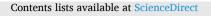
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Functional role of metabolic suppression in avian thermoregulation in the heat

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ABSTRACT

Hypometabolism arising from active metabolic suppression occurs in several contexts among endotherms, particularly during heterothermic states such as torpor. However, observed $Q_{10} \approx 1$ for avian resting metabolic rate within the thermoneutral zone, values far below the $Q_{10} = 2-3$ expected on the basis of Arrhenius effects, suggests hypometabolism also plays a role in birds' thermoregulation at environmental temperatures approaching or exceeding normothermic body temperature (T_b) . We evaluated the occurrence of hypometabolism during heat exposure among birds by re-analysing literature data to quantify changes in T_b and resting metabolic rate (RMR) near the upper boundary of the thermoneutral zone, at air temperatures (T_{air}) between the inflection above which T_b increases above normothermic levels ($T_{b.inf}$) and the upper critical limit of thermoneutrality (T_{uc}). Among the ~55 % of species in which $T_{uc} - T_{b.inf} > 0$, $Q_{10} < 2-3$ occurred in nine of 10 orders for which suitable data exist, indicating that hypometabolism during heat exposure is widespread across the avian phylogeny. Values of $Q_{10} < 2-3$ were not restricted to small body mass, as previously proposed. Our findings support the idea that metabolic suppression reduces avian metabolic heat production and hence evaporative cooling requirements during heat exposure, with reductions of 20–30 % in RMR in some species. Moreover, these findings add to evidence that hypometabolism is an important component of heat tolerance among endotherms such as birds and tropical arboreal mammals.

1. Introduction

Temperature has important and pervasive effects on the rates of biochemical reactions and overall metabolic rate (MR) via the Arrhenius effect, with a 10 $^{\circ}$ C increase in body temperature ($T_{\rm b}$) typically associated with a 2-3-fold increase in MR (Arrhenius, 1889; Christophersen, 1973). Arrhenius effects are quantified using the Q_{10} coefficient, the fractional change in rate associated with a 10 °C difference in $T_{\rm b}$ (Arrhenius, 1889; Christophersen, 1973). Although Arrhenius effects are best understood among ectotherms, in which the close coupling between environmental temperature (T_{env}) and body temperature (T_b) results in strong temperature-dependence of physiological and behavioural processes (Angilletta Jr., 2009; Christophersen, 1973; Huey et al., 2009), nevertheless, T_b-dependence of MR is also a feature of avian and mammalian thermoregulation (e.g., Geiser, 1988; Lyman et al., 1982; Lovegrove et al., 2014; Reher et al., 2018). During torpor and hibernation, for instance, Q_{10} for MR is often 2–3 when T_{env} is above the reduced T_b set point defended during heterothermy and animals are thermoconforming (Geiser 2004; Lyman et al., 1982; Lovegrove et al., 2014). Under these conditions, warmer T_{env} can result in increased MR, which may negatively affect the ability of hibernators, such as non-migratory bats at temperate and boreal latitudes, to survive the winter using their stored fat reserves (e.g., Humphries et al., 2002).

Recent evidence suggests some endotherms use hypometabolism to avoid Arrhenius effects on MR when T_b is elevated during heat exposure (Lovegrove et al., 2014; Reher et al., 2018; Reher and Dausmann, 2021). Malagasy bats (Macronycteris commersoni), for instance, decreased MR during hyperthermic torpor on very hot days when skin temperature tracked T_{env} , even when T_{b} exceeded normothermic levels (Reher and Dausmann, 2021). Diurnal hyperthermic torpor was also observed in four nocturnal arboreal mammals, including tarsiers (Tarsius syrichta), leading Lovegrove et al. (2014) to posit that T. syrichta uses active metabolic suppression to avoid Arrhenius effects on MR during the heat of the day. Reductions in the magnitude of Arrhenius effects on metabolic heat production may confer benefits by avoiding or delaying lethal hyperthermia and dehydration (Levesque et al., 2018; Lovegrove et al., 2014; Weathers, 1981), potentially from increased heat storage capacity or the use of non-evaporative heat dissipation avenues (Dawson and Hudson, 1970; Gerson et al., 2019; Williams and Tieleman, 2005).

