



## Metabolic rate, latitude and thermal stability of roosts, but not phylogeny, affect rewarming rates of bats



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### HIGHLIGHTS

- We examined ecological, behavioral and physiological drivers of rewarming rates of 45 bat species.
- After controlling for phylogeny, high basal metabolic rate was associated with rapid rewarming.
- Species that live at higher absolute latitudes, and in less thermally stable roosts, also rewarmed most rapidly.
- Results suggest some species rely on passive means to reduce costs of rewarming, but others rely on faster metabolism as an alternative.
- This reinforces the importance of local climate, physiology, and behaviour in the proliferation of heterothermic endotherms

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### ABSTRACT

Torpor is an adaptation that allows many endotherms to save energy by abandoning the energetic cost of maintaining elevated body temperatures. Although torpor reduces energy consumption, the metabolic heat production required to arouse from torpor is energetically expensive and can impact the overall cost of torpor. The rate at which rewarming occurs can impact the cost of arousal, therefore, factors influencing rewarming rates of heterothermic endotherms could have influenced the evolution of rewarming rates and overall energetic costs of arousal from torpor. Bats are a useful taxon for studies of ecological and behavioral correlates of rewarming rate because of the widespread expression of heterothermy and ecological diversity across the > 1200 known species. We used a comparative analysis of 45 bat species to test the hypothesis that ecological, behavioral, and physiological factors affect rewarming rates. We used basal metabolic rate (BMR) as an index of thermogenic capacity, and local climate (i.e., latitude of geographic range), roost stability and maximum colony size as ecological and behavioral predictors of rewarming rate. After controlling for phylogeny, high BMR was associated with rapid rewarming while species that live at higher absolute latitudes and in less thermally stable roosts also rewarmed most rapidly. These patterns suggests that some bat species rely on passive rewarming and social thermoregulation to reduce costs of rewarming, while others might rely on thermogenic capacity to maintain rapid rewarming rates in order to reduce energetic costs of arousal. Our results highlight species-specific traits associated with maintaining positive energy balance in a wide range of climates, while also providing insight into possible mechanisms underlying the evolution of heterothermy in endotherms.

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### 1. Introduction

Endothermic animals defend a constant, high body temperature ( $T_b$ ) via endogenous, metabolic heat production, which can allow for

sustained activity across a range of conditions. However, maintaining an elevated  $T_b$  is energetically expensive, especially during cold weather, and can become challenging during periods of low resource availability [1]. To offset these high energy demands, many endotherms employ facultative heterothermy or torpor [2]. Torpor is characterized by a controlled reduction of  $T_b$ , metabolic rate (MR), and other physiological functions, greatly reducing energy consumption [2]. Short bouts of torpor during certain parts of the day (i.e., daily torpor) and/or longer bouts at specific times of year (i.e., seasonal torpor or hibernation) allow endotherms to maintain positive energy balance across a range of environmental conditions and timescales, and can also result in other ecological benefits (reviewed by Geiser and Brigham [3]).

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Bouts of torpor can be separated into three phases: 1) the cooling phase, during which an individual decreases  $T_b$  and MR to a new, lower set-point, 2) the torpid phase, during which the individual defends a reduced  $T_b$  set-point and torpid metabolic rate (TMR, which can be less than 1% of resting MR) [4], and, 3) the warming or arousal phase, during which the individual actively terminates the torpor bout and rewarms to normothermic  $T_b$ . The arousal phase is considered one of the major disadvantages of heterothermy because rearming from torpid to euthermic  $T_b$  is energetically expensive and requires extensive heat generation capacity [2,5]. High arousal costs negate potential energy savings of short torpor bouts, and therefore may represent an important factor in the cost-benefit tradeoff influencing torpor expression. Ultimately, these costs could represent a substantial portion of the long-term energy budget. Thus, mechanisms for mitigating costs of arousing from torpor are likely important drivers of the ecology, behavior, and life histories of heterothermic endotherms.

Somewhat counter-intuitively, rapid rearming is less energetically demanding than rearming slowly because the need to balance heat production with heat loss to the environment is less costly over shorter time intervals [6]. This predicts that selection should favor rapid rearming or, alternatively, other energy-saving mechanisms such as social thermoregulation or passive rearming to minimize arousal costs. If arousal from torpor depends on metabolic heat production, heterothermic species with greater thermogenic capacity (potentially higher basal and/or summit metabolic rate) should be capable of rearming more rapidly [7]. However, the evolutionary relationship between MR and rearming rate remains unclear. Geiser and Baudinette [7] found a nearly perfect correlation between BMR and rearming rate across all mammalian taxa but few subsequent analyses exist and, those studies that do have found equivocal patterns within a single taxon or group [8,9].

