are, to the best of our knowledge, the lowest yet recorded among birds. In free-ranging common poorwills (*Phalaenoptilus nuttallii*), minimum  $T_b = 4.3$ °C was

**Table 1.** Comparison of generalized linear multilevel models of minimum body temperature ( $T_b$ ) and percent overnight body mass ( $M_b$ ) loss. Models varied in whether they incorporated species, phylogenetic, both or no random effects. Estimated effect sizes and 95% highest posterior density (HPD) are provided. Proportion of variance explained and 95% HPD are indicated for models with species and phylogenetic random effects. Model fit was assessed using leave-one-out cross-validation (LOOIC). The difference between each model and the best-fit model is shown as  $\Delta$ elpd (expected log predictive density) with standard error (s.e.). The structure of the full models are M in  $T_b \sim bout$  duration + species + phylogeny and  $M_b$  loss  $\sim bout$  duration + species + phylogeny.

response	fixed effect bout duration	random effects		% variance explained		
		species	phylogeny	species	phylogeny	$\Delta$ elpd (s.e.)
Min. $T_{\rm b}$	-0.60 (-0.80, -0.40)	1.54 (0.08, 4.75)	0.39 (0.02, 1.29)	0.51 (0.004, 0.95)	0.12 (0.0002, 0.57)	0
Min. T <sub>b</sub>	-0.58 (-0.78, -0.38)	<del></del>	0.47 (0.15, 1.24)	_	0.15 (0.01, 0.53)	-0.1 (0.3)
Min. T <sub>b</sub>	$-0.60 \ (-0.80, \ -0.41)$	1.78 (0.65, 4.24)	<del></del>	0.63 (0.21, 0.94)	<u> </u>	-0.1 (0.3)
Min. T <sub>b</sub>	-0.66 (-0.85, -0.47)	<del></del>	<del></del>	_	<u> </u>	<b>-7.1</b> (3.5)
$M_{\rm b}$ loss	-0.64 (-0.96, -0.32)	<del></del>	<del></del>	_	<del>-</del>	0
$M_{\rm b}$ loss	-0.60 (-0.97, -0.23)	1.18 (0.04, 3.85)	<del></del>	0.19 (0.0002, 0.69)	_	-1.0 (1.9)
$M_{\rm b}$ loss	-0.57 (-0.95, -0.19)	<del></del>	0.32 (0.01, 1.05)	_	0.03 (0.00002, 0.15)	-1.1 (2.1)
$M_{\rm b}$ loss	-0.56 (-0.96, -0.14)	1.32 (0.05, 4.60)	0.36 (0.01, 1.27)	0.21 (0.0003, 0.78)	0.03 (0.00002, 0.21)	-1.6 (2.6)

inferred from a skin temperature ( $T_{\rm skin}$ ) datum of 2.8°C [25], with similar values reported more recently [32]. Moreover, the  $T_{\rm b}$  minima for M. phoebe appears to be the lowest reported for any avian or mammalian daily heterotherm, with  $T_{\rm b}$  less than 5°C otherwise restricted to hibernators [15].

Hummingbirds rewarmed from deep torpor surprisingly rapidly, with the maximum observed rate for *P gigas* equivalent to 168% of the value reported under laboratory conditions [33]. Observed maximum rates for the smaller species were equivalent to 163–194% of allometrically expected values [34], consistent with hummingbirds' metabolic rates while rewarming approaching those during hovering flight [35]. Rapid rewarming may maximize the time spent in deep torpor before commencing foraging [24].

Our data supported the prediction that energy expenditure is directly related to time spent torpid, with overnight  $M_{\rm b}$  loss negatively related to bout duration. Similar findings were reported for three Brazilian lowland species [18]. Rates of overnight  $M_{\rm b}$  loss for our study species were comparable to those reported by Bech *et al.* [18], despite the much colder environment of the present study. Both hummingbird communities achieved similar overnight energy savings despite differences in  $T_{\rm b}$  and  $T_{\rm a}$  of approximately 20°C, likely reflecting greater costs of rewarming under colder conditions.

The relationship between torpor bout duration and minimum  $T_b$  we observed likely reflects how costs of rewarming constrain overall energy savings. The negative, approximately linear effect of torpor depth on rewarming costs [34] combined with the nonlinear, Arrhenius effect on metabolic rate while thermoconforming [36], leads to the prediction that energy savings are maximized when bout duration increases with decreasing torpor  $T_b$ . Our results are consistent with recent findings that bout duration is the primary determinant of energy savings during overnight torpor in hummingbirds [24].

Individuals in our study fasted for just 30 min before dark but entered torpor routinely, suggesting that torpor use is less tightly coupled to individuals' energy reserves as often reported for hummingbirds in other environments [18,22,37]. However, several authors have documented intense feeding immediately before dark [38,39] and the extent to which torpor in high Andean hummingbirds is a routine component of thermoregulation or an 'emergency' response (e.g. [20]) requires further investigation.

In conclusion, we found that tropical hummingbird species living at elevations approaching 4,000 m.a.s.l. have evolved pronounced, but variable, capacities for torpor, with minimum  $T_b$  rivalling that of temperate- and boreal-latitude mammalian hibernators. Although avian hibernation (i.e. multi-day torpor) has been reported only in one caprimulgid [32,40], the depth of overnight torpor we document here raises the possibility that some high-elevation hummingbirds may hibernate during periods of inclement weather. Regardless, the energy savings associated with pronounced torpor are one of the major reasons why these tiny birds can persist in these harsh, physiologically challenging environments. Our finding that phylogenetic relationships are linked to torpor energy savings among co-occurring species suggests that differential evolutionary colonization of mountains [11,12] may have resulted from deeply conserved physiological differences among hummingbird clades.

Ethics. All procedures were conducted under permits 0280-2014-MIN-AGRI-DGFFS/DEGFFS and 405-2017-SERFOR/DGGSPFFS and University of New Mexico Institutional Animal Care and Use Committee protocols 14-101168-MC and 16-200418-MC. Ethical approval was also obtained from the Animal Ethics Committee of the University of Pretoria (NAS473/2019) and Research Ethics and Scientific Committee of the South African National Biodiversity Institute (P19/27).

Data accessibility. Data are available from the Dryad Digital Repository at: https://doi.org/10.5061/dryad.vx0k6djp6 [41].

Authors' contributions. B.O.W. and C.C.W. designed the research. B.O.W., C.J.S., A.B.J. and C.C.W. collected the data, which were analysed by A.E.M., C.J.S. and Z.J.C. A.E.M. led the writing of the manuscript; C.J.S., Z.J.C., B.O.W., A.B.J. and C.C.W. contributed. All authors gave final approval for publication and agree to be held accountable for the work performed therein.

Competing interests. We declare we have no competing interests.

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