Although most evidence for a role of metabolic suppression in reducing Arrhenius effects on MR comes from tropical mammals, there is also some evidence for $Q_{10} < 2-3$ in birds. Weathers (1981) reported

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 $Q_{10} \approx 1$ in three of four species he investigated, in individuals whose T_b increased within their thermoneutral zone (TNZ) and before the onset of thermoregulatory behaviours such as panting and the accompanying approximately linear increases in resting metabolic rate (RMR) that occur at T_{air} above the upper critical limit of thermoneutrality (T_{uc}). Lower-than-expected Q_{10} was also observed in budgerigars (*Melopsittacus undulatus*) (Weathers and Schoenbaechler, 1976). These authors reviewed the literature and noted that evidence for hypometabolism existed for 11 other bird species representing five orders (Weathers and Schoenbaechler, 1976), observations suggesting some birds employ active metabolic suppression during heat exposure to offset Arrhenius effects on MR.

Despite Weathers' observations of avian $Q_{10} < 2-3$ and the increasing evidence for a role of metabolic suppression during heterothermy among tropical mammals, no meta-analyses have been conducted to assess whether T_b -independence of MR at T_{env} approaching or exceeding normothermic T_b occurs across the avian phylogeny or whether it is limited to a subset of taxa. A priori, avoiding T_b -related increases in metabolic heat production within the TNZ may be expected to confer adaptive benefits by reducing water requirements for evaporative cooling as T_{env} approaches T_b . For this reason, we hypothesised that hypometabolism during heat exposure is widespread across the avian phylogeny and predicted that increases in T_b within the TNZ are associated lower-than-expected increases in RMR, evident as Q_{10} below the range of 2–3 associated with typical Arrhenius effects.

2. Methods and materials

2.1. Data sources

Data from the literature and unpublished data sources (Appendix, Table S1) were compiled for 106 bird species ranging in body mass (M_b) from \sim 7 g to \sim 700 g and representing 17 orders (Appendix, Table S1). The study sites for these species spanned various arid locations in South Africa, North America (USA), and Australia, as well as montane and coastal lowland areas in South Africa (multiple datasets exist for some species, for instance when conspecific populations were examined in >1 climatic region, yielding 115 datasets). Details of capture, holding conditions, and gas exchange measurements using flow-through respirometry with low chamber humidity (~1g H₂O m⁻³; Freeman et al., 2020, 2022, 2024) are provided in each of the published studies (Appendix, Table S1), all of which followed the methodologies described by Whitfield et al. (2015) and Czenze et al. (2020), ensuring comparability studies like the present one. In all these studies, for intraperitoneally-injected passive integrated transponder (PIT) tags were used to measure core T_b during respirometry trials (e.g. Czenze et al., 2020; Freeman et al., 2020, 2022, 2024; Whitfield et al., 2015). Furthermore, for each species in these studies food was withheld prior to gas exchange for periods greater than the gut passage time predicted from body mass by Karasov (1990). We excluded published datasets not collected using comparable methodology, particularly the maintenance of low chamber humidity by using high flow rates. For each species/population, we re-analysed T_b and RMR measured between $T_{air} =$ 25 °C-30 °C and the maximum Tair associated with thermal endpoints, as well as inflection T_{air} values for T_b ($T_{b.inf}$) and RMR (i.e., T_{uc}).

Our primary data inclusion criterion for subsequent analyses was an increase in T_b within the TNZ (i.e, $T_{uc} - T_{b.inf} > 0$), as for most species a Q_{10} value for MR cannot be estimated at $T_{air} > T_{uc}$ because of the obscuring effects of MR increases associated with active heat dissipation processes such as panting (Weathers and Schoenbaechler, 1976; Weathers, 1981). We also included the following species lacking a discernible T_{uc} : Namaqua sandgrouse (*Pterocles namaqua*), Freckled nightjar (*Caprimulgus tristigma*) and Rufous-cheeked nightjar (*Caprimulgus tristigma*) and Rufous-cheeked nightjar (*Caprimulgus tristigma*). Among species for which a T_{uc} was evident, values of $T_{uc} - T_{b.inf}$ were normally distributed (Shapiro-Wilk normality test, p = 0.29;