The lack of a consistent relationship between MR and rearming rate could also reflect the fact that rearming in many species is largely passive [10,11,12] and does not rely solely on metabolic heat production. For species employing passive strategies, the energetic costs associated with rearming may be heavily influenced by behavioral and environmental factors that allow individuals to take advantage of external heat sources for passive rearming [6,11,13]. The availability of solar radiation and fluctuations in ambient temperature ( $T_a$ ) are important determinants of the energetic cost of rearming [10,11,12,13]. Many heterothermic endotherms have limited exposure to sunlight and/or variation in natural  $T_a$  in their burrows, nests, or roosts (e.g., underground burrows, caves, well-insulated trees), minimizing opportunities for passive rearming. In addition, variation in the rate at which heterothermic endotherms rearm from torpor might also reflect local behavioral and physiological adaptation to different climatic regimes. Many other energetic and physiological traits vary along latitudinal gradients [14–17], including BMR, capacity for non-shivering thermogenesis, heterothermy index (i.e., continuous metric of heterothermy [18]) and thermoregulatory scope (i.e., range of body temperatures exhibited by a species, mean  $T_b$  – minimum  $T_b$  [17]). Species residing at higher latitudes demonstrate greater thermogenic capacity (i.e., higher MR [15] and greater non-shivering thermogenesis [16]) to cope with unpredictable, occasionally extreme changes in environmental conditions. Temperate heterotherms might also exhibit faster rearming rates to reduce costs of arousal when  $T_a$  is cold and access to passive heat sources is limited.

Social thermoregulation is another behavioral strategy that could influence rearming rate. Some heterotherms huddle with conspecifics to reduce heat loss and/or increase  $T_a$  of their immediate microclimate [19,20]. Huddling is common in species that hibernate, roost, or nest in enclosed burrows where opportunities for passive heating are reduced. If groups of individuals can share thermoregulatory costs or reduce heat loss by huddling, this could reduce selection pressure favoring rapid warming. On the other hand, solitary species or those

in smaller groups may depend on endogenous thermogenic capacity and rapid rearming to keep the cost of arousal low [6,21].

Bats are a useful model taxon for comparative studies of heterothermy and rearming. They are among the most ecologically diverse vertebrates and many species from both temperate and tropical regions employ daily and/or seasonal torpor, with some temperate species dependent on months of hibernation for over-winter survival [22]. Different bat species roost in a range of habitats, from caves and mines with high thermal stability, to tree hollows with moderate thermal stability to shedding bark or exposed foliage with high thermal variability. In addition, there is enormous variation in the potential for social thermoregulation among bats with some species roosting in large colonies of up to hundreds of thousands or millions of individuals and others roosting solitarily for most or all of their annual cycle [23]. This variation in ecology and behavior provides an excellent opportunity to test hypotheses about the evolution of rearming rates in heterothermic endotherms.

We used comparative analyses to assess how physiological, behavioral and environmental variation affect rearming rates of bats. First, we tested the hypothesis that metabolic rate is associated with rearming rate in bats and predicted that species with higher BMR would exhibit faster maximum rearming rates. Second, we tested the hypothesis that latitudinal variation in climate affects the evolution of rearming rates. We predicted that bat species living at higher, temperate latitudes might be capable of faster maximum rearming rates compared to low latitude species because of selective pressure imposed by colder, less predictable environmental conditions. Finally, we tested the hypothesis that roosting behavior affects the evolution of rearming rates and predicted that species living in smaller groups and in less thermally stable roosts would exhibit the highest rearming rates because these species have limited opportunities for passive rearming and, therefore, should have faced selection pressure favoring rapid rearming.

## 2. Methods

### 2.1. Field measurements of rearming rate

We collected data for silver-haired (*Lasiorycteris noctivagans*) and northern long-eared bats (*Myotis septentrionalis*) at the Sandilands Forest Discovery Centre in Manitoba, Canada (49.67°N, 95.90°W), from July 27th to August 1st, 2012 and 2014. Bats were captured using mist nets (12 m by 6 m), held in cloth bags and transported less than 1 km on foot to a nearby field laboratory. Here, bats were kept in their holding bags in a quiet room with screen-windows, under natural photoperiod (16 h light, 8 h dark) and natural  $T_a$  (minimum 11 °C–maximum 33 °C) for up to 24 h. At least two times a day, bats were provided with water, using a disposable pipette but were not fed because captivity was so brief.

As part of a concurrent respirometry study, every night up to four bats were placed in 100 mL transparent, acrylic chambers within a temperature-controlled cabinet set at 15 °C (i.e., well below the lower critical temperature) to encourage bats to enter torpor. The following day, between 13:00 and 22:00 individuals were removed from their chambers one at a time, and  $T_b$  was measured immediately ( $T_{b1}$ ), by inserting a lubricated, 1 mm diameter thermocouple approximately 3 mm into the rectum until the reading was stable. After the first measurement, the bat was placed in a handling bag for approximately 5 min to minimize handling stress, after which a second  $T_b$  measurement was recorded ( $T_{b2}$ ). Rewarming rate was calculated by subtracting  $T_{b1}$  from  $T_{b2}$  and dividing by the time interval between the two  $T_b$  measurements to give a rate in °C/min. Following  $T_b$  measurements, bats were weighed, provided water with a disposable pipette and released at the site of capture at dusk. These procedures and fieldwork were conducted under a Manitoba Conservation Wildlife Scientific Permit and were approved by the University of Winnipeg Animal Care Committee.

We used unpublished temperature radio-telemetry data for hoary bats (*Lasiurus cinereus*) and Eastern red bats (*Lasiurus borealis*) collected as part of a field study by McGuire et al. [24]. These data were not reported by McGuire et al. [24], but were collected at the same time, and using identical methods as their data for silver-haired bats. Briefly, bats were captured at Long Point, Ontario, Canada using mist nets. A small patch of fur was trimmed in the interscapular region and a temperature-sensitive radiotransmitter ( $\pm 0.1$  °C, 0.38 g, <4% of body mass; Pip3, Lotek Wireless, Newmarket, Ontario, Canada) was affixed with ostomy bonding cement (Torbot; Cranston, RI, USA). Each bat was immediately released and later tracked to the day roost, where a datalogging receiver (SRX400; Lotek Wireless) recorded  $T_{sk}$  every ~30 s throughout the day. We extracted arousal rate from telemetry data, using  $T_{sk}$  timecourses to visually identify periods of arousal from torpor. We then calculated maximum rewarming rate from the steepest portion of the  $T_{sk}$  trace.

## 2.2. Rewarming rate database

We reviewed the literature and tabulated rewarming rates for 41 bat species from previously published studies and combined this pre-

existing data with our own field measurements to create a database of 45 species (Table 1). When possible, we restricted our analysis to warming rates based on  $T_b$  instead of  $T_{sk}$ . However, a number of studies of bats have used external temperature-sensitive transmitters to record skin temperature ( $T_{sk}$ ) in the lab and field. Acknowledging there can be differences between absolute values of  $T_b$  and  $T_{sk}$  [see [24–26]], we argue that  $T_{sk}$  is an adequate proxy for  $T_b$  in the context of rewarming; Willis and Brigham [26] showed that, when bats are rewarming at their maximum rate, the slopes of the lines are virtually indistinguishable, especially at warmer temperatures (i.e., our temperature range of 18–22 °C). We performed a calibration on seven silver-haired bats and found a tight correlation between rewarming rates calculated using  $T_{sk}$  vs.  $T_b$  ( $R^2 = 0.86$ ,  $F_{1,6} = 36.3$ ,  $p < 0.001$ ). We used the maximum reported rewarming rate for each species or extracted the maximum rewarming rate from  $T_{sk}$  traces as described above [9]. Since both rewarming rates and differences between  $T_{sk}$  and  $T_b$  are affected by  $T_a$  [7,24,25,26], we only included data for bats recorded arousing from torpor when  $T_a$  was between 18 and 22 °C.

BMR values were available from the literature for 23 of the 45 species in our analysis (see Table 1). Body mass accounted for over 86% of

**Table 1**

Data used in our analysis including latitude representative of the geographic range, maximum colony size, stability of roost temperature on a three-point scale (i.e., roost score, see Methods section), body mass, basal metabolic rate (BMR) and rewarming rate for 45 bat species. Latitude was calculated based on centroid coordinate values (i.e., latitude and longitude) from geographic data obtained for each species' distribution. Colony size is the maximum colony size reported in the literature. Roost score represents the temperature stability of roosts on a three-point scale ranging from 0 (highly stable roosts; e.g., caves) to 2 (highly unstable roosts; e.g., foliage). Body mass values were taken from the original sources for the rewarming rate data.

Species	Latitude (°)	Colony Size	Roost Score	Mass (g)	BMR (mL O <sub>2</sub> /h)	Source for BMR	Rewarming Rate (°C/min)	Source for Rewarming Rate
<i>Antrozous pallidus</i>	33.69	139	2	28.0	21.20	[48]	1.00	[63]
<i>Asellia tridens</i>	22.70	5000	0	15.0	–	–	0.42	[64]
<i>Chaerephon pumila</i>	–0.29	300	2	18.0	–	–	1.05	[9]
<i>Chalinolobus gouldii</i>	–26.01	40	2	12.0	25.20	[49]	1.40	[64]
<i>Chalinolobus picatus</i>	–28.40	15	0	6.0	–	–	1.40	[64]
<i>Corynorhinus rafinesquii</i>	33.90	118	1	10.5	–	–	0.83	[9]
<i>Corynorhinus townsendii</i>	37.65	300	0	10.5	–	–	1.52	[63]
<i>Eptesicus fuscus</i>	38.50	700	2	16.0	17.00	[50]	1.50	[50]
<i>Eptesicus serotinus</i>	41.33	20	2	25.0	43.10	–	1.30	[65]
<i>Glossophaga soricina</i>	5.20	1000	1	9.5	21.60	[51]	0.35	[66]
<i>Hipposideros armiger</i>	22.45	1000	0	53.0	32.86	[52]	0.51	[52]
<i>Hipposideros speoris</i>	16.50	1200	1	9.0	–	–	0.35	[64]
<i>Lasionycteris noctivagans</i>	43.80	50	2	10.5	–	–	2.70	This study
<i>Lasiurus borealis</i>	38.51	1	2	13.0	–	–	1.78	This study
<i>Lasiurus cinereus</i>	27.79	1	2	25.0	–	–	2.14	This study
<i>Macroglossus minimus</i>	–1.67	1	2	16.0	21.00	[53]	0.31	[53]
<i>Miniopterus schreibersii</i>	39.61	10,000	0	15.0	25.42	[27]	1.40	[64]
<i>Mops condylurus</i>	–25.02	700	2	36.0	31.09	[54]	0.80	[64]
<i>Mormopterus loriae</i>	–16.10	300	2	8.5	–	–	0.90	[64]
<i>Mormopterus planiceps</i>	–30.00	150	2	9.5	–	–	1.40	[64]
<i>Myotis adversus</i>	4.12	300	2	8.0	–	–	1.20	[64]
<i>Myotis californicus</i>	35.96	52	2	5.0	–	–	1.29	[63]
<i>Myotis evotis</i>	43.85	30	2	7.3	–	–	0.67	[67]
<i>Myotis lucifugus</i>	49.73	3000	1	7.6	9.30	[55]	0.80	[68]
<i>Myotis myotis</i>	44.92	1000	1	25.0	25.00	[56]	1.50	[64]
<i>Myotis nattereri</i>	48.08	200	2	8.0	–	–	1.05	[64]
<i>Myotis septentrionalis</i>	47.15	60	1	8.2	–	[27]	1.99	This study
<i>Myotis thysanodes</i>	34.86	300	1	6.5	17.40	[57]	0.91	[63]
<i>Nyctalus noctula</i>	47.63	234	2	27.0	39.69	[27]	1.58	[64]
<i>Nyctimene albiventer</i>	–5.15	1	2	28.0	26.30	[58]	0.60	[69]
<i>Nyctophilus bifax</i>	–18.00	7	2	10.1	13.10	[59]	0.75	[70]
<i>Nyctophilus geoffroyi</i>	–26.20	20	2	7.8	10.99	[60]	1.34	[71]
<i>Nyctophilus gouldii</i>	–30.40	20	2	8.8	10.39	[61]	0.61	[72]
<i>Pipistrellus hesperus</i>	31.80	12	2	4.0	–	–	1.20	[73]
<i>Pipistrellus pipistrellus</i>	42.22	300	1	6.0	12.10	[27]	0.95	[64]
<i>Plecotus auritus</i>	51.09	30	1	12.0	11.20	[62]	0.70	[64]
<i>Rhinolophus ferrumequinum</i>	36.86	300	0	18.0	46.50	[27]	0.75	[64]
<i>Rhinolophus hipposideros</i>	40.39	100	1	6.0	–	–	1.05	[64]
<i>Rhinolophus megaphyllus</i>	–22.10	2000	0	8.0	25.84	[9]	0.84	[64]
<i>Rhinopoma hardwicki</i>	21.80	5000	0	10.0	–	–	0.50	[64]
<i>Rhinopoma microphyllum</i>	20.23	5000	0	19.0	–	–	0.30	[64]
<i>Tadarida brasiliensis</i>	2.83	200,000	0	12.0	15.30	[48]	0.73	[63]
<i>Tadarida teniotis</i>	38.03	100	2	30.0	31.52	[27]	0.78	[64]
<i>Taphozous australis</i>	17.30	100	0	22.0	–	–	0.57	[64]
<i>Taphozous melanopogon</i>	13.92	4000	0	26.0	–	–	0.15	[64]