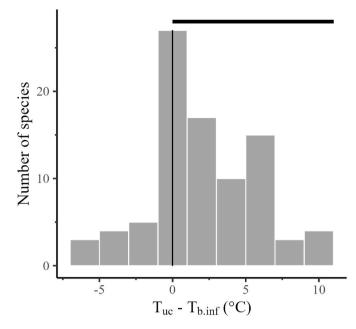


Fig. 1. Distribution of the difference between air temperature (T_{air}) inflections for body temperature (T_b) and resting metabolic rate ($T_{b.inf}$ and T_{uc} , respectively) for the 90 avian species or populations which had identified T_{uc} and $T_{b.inf}$. Data sets for which T_{uc} and/or $T_{b.inf}$ could not be identified were excluded from this analysis, as were outliers. The horizontal line above the histogram indicates data sets for which $T_{uc} - T_{b.inf} > 0$ and therefore met the first data inclusion criterion for this study.

Fig. 1). Data sets for which $T_{uc} - T_{b.inf} \leq 0$ were excluded from further analyses, as were those with sample sizes of ≤ 6 individuals.

2.2. Q_{10} calculations and analyses

For each species in the remaining subset of studies (Table 1), relationships between T_{air} , RMR and T_b were evaluated using generalised linear mixed models (GLMMs) with individual included as a random factor, using the *nlme* package version 3.1–163 (Pinheiro et al., 2023) in R 4.3.1 (R Core Team, 2023) following Freeman et al. (2022). Values of $T_{\rm b.inf}$ and $T_{\rm uc}$ were estimated using the package segmented (Muggeo, 2008) and served as lower and upper bounds of the T_{air} range over which we estimated Q10, unless Tbinf was below the lower limit of thermoneutrality ($T_{\rm lc}$), in which case the $T_{\rm lc}$ was used as the lower bound, or if there was no discernible T_{uc} , in which case the highest T_{air} setpoint was used as the upper bound. Average RMR and $T_{\rm b}$ values were then calculated for these Tair values using the GLMMs using the slopes for RMR at $T_{air} \leq T_{uc}$ and the T_b slope at $T_{air} \geq T_{b,inf}$ for $T_{air} \geq 28$. This approach was necessary because T_{b.inf} and T_{uc} usually did not occur exactly at one of the experimental T_{air} setpoints and therefore GLMMs were used to have a standardised, repeatable method of calculating Q_{10} for each species, although in some cases our estimated Q₁₀ may vary from those calculated using average RMR values at the nearest (± 1 °C) T_{air} setpoints (e.g., McKechnie et al., 2016a). The Q₁₀ for each species was calculated using the standard equation describing the Arrhenius effect (Lovegrove et al., 2014; Snyder and Nestler, 1990; Withers, 1992):

$$Q_{10} = \left(\frac{R_2}{R_1}\right)^{10/(T_2 - T_1)}$$

where R_1 and R_2 correspond to RMR (in Watts) at the $T_{b,inf}$ and T_{uc} , respectively, and T_1 and T_2 correspond to T_b at each of these respective T_{air} values. For several species, the $T_{b,inf}$ and T_{uc} occurred between experimental T_{air} setpoints and consequently, there were few (if any) RMR and T_b data between these inflection points. Therefore, to ensure

Table 1

 Q_{10} values, change in body temperature (ΔT_b) and body mass (M_b) of bird species in which T_b increased within the thermoneutral zone (TNZ). The magnitude of change in T_b (>1 °C) and the number of data points ($n \ge 7$) for resting metabolic rate (RMR) within this range were used as additional criteria for subsequent phylogenetic analyses. Species included in the final analyses are indicated in bold font.