the variation in BMR for these species ( $F_{1,21} = 21.7$ ,  $p = 0.0001$ ,  $r^2 = 0.86$ ). Therefore, to account for effects of body mass on BMR we used the residuals of this linear regression between body mass and BMR as a predictor variable in subsequent analyses (hereafter, mass-corrected BMR).

We assessed ecological and behavioral predictors of rewarming rate using three metrics: latitude, maximum colony size, and roost stability. We identified maximum colony size and roost preferences from the literature [9,23] or from the International Union for the Conservation of Nature (IUCN) Red List of Threatened Species (version 2015.2). Due to a lack of direct measurements of roost temperature in the literature, we classified species according to roost stability on a three-point, categorical scale [9]: 0 representing highly stable microclimates (e.g., caves and mines), 1 representing less thermally stable roosts (e.g., tree hollows and buildings) or species that roost in caves for part of their annual cycle but also switch to trees or other structures, and 2 representing thermally unstable roosts characterized by large fluctuations in  $T_a$  (e.g., foliage, shedding bark). We obtained geographic data through IUCN Red List of Threatened Species in the form of two-dimensional shapefiles (i.e., geospatial vector data of points on a map). We calculated the centroid coordinate (latitude, longitude) for each species-specific shapefile, and then calculated the weighted latitude (i.e., larger shapefile vectors were weighted more heavily) for all species in our dataset using the 'mapproj' package (version 0.8-29, [28]) in R (v3.1.1; [29]). An absolute value of latitude for each species was taken to provide a measure of distance from the equator without differentiating between northern and southern hemispheres. Body mass values for each species were obtained from the original sources of rewarming rate data and we used this variable as a predictor in subsequent analyses.

### 2.3. Analyses

All analyses were conducted in R [29]. Prior to analysis, we tested for normality and equality of variance for all variables and log<sub>10</sub> transformed variables when appropriate. Significance was assessed at  $\alpha = 0.05$ .

We used two analytical approaches to account for common ancestry. We first analyzed our dataset using a Phylogenetic Generalized Least-Squares model (PGLS; [30,31]) in R, using the 'ape' (v. 3.1.2; [32]) and 'caper' (v. 3.1.2; [33]) packages. We obtained data on the phylogenetic distances between bat species from the phylogenetic tree provided by Agnarsson et al. [34]. In PGLS models the value of the phylogenetic signal ( $\lambda$ ) was estimated using maximum likelihood to optimally adjust the degree of phylogenetic correlation in the data. PGLS models allow for flexibility of  $\lambda$ , which can range from 0 to 1, where 0 indicates no phylogenetic signal and 1 represents a very strong phylogenetic signal [35]. Values of 1 are similar to phylogenetic independent contrasts (PIC; [36]) and assume a 'Brownian motion' model where trait data are fully explained by phylogeny. Values of 0 are similar to ordinary least squares models (e.g., linear regression) and assume no phylogenetic signal. Intermediate values of  $\lambda$ , calculated using PGLS models, indicate that phylogeny is corrected for but do not follow a full 'Brownian motion' evolutionary model [37]. Therefore, the use of PGLS models and maximum-likelihood estimation of  $\lambda$  allowed us to optimize the phylogenetic correction for our dataset [37].