Species	Order	Climate	Location	M _b (g)	Q10	ΔT_b (°C)	n	Ref
Apus affinis	Apodiformes	Lowland	South Africa	25.9	0.8	0.3	5	5
Apus caffer	Apodiformes	Montane	South Africa	27.3	0.8	0.4	7	5
Bycanistes bucinator	Bucerotiformes	Lowland	South Africa	702.7	1.1	0.1	1	1
Tockus leucomelas	Bucerotiformes	Arid	South Africa	219.6	1.2	2.0	33	17
Aegotheles cristatus	Caprimulgiformes	Arid	Australia	44	1.4	1.4	16	15
Caprimulgus rufigena	Caprimulgiformes	Arid	South Africa	52.2	1.3	2.2	64	12
Caprimulgus tristigma	Caprimulgiformes	Arid	South Africa	67.3	1.1	2.3	75	12
Chordeiles acutipennis	Caprimulgiformes	Arid	South Africa	50.6	1.3	1.7	18	15
Phalaenoptilus nuttallii	Caprimulgiformes	Arid	North America	44.3	1.3	1.5	38	15
Burhinus capensis	Charadriformes	Arid	South Africa	434.3	0.9	1.6	50	3
Colius colius	Coliiformes	Arid	South Africa	36.2	0.1	0.9	11	7
Oena capensis	Columbiformes	Arid	South Africa	37.1	1.1	2.4	51	9
Streptopelia capicola	Columbiformes	Arid	South Africa	147.5	1.4	1.3	24	9
Coracias caudatus	Coraciiformes	Arid	South Africa	94.5	0.7	1.5	38	13
Halcyon albiventris	Coraciiformes	Lowland	South Africa	62.1	0.6	0.8	10	5
Merops persicus	Coraciiformes	Lowland	South Africa	46.3	0.1	1.0	11	5
Cuculus gularis	Cuculiformes	Arid	South Africa	110	3.7	2.0	8	13
Acanthagenys rufogularis	Passeriformes	Arid	Australia	41.7	1.3	1.7	15	10
Andropadus importunus	Passeriformes	Lowland	South Africa	30.6	1.1	0.5	9	5
Calandrella cinerea	Passeriformes	Montane	South Africa	25.5	1.3	2.4	42	5
Calendulauda africanoides	Passeriformes	Arid	South Africa	23	0.7	0.1	2	2
Campicoloides bifasciatus	Passeriformes	Montane	South Africa	34.8	0.2	0.9	7	5
Certhilauda albescens	Passeriformes	Arid	South Africa	28.2	0.5	0.1	3	6
Certhilauda subcoronata	Passeriformes	Arid	South Africa	40	0.8	1.3	20	2
Chersomanes albofasciata	Passeriformes	Montane	South Africa	28	0.8	2.0	32	5
•	Passeriformes	Arid	South Africa	28 25			32 16	2
Chersomanes albofasciata				25 29.6	0.5	1.2	30	2 5
Cossypha natalensis	Passeriformes	Lowland	South Africa		1.5	2.0		
Cracticus torquatus	Passeriformes	Arid	Australia	86	1.2	1.9	19	10
Emberiza capensis	Passeriformes	Arid	South Africa	19.7	0.8	0.5	6	6
Emberiza impetuani	Passeriformes	Arid	South Africa	14.2	0.5	1.5	15	5
Eremopterix verticalis	Passeriformes	Arid	South Africa	16	0.8	0.9	13	2
Euplectes capensis	Passeriformes	Arid	South Africa	30.6	0.7	1.3	16	6
Euplectes orix	Passeriformes	Lowland	South Africa	16.8	0.9	0.7	10	5
Hypargos margaritatus	Passeriformes	Lowland	South Africa	12.5	0.5	0.8	8	6
Laniarius atrococcineus	Passeriformes	Arid	South Africa	46.2	0.6	0.8	9	2
Lanius collaris	Passeriformes	Arid	South Africa	37.3	0.1	0.6	15	2
Lanius collaris	Passeriformes	Lowland	South Africa	40.6	0.3	0.7	10	5
Lanius collaris	Passeriformes	Montane	South Africa	39.7	0.3	0.5	17	5
Myrmecocichla formicivora	Passeriformes	Montane	South Africa	48.3	1	2.2	25	5
Petrochelidon spilodera	Passeriformes	Montane	South Africa	21	0.2	0.9	22	5
Ploceus ocularis	Passeriformes	Lowland	South Africa	26.1	1.3	1.0	11	5
Ploceus subaureus	Passeriformes	Lowland	South Africa	28.5	1	0.6	6	5
Ploceus velatus	Passeriformes	Montane	South Africa	28.9	0.7	1.1	11	5
Pomatostomus ruficeps	Passeriformes	Arid	Australia	52	0.8	0.1	2	10
Prinia maculosa	Passeriformes	Arid	South Africa	7.4	0.5	0.6	6	6
Prinia subflava	Passeriformes	Lowland	South Africa	9.1	0.2	0.1	0	5
Pycnonotus tricolor	Passeriformes	Lowland	South Africa	36.8	0.7	0.4	4	5
Pycnonotus tricolor	Passeriformes	Montane	South Africa	40.2	1.2	1.7	22	5
Quelea quelea	Passeriformes	Montane	South Africa	17.9	0.8	2.9	81	4
Serinus albogularis	Passeriformes	Arid	South Africa	22.1	1.1	1.0	14	6
Spermestes cucullata	Passeriformes	Lowland	South Africa	9.1	0.7	1.6	19	5
Spizocorys starki	Passeriformes	Arid	South Africa	17.9	0.7	0.9	18	2
Spreo bicolor	Passeriformes	Montane	South Africa	99.3	0.8	0.1	1	5
Felophorus zeylonus	Passeriformes	Montane	South Africa	66.8	0.5	0.9	11	5
Telophonus zeytonus Toxostoma curvirostre	Passeriformes	Arid	North America	70.6	1.1	1.3	20	14
Zosterops pallidus	Passeriformes	Arid	South Africa	7.7	0.6	0.6	20 9	2
Zosterops pallidus Zosterops pallidus	Passeriformes Passeriformes	Arid Arid	South Africa	7./ 9.6		0.6 2.0	9 21	2 6
	Passeriformes	Lowland	South Africa		0.7			5
Zosterops virens Friebolaema laucomelae				11	0.7	0.3	1	
Fricholaema leucomelas	Piciformes	Arid	South Africa	34.2	0.6	0.2	7	8
Psephotellus varius	Psittaciformes	Arid	Australia	54.6	0.9	1.3	17	11
Pterocles burchelli	Pterocliformes	Arid	South Africa	192.9	1.2	2.4	26	8
Pterocles namaqua	Pterocliformes	Arid	South Africa	165.8	1.4	4.2	94	12
Megascops kennicottii	Strigiformes	Arid	North America	101	1.7	2.6	34	16