Initial PGLS analyses generated low  $\lambda$  values (0.00 to 0.40), indicating that phylogeny was not a strong predictor of rewarming rate [38]. Therefore, we conducted a second analysis using a series of non-phylogenetically-corrected general additive models (GAMs), to account for non-linear relationships between rewarming rate and colony size that we observed in our preliminary analysis. We included colony size (log<sub>10</sub> transformed) as a non-linear term in our models and absolute latitude, roost preference score and mass-corrected BMR residuals (log<sub>10</sub> transformed; due to negative values we added a constant to every residual value before transforming) as linear terms. We evaluated the relative support for each analysis using the Akaike Information Criteria

corrected for small sample sizes ( $AIC_c$ ). The model with the lowest  $AIC_c$  value was considered to be most appropriate for our data [39].

Since metabolic data were only available for a subset of the species, we conducted two separate sets of analyses: a global analysis with all 45 species, excluding BMR, as well as a partial analysis, including only the 23 species for which BMR data were available. The partial PGLS model included mass-corrected BMR, absolute latitude, colony size, and roost score as predictors of rewarming rate. The global analyses included body mass, absolute latitude, colony size, and roost score as predictors of rewarming rate. For each series of models we removed variables using model decomposition based on  $AIC_c$  values. All models within  $\Delta AIC_c \leq 2$  are reported, as they were considered to have substantial support [39]. In total, we compared results from 30 candidate models (15 for PGLS, 15 for GAM). We quantified  $AIC_c$  weights ( $w_i$ ), which sum to one across all models and represent the probability that each model provides the best fit compared to all other candidate models. From these  $AIC_c$  weights, we also calculated the variable weights (vw) of each predictor variable included in our models (i.e., mass-corrected BMR, mass, latitude, colony size, roost score) to assess the relative importance of each predictor (i.e., the predictor with the largest vw is estimated to be the most important) [39].

### 3. Results

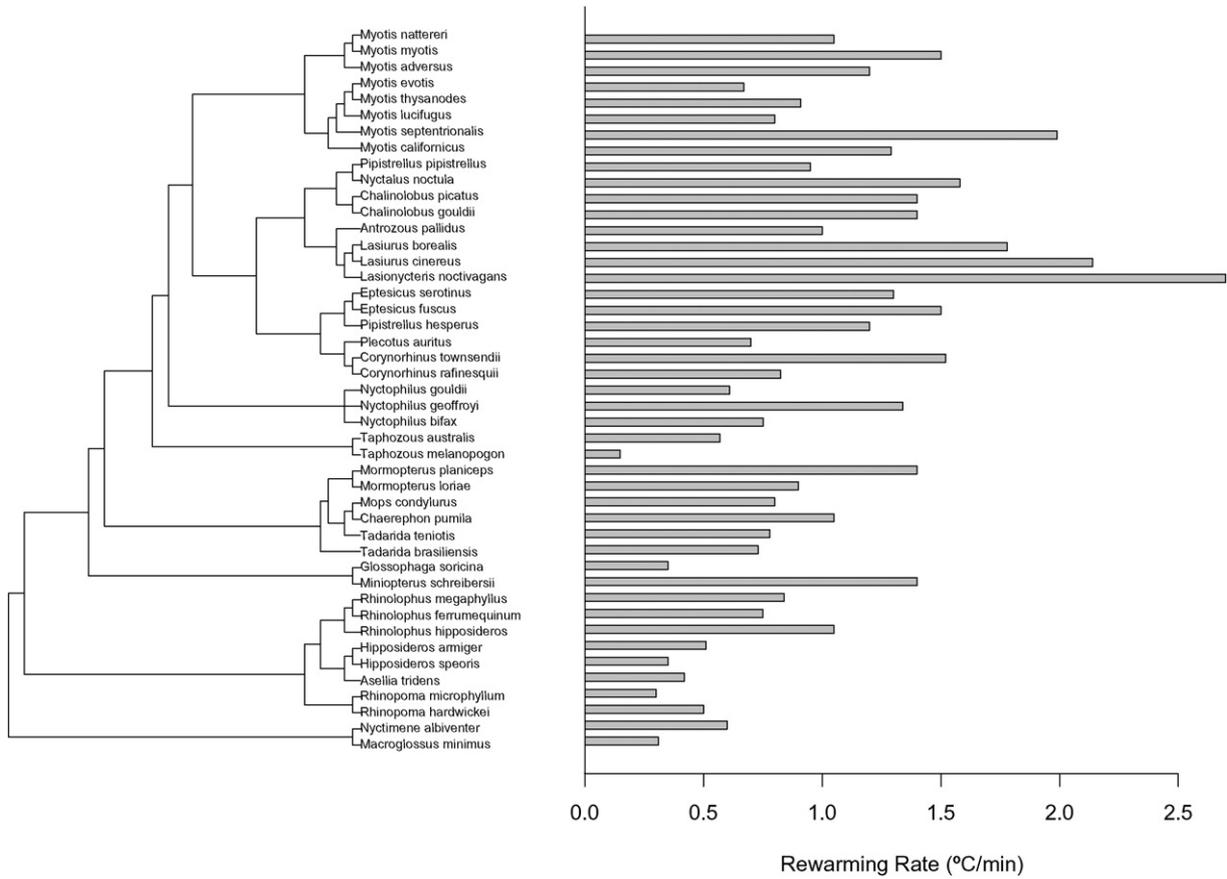
We found maximum rewarming rates for 41 bat species from eight families in the literature and added new data for four species based on our own measurements (Fig. 1). Mean maximum rewarming rate was  $1.02 \pm 0.53$  °C/min but was highly variable, ranging from 0.15 to 2.7 °C/min (Table 1). The most parsimonious global PGLS model included only latitude (Table 2), while other top models included latitude, roost score and mass, explaining between 8% and 32% of variation in rewarming rate. The most parsimonious partial PGLS model included all variables, explaining 66% of the variation in rewarming rate, while the other top model included BMR and latitude, explaining 47% of the variation in rewarming rate (Table 2).

In our full models, latitude appeared in all of the top PGLS models (vw = 0.88) and the top GAM model (vw = 0.99) (Table 2). Roost score was moderately important as a predictor variable as it appeared in almost all PGLS (vw = 0.44) and GAM models (vw = 0.79) (Table 2). Colony size did not appear in any of the top global models, indicating it was not a strong predictor of rewarming rate. Mass appeared in three of the reported models (vw = 0.36 for PGLS, vw = 0.30 for GAM) (Table 2).

In our partial PGLS models, BMR and latitude appeared in both of the top models (vw = 0.55, 0.71 respectively) and were positively correlated with rewarming rate (Figs. 2, 3). Colony size (vw = 0.53) and roost score (vw = 0.53) were moderately important predictors, appearing in the top model (Table 2).

### 4. Discussion

Our results suggest that physiology, behavior and climate have influenced the evolution of rewarming rates in bats. In contrast to an earlier, smaller analysis on bats [9], but consistent with trends across mammals in general [7], species with high BMR (i.e., potentially greater capacity for metabolic heat production) had relatively high rewarming rates (as seen in [7]). Species at higher latitudes (i.e., in temperate regions) also tended to exhibit higher rewarming rates, possibly as a result of climate-mediated selection. Species roosting in thermally unstable roost sites had higher rewarming rates than those roosting in stable microclimates. Finally, we observed a weak correlation between colony size and rewarming rate, with species roosting in smaller colonies exhibiting higher rewarming rates, suggesting a potential co-evolutionary relationship between social behavior and thermoregulatory energetics of torpor. Taken together these results suggest that, for heterothermic endotherms, the ability to maintain positive energy balance has been



**Fig. 1.** Phylogenetic tree (left; adapted from [34]) used in the Phylogenetic Least Squares (PGLS) analysis alongside the distribution of rewarming rate values (right) for the 45 bat species used in our analyses.

shaped by thermogenic capacity, adaptation to local climate and various behavioral mechanisms.

Consistent with our first hypothesis, we identified a correlation between rewarming rate and BMR for 23 species, suggesting that heat production during rewarming from torpor is related to metabolic rate [7,8] (Fig. 2). Although BMR is a metric of “maintenance” costs and not necessarily representative of thermogenic capacity, rewarming from torpor requires intense metabolic output and BMR does correlate with overall metabolic performance [7,40]. Rewarming rates could also be strongly influenced by other metrics of metabolism (e.g., summit metabolism)

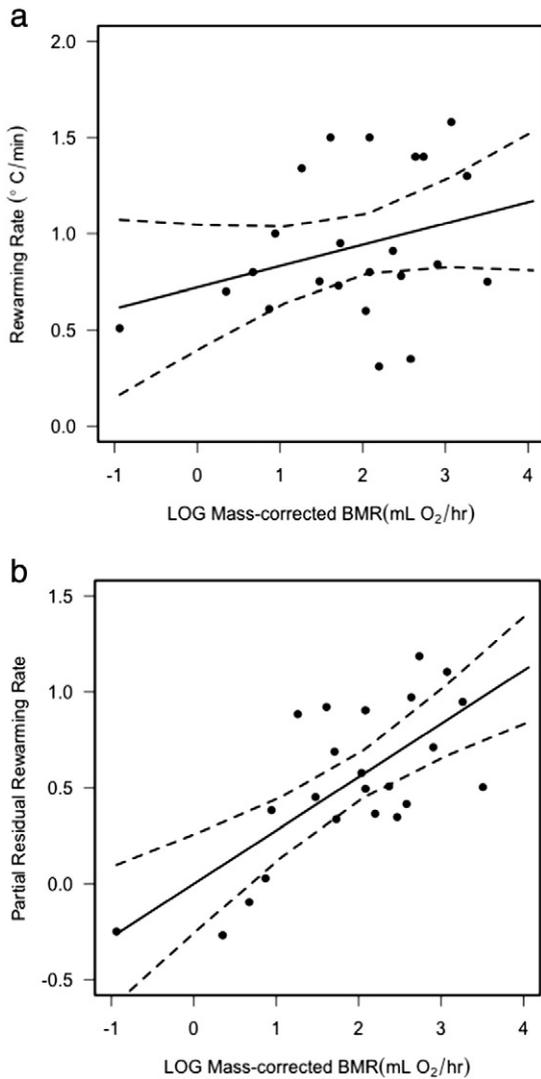
**Table 2**

Model selection parameters (AICc = Akaike’s Information Criterion corrected for small sample size;  $w_i$  = Aikaike weight) of (A) global and (B) partial Phylogenetic Least Squares models (PGLS) and (C) global General Additive Models (GAM), testing the effects of body mass, absolute latitude, colony size, roost stability and mass-corrected basal metabolic rate (BMR) on rewarming rate. Models listed are those considered the most parsimonious ( $\Delta AICc$  values <2). In PGLS models,  $\lambda$  is an indicator of the phylogenetic correlation in our data, and can range from 0 (no phylogenetic signal) to 1 (very strong phylogenetic signal).