Sources: (1) Coulson et al., unpubl; (2) Czenze et al. (2020); (3) Czenze et al. (2021); (4) Freeman et al. (2020); (5) Freeman et al. (2022); (6) Freeman et al. (2024); (7) Freeman et al., unpubl; (8) McKechnie et al. (2016a); (9) McKechnie et al. (2016b); (10) McKechnie et al. (2017); (11) McWhorter et al. (2018); (12) O'Connor et al. (2017); (13) Smit et al. (2018); (14) Smith et al., 2017; (15) Talbot et al. (2017); (16) Talbot et al. (2018); (17) van Jaarsveld et al. (2021).

confidence in the Q_{10} values we excluded species with ≤ 6 data points between T_{uc} and $T_{b.inf}$ or for which T_b changed by < 1 °C (Fig. 2). The Q_{10} data included in the final data set were then mapped onto an avian maximum likelihood tree constructed using phylogenies from www.bir

dtree.org (Jetz et al., 2012) using the Hackett backbone (Hackett et al., 2008) following Freeman et al. (2022), using *Mesquite* (Maddison and Maddison, 2014).

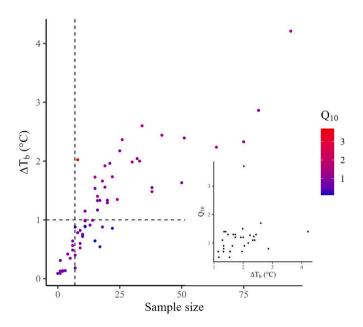


Fig. 2. Relationship between change in T_b (°C) and the number of data points. Inclusion criterion for Q_{10} values based on the sample size ($n \geq 7$, horizontal dashed line) of data points within the range over which Q_{10} was calculated and the change in T_b ($\Delta T_b \geq 1$, vertical dashed line). Each species' Q_{10} is indicated by colour, with higher values indicated in red and low values shown in blue. with The inset graph shows the nonsignificant relationship (p=0.17) between the calculated Q_{10} and change in T_b for $\Delta T_b \geq 1$. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

3. Results

Values of $T_{\rm uc} - T_{\rm b.inf}$ among species for which a clear $T_{\rm uc}$ exists ranged from $-7 \,^{\circ}$ C to 11 $^{\circ}$ C (Fig. 1) with the Namaqua dove (*Oena capensis*) as an outlier at 17.7 $^{\circ}$ C, whereas for the sandgrouse and two caprimulgids without a $T_{\rm uc}$ the corresponding range was 19.6 $^{\circ}$ C-32 $^{\circ}$ C. Of the 106 species for which data were available, 58 species representing 13 orders met the criterion of $T_{\rm uc} - T_{\rm b.inf} > 0$ (for three of which data were available for populations in >1 climatic region; Table 1; Fig. 2). Of these 58 species, 30 (with one, *Chersomanes albofasciata*, occurring in both arid and montane regions), also met our second data inclusion criterion

(Table 1; Figs. 2-4). Among these 30 species, representing 10 orders with $M_{\rm b}$ ranging from ~9 to ~430g, Q_{10} ranged from 0.5 to 3.8, with Q₁₀ < 1 in species including bronze mannikin (Spermestes cucullata) and Cape turtle dove (Streptopelia capicola) and $Q_{10} = 1-2$ in species including Namaqua sandgrouse (Pterocles namaqua), spotted thick-knee (Burhinus capensis) and Western screech owl (Megascops kennicottii). In only one species, African cuckoo (*Cuculus gularis*), was $Q_{10} > 2$ ($Q_{10} =$ 3.7, Fig. 4). When C. gularis was excluded, the mean \pm SD Q₁₀ for all included species was 1.02 \pm 0.32. The proportional change in RMR between the $T_{\rm b.inf}$ and $T_{\rm uc}$ ranged from \sim -50 – 150 % (Fig. 3) indicating that RMR decreased for some species (i.e., $Q_{10} < 1$), but increased on others ($Q_{10} > 1$). However, variation in Q_{10} did not show any obvious phylogenetic pattern (Fig. 4), with a wide range of values in some orders, including passerines ($Q_{10} = 0.5-1.5$). We found that Q_{10} values were not significantly related to body mass (p = 0.38) or change in T_b (p = 0.17).

4. Discussion

Our analyses support the hypothesis that metabolic suppression and reduced Arrhenius effects are a general feature of avian thermoregulation in the heat, with smaller-than-expected increases in RMR with increasing $T_{\rm b}$ within the TNZ evident in species representing nine orders. These findings confirm earlier observations by Weathers (1981) and Weathers and Schoenbaechler (1976), but also argue against these authors' conclusion that $T_{\rm b}$ -independent thermoneutral RMR is confined to species with $M_{\rm b} < 150$ g; the four species in our dataset with $M_{\rm b} > 150$ g showed Q_{10} between 0.9 and 1.4. The phylogenetic diversity of species in which metabolic suppression occurs during heat exposure raises the possibility of this being a pleisiomorphic trait among birds.

Our results also support the notion that hypometabolism during heat exposure reduces the amount of metabolic heat produced and hence heat dissipation requirements (Weathers and Schoenbaechler, 1976). In red-billed queleas (*Quelea quelea*, $Q_{10} = 0.8$), for example, the observed RMR of 0.44 W at $T_{air} = 43.8$ °C is 72 % of the 0.61 W expected if $Q_{10} = 2.5$. Similarly, the RMR = 1.06 W of Namaqua sandgrouse (*Pterocles namaqua*, $Q_{10} = 1.4$) at $T_{air} = 60$ °C is equivalent to 79 % of the RMR = 1.34 W expected if $Q_{10} = 2.5$. Avoiding increases in metabolic heat production with increasing T_b has significant implications for birds' water economy by decreasing evaporative cooling requirements (Weathers, 1981; Weathers and Schoenbaechler, 1976). These savings are greatest for species that lack a T_{uc} ; in *P. namaqua* and *C. rufigena*, $Q_{10} = 2.5$ for RMR would result in rates of EWL equivalent to 21 and 11 %

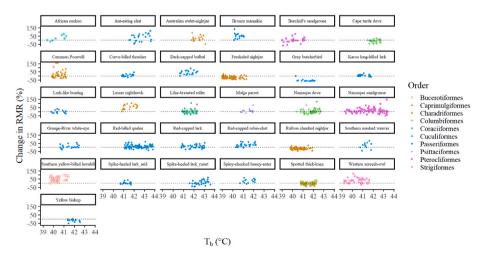


Fig. 3. Relative change in resting metabolic rates (RMR) as body temperature (T_b) increases for each of the 30 species which had a T_b inflection below their upper critical temperature of thermoneutrality (T_{uc}) and seven or more weathers data points within this range of air temperatures. Each species' order is indicated by colour and shape of the data points. Data are included for two populations of spike-heeled lark, one inhabiting a desert habitat ("_arid") and the other montane grasslands ("_mont"). (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

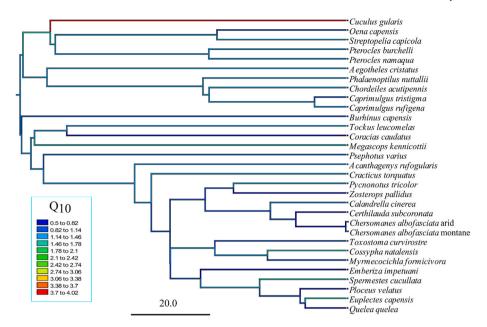


Fig. 4. The phylogenetic tree of the 30 analysed species, pruned from the maximum-likelihood phylogeny (Hackett et al., 2008). Line colour indicates the Q_{10} value of the species, with red indicating the highest Q_{10} values (\geq 3.7) and dark blue the lowest (0.5 – 0.84). (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

higher, respectively, compared to observed EWL at the highest T_{air} these species reached (Czenze et al., 2021; O'Connor et al., 2017), assuming unchanged ratios of evaporative heat loss to metabolic heat production. The physiological significance of lower-than-expected avian Q_{10} in terms of water balance was dismissed by Tieleman and Williams (1999). However, their arguments in this regard do not stand up to close scrutiny, as they were modelled closely on those of Snyder and Nestler (1990) and Heldmaier and Ruf (1992) against the use of Q_{10} values for inferring mechanisms involved in entry into hibernation or torpor, making their applicability to thermoregulation during heat exposure questionable. In reality, avoiding Arrhenius effects on metabolic heat production can equate to significant water savings.

In addition to confirming that reduced Arrhenius effects occur widely among birds, our analysis reveals unexpected interspecific variation in Q_{10} values. Among ectotherms, variation in Q_{10} for MR is thought to vary with body size (Barlow 1961; Hölker, 2006; Rao and Bullock, 1954, but see also Scholander et al., 1953). Additionally, variation in ectotherm Q_{10} has been attributed to acclimatisation to T_{env} or climate adaptation (DeLiberto et al., 2022; Rao and Bullock, 1954; Addo-Bediako et al., 2002; Nielsen et al., 1999). Sources of variation in the Q_{10} of MR within the TNZ in endotherms has received far less attention. Our findings here do not support the argument of Weathers and Schoenbaechler (1976) that Q_{10} values vary with body mass. The possibility of climate adaptation in endotherm Q_{10} remains to be investigated, but would require a larger data set spanning a wider range of latitudes and habitats than included in our present analysis.