(A) Global PGLS Models	$\lambda$	R <sup>2</sup>	AICc	$\Delta AICc$	$w_i$
Latitude	0.40	0.08	56.1	0.0	0.32
Latitude + Roost Score	0.00	0.31	57.1	1.0	0.20
Latitude + Roost Score + Mass	0.00	0.32	58.1	2.0	0.12
Latitude + Mass	0.39	0.07	58.1	2.0	0.12
(B) Partial PGLS Models					
BMR + Latitude + Roost Score + Colony	0.00	0.66	10.94	0.0	0.34
BMR + Latitude	0.00	0.47	11.27	0.3	0.29
(C) Global GAM Models					
Latitude + Roost Score	–	0.31	63.3	0.0	0.45
Latitude + Roost Score + Mass	–	0.30	64.9	1.6	0.20

or thermoregulatory parameters (e.g., thermal conductance, which is also strongly correlated to BMR [41]). However, due to a lack of published data on these parameters for bats we were unable to include them in our analysis. As more data become available, incorporating thermal conductance and other measures of thermogenic capacity (e.g., summit metabolism) could be useful for further resolving the relationship between heat generation capacity and rewarming.

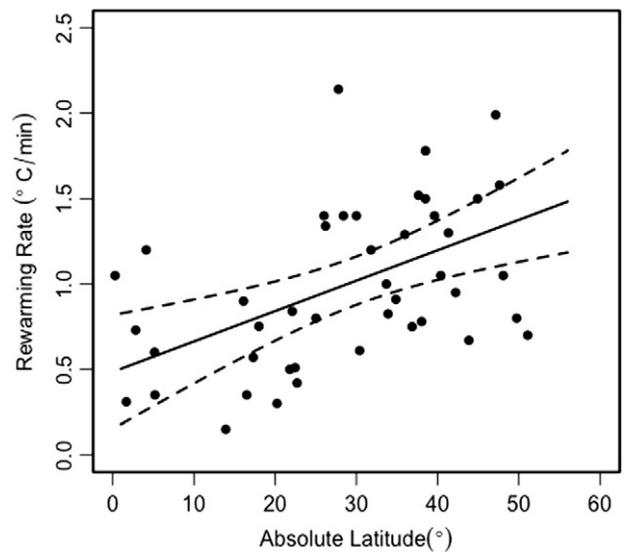
Consistent with our second hypothesis, we identified a positive relationship between latitude and rewarming rate (Fig. 3). Our results suggest that variation in rewarming rates may reflect an adaptation to regional climate, highlighting the importance of seasonality and temperature as selective forces on both thermogenesis and behavior in small mammals [14–17]. A handful of physiological traits associated with thermogenesis are known to vary along latitudinal gradients. For example, species that evolved in cooler, more seasonal climates (i.e., higher absolute latitude) exhibit higher MR (both BMR and summit MR; [15,42,43]) and greater capacity for non-shivering thermogenesis [16]. To our knowledge, ours is the first study to identify a similar latitudinal gradient for maximum rewarming rate in endotherms. To cope with unpredictable climate and extreme variation in  $T_a$  associated with higher latitudes, individuals of temperate species should benefit from the capacity to rewarm rapidly. In contrast, maximum rewarming rates of tropical species were relatively low (Table 1, Fig. 3) even though all the tropical species in our analysis would regularly experience temperatures well below thermoneutrality and all readily express torpor. Despite their use of torpor, however, lower rewarming rates for tropical species likely reflect a relatively low thermoregulatory scope (i.e., lower range of  $T_b$  variation; [17]) compared to temperate species. The costs of slower rewarming in the tropics may be lower because of a smaller  $T_b$  differential between torpor and normothermia. It is clear that latitude is an important predictor of various physiological parameters [14,15,



**Fig. 2.** (a) Positive relationship between mass-corrected BMR residuals (mL O<sub>2</sub>/h; log<sub>10</sub> transformed) and rewarming rate (°C/min) for 23 bat species. (b) Partial residual plot demonstrating the effect of mass-corrected BMR (mL O<sub>2</sub>/h; log<sub>10</sub> transformed) on partial residual rewarming rate, taking the effect of absolute latitude into account. Dashed lines represent 95% confidence interval around the regression line

17] and the relationship between rewarming rate and latitude we observed suggests another example of adaptation to local climate and the importance of temperature as a selective force on energy expenditure and behavior of small mammals.