Our data confirm metabolic suppression during heat exposure is widespread across the avian phylogeny, however, too few data are currently available to convincingly test hypotheses concerning adaptive variation in this trait. For instance, in tropical and subtropical lowland habitats characterized by high humidity, where birds have evolved pronounced hyperthermia tolerance (Weathers 1981; Freeman et al., 2022), avoiding Arrhenius effects on metabolic heat production may significantly benefit heat balance during hot weather. One constraint on the availability of suitable data concerns the fact that studies we reviewed for this analysis focussed more strongly on thermoregulation at $T_{air} > T_b$ than on relationships between T_b and RMR within the TNZ; whereas increments of 2 °C in experimental T_{air} were used at $T_{air} > 40$ °C, larger increments of 4 °C below $T_{air} = 40$ °C hindered our ability

to accurately estimate Q_{10} within the TNZ (e.g., Czenze et al., 2020; Whitfield et al., 2015, Table 1, Appendix, Table S1). We recommend future studies exploring Arrhenius effects on avian metabolic rates in the TNZ involve physiological measurements over smaller increments in experimental T_{air} .

Metabolic heat production associated with heat dissipation pathways, particularly panting, is the primary constraint on the avian capacity to avoid lethal hyperthermia during extreme heat exposure when Tenv exceeds Tb (McKechnie et al., 2021). Avoidance of Arrhenius effects on RMR, similar to those we have demonstrated here within the TNZ, are also likely to confer adaptive benefits at $T_{env} > T_{uc}$, even during bouts of activity. However, estimating $Q_{10}\ \text{for RMR}$ under these conditions is challenging, as doing so requires that increases in overall MR be partitioned into those reflecting increasing $T_{\rm b}$ versus those associated with panting or other evaporative cooling pathways. Partitioning increases in RMR in this way may be possible when bouts of panting are intermittent and rates of gas exchange are measured in respirometry systems with rapid equilibrium times (Lasiewski et al., 1966) achieved by combining a small chamber with rapid flow rates (see, for example, the short-term fluctuations in CO₂ production and EWL in Figure 5D of W.A. Talbot et al., 2017).

The evidence we present here for widespread avian hypometabolism during heat exposure confirms that metabolic suppression plays a role in heat tolerance among both birds (Weathers and Schoenbaechler 1976; Weathers 1981) and mammals (Lovegrove et al., 2014; Reher et al., 2018). It also supports arguments that the capacity for hypometabolism is a plesiomorphic trait among both classes (Lovegrove et al., 2014). Although, many unanswered questions about the role of avian hypometabolism during heat exposure remain. First, why do species vary in terms of whether $T_{\rm b}$ starts increasing within the TNZ, or only from the $T_{\rm uc}$ upwards? The lack of a bimodal distribution in $T_{\rm uc}$ - $T_{\rm b.inf}$ (Fig. 1) suggests a physiological continuum rather than discrete responses. Second, how does the functional role of hypometabolism vary with biotic and abiotic variables? For instance, is there stronger selection for active metabolic suppression in species that forage in exposed microsites and experience higher T_{env} than species foraging in shaded locations? An additional consideration would be including a wider array of phylogenetically diverse species with an increased range of body masses. Such an approach may increase our understanding around the relationship between functional traits and hypometabolism. Finally, all the data we included in our analyses here were collected under conditions of low humidity. However, reductions in metabolic production and evaporative cooling requirements are, *a priori*, expected to confer benefits both in terms of water conservation in arid environments as well as hyper-thermia avoidance in humid environments where evaporative heat dissipation is impeded, making humidity a variable of interest for future research on avian hypometabolism during heat exposure.

Data availability

The data included in this analysis are available at https://data.me ndeley.com/datasets/7283n2png5/1.

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CRediT authorship contribution statement

Jochen J. Voges: Writing – original draft, Formal analysis, Data curation. Marc T. Freeman: Writing – review & editing, Supervision, Formal analysis, Conceptualization. Blair O. Wolf: Writing – review & editing, Supervision. Andrew E. McKechnie: Writing – review & editing, Supervision, Funding acquisition, Conceptualization.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

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