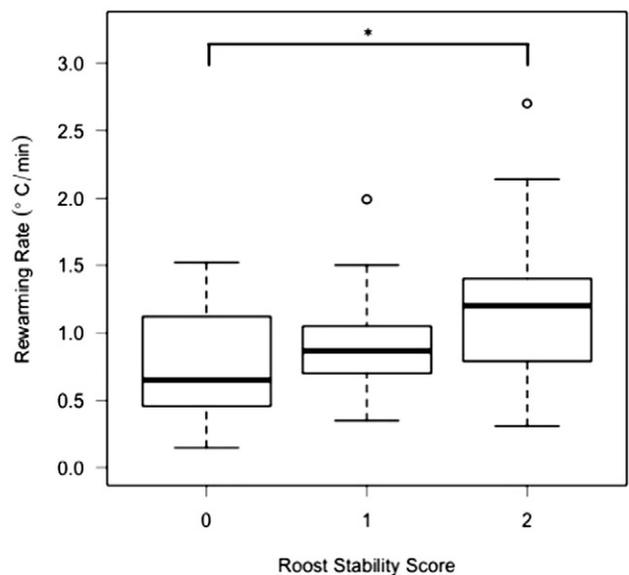
Consistent with our third, behavioral hypothesis, roost preference was correlated with rewarming rate (Table 2, Fig. 4). Although we predicted less thermally stable roost sites might allow for greater passive rewarming opportunities, we found that species roosting in less stable environments have slightly higher rewarming rates than those roosting in the most thermally stable environments. Well insulated roost sites that maintain relatively stable microclimates could be better suited to social thermoregulation if they retain more heat generated by the bats themselves. In some circumstances, this could provide greater energy savings than the passive rewarming opportunities provided by fluctuating T<sub>a</sub> in less thermally stable roost sites. However, this result should be treated cautiously because the need to treat roost stability as a categorical variable in our analysis likely reduced our ability to quantify its relationship to rewarming rate. For example, bats could also exploit spatial variation in microclimate within the roost structure to maximize rewarming opportunities but our analysis would not be able to detect this pattern. We suggest that future analyses of roost and/or



**Fig. 3.** Positive correlation between absolute latitude and rewarming rates (°C/min) for 45 bat species. Absolute latitude values for each species were taken to provide a measure of distance from the equator without differentiating between the two hemispheres. Dashed lines represent 95% confidence interval around the regression line.

microclimate preferences of bats would benefit from more precise T<sub>a</sub> measurements for a broad range of roost types and better data on spatial and temporal variation in roost temperature.

Contrary to our behavioral predictions, colony size was not an important predictor of rewarming rate. It is known that huddling with a large number of conspecifics can reduce heat loss to the environment, increase local T<sub>a</sub>, and facilitate passive rewarming for individuals via heat generated by huddle-mates, thereby minimizing the energetic cost of arousal [44,45]. One explanation for only a weak effect of colony-size in our analysis is that thermoregulatory capacity, local climate and thermal stability of roosts outweigh the effects of social thermoregulation. Perhaps only bat species that reside in temperate, thermally unstable roost sites or that lack sufficient thermogenic capacity are obligated to overcome thermoregulatory demands by clustering with



**Fig. 4.** Relationship between roost stability (i.e., roost score) and rewarming rate (°C/min) for 45 bat species. Roost score represents the temperature stability of roosts on a three-point scale ranging from 0 (highly stable roosts; e.g., caves) to 2 (highly unstable roosts; e.g., foliage). The \* represents a significant difference ( $p < 0.05$ ) between roost stability scores.

other individuals. We were restricted to the use of maximum colony size, rather than the average, because estimating mean colony size would require an accurate estimate of a large sample of different colony sizes of a given species. For many species this would be next to impossible and, therefore, so few species accounts report an “average” colony size and, instead, report minimum or maximum colony sizes. It is also possible that a measurement of maximum colony size is not ideal for capturing the thermoregulatory consequences of group size. This measure may not describe the most common colony size of a species, but may represent a “best-case scenario”.

Past studies of rewarming rate have either not controlled for phylogeny [7] or have corrected for phylogeny using independent contrasts which assume a significant impact of phylogeny on the trait of interest [8,9]. Ours is the first analysis to quantify the effect of phylogeny on rates of arousal from torpor across a range of bat species and our data suggest the effect of phylogeny is relatively small (Table 2;  $\lambda$  ranging from 0 to 0.40 [35]). However, intermediate values of  $\lambda$ , such as those reported here, are difficult to interpret because the “strength” of  $\lambda$  is context-dependent and can be influenced by both the scale of the dataset and/or convergent evolution [38,46]. Therefore, although we found values of  $\lambda$  approaching 0, we cautiously interpret phylogeny as having at least some limited effects on rewarming rates in bats (Table 2), a result which is qualitatively supported through visual inspection of Fig. 1 (i.e. rewarming rates range within clades). However, the relatively weak phylogenetic signal we observed, the large agreement between phylogenetic and non-phylogenetic analyses, and the strong influence of BMR, latitude, and roost preference supports the hypothesis that mammalian heterothermy has undergone extensive adaptive modification in response to climate and local environment, resulting in a range of thermoregulatory strategies spread throughout the phylogeny [17,44,47]. While phylogeny may determine the historic origin and evolution of heterothermy, the contemporary expression of torpor-related traits, like rewarming rate, appear to have been more strongly shaped by local climate, physiology, and behavior [e.g., 17].

In conclusion, our findings suggest that the costs of arousal from torpor exert selection pressure on rewarming capacity and that behavior and distribution of heterothermic endotherms influences the evolution of traits associated with thermoregulation. Some species can benefit from behavioral mechanisms and environmental conditions that favor passive rewarming [10,11] whereas others have evolved more rapid endogenous warming rates to reduce energetic costs. Our results contribute to the growing literature on the evolution of species-specific energetic traits in bats as well as other taxa and highlight that local climate, physiology, and behavior are likely mechanisms associated with the proliferation of heterothermic endotherms.